IMA-2021
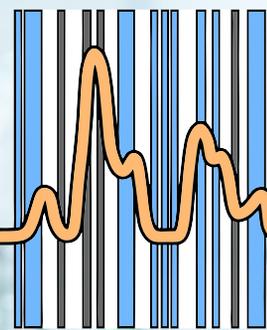
12th International Conference on Instrumental Methods of Analysis
Modern Trends and Applications

20-23 September 2021 | VIRTUAL EVENT

BOOK OF ABSTRACTS

Organized by

ARISTOTLE UNIVERSITY OF THESSALONIKI

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CONFERENCE DETAILS

The 12th International Conference on "Instrumental Methods of Analysis" (IMA-2021) takes place between 20-23 September 2021 as a Virtual event organized by the Aristotle University of Thessaloniki and the National Technical University of Athens. IMA is a biannual series of conferences that started in 1999 and covers all areas of Chemical Analysis, including the development of new techniques, modern trends, and applications in a wide range of scientific disciplines. Up to date several leading analytical chemists from Greece and abroad have presented their research work at the previous IMA meetings. Considering existing pandemic conditions, the relevant uncertainty on international travelling and the difficulty of gathering large meeting groups, our decision to organize IMA-2021 with virtual presentations and attendance ensures safety for all participants while at the same time it provides the opportunity for high-level analytical scientists from all around the world to promote their relevant research.

For the 12th IMA conference (first time in Virtual format), it is our ambition to bring together some of the most talented and innovative analytical chemists from all over the world for an excellent scientific online conference. The scientific program consists of keynote lectures, oral and poster presentations, given by a wide mix of scientists and engineers, ranging from the established academics and world leaders in analytical chemistry to the dynamic and ambitious postgraduate students. Furthermore, contributions from companies, including presentations of new instrumentation, new applications, and assessment of future commercial trends and opportunities are also included.

TOPICS

- Spectrometric and Electrometric analysis
- Chromatographic, Mass Spectrometric and Microscopic analysis
- Thermal analysis
- Proteomics, Metabolomics, Metallomics and Elemental Speciation analysis
- Chemical- and bio- sensors
- Field analysis - Mobile analytical instruments
- Miniaturized analytical systems (Lab-on-a-Chip)
- Micro- and Nano- fluidics
- Immunoassays
- Electrophoretic separation techniques
- Sampling techniques and strategies
- Sample handling and preparation
- Robotics and Automation
- Quality control-quality assurance in analysis
- Metrology
- Environmental, Biomedical, Pharmaceutical, Food analysis

- Materials Analysis (Nanomaterials, Smart/Advanced Materials)Archaeometry
- Analytical chemistry markets and possibilities for commercialization

Special sessions on the following themes

- AEROSOL METROLOGY
- ADVANCED X-RAY TECHNIQUES
- VIRUS SPREAD ANALYTICS

AWARDS

The most outstanding presentations during IMA-2021 will receive one of the awards from the following categories:

- **Best presentation related to X-ray Spectroscopy.** The best oral presentation with X-ray Spectroscopy research will receive an award sponsored by the European X-ray Spectrometry Association.
- **Best poster award prize.** The best poster presentation will receive an award of 500 CHF sponsored by the new open-access journal MDPI Analytica. Apart from the significance and the originality of the poster, the presentation impact will be also evaluated through a short 3-min presentation of the work by the presenting author during the poster session. Therefore, participating with a short video or live presentation is a requirement to be included in the competition for this award.
- **Best oral presentation award in the field of sensors, instrumentation and multiphysics simulation.** The best oral presentation dealing with Chemical and biosensors, Multiphysics phenomena simulation, Field analysis—mobile analytical instruments, Robotics and automation, Miniaturized analytical systems (Lab-on-a-Chip), micro and nanofluidics, Multiscale modelling and characterization of materials and processes, Design and optimization of devices and systems with numerical approaches/techniques, will receive an award of 500 $ sponsored by the open-access journal MDPI Micromachines.

Presented papers will be evaluated by the Award Committee according to the following criteria: Significance of the paper, Originality, Presentation impact, Young researcher. The winners of IMA-2021 awards will be announced during the closing ceremony of the conference.

PROCEEDINGS

A book of abstracts including submitted abstracts will be shared to the participants. In addition, all accepted abstracts will be published as open-access documents in the MDPI Applied Sciences (impact factor 2.679) Special issue dedicated to IMA-2021.

PUBLICATION IN JOURNALS

IMA-2021 provides several possibilities for the publication of presented studies in one of the collaborating peer-review journals.

Participants have the option to submit their full papers in the Special Issue "Selected Papers from the 12th International Conference on "Instrumental Methods of Analysis" (IMA-2021)" of the new open-access journal **MDPI Analytica**. All submissions through the conference will receive a full-waiver for article processing charges.

Participants have the option to submit their full papers in the Special Issue "12th International Conference on Instrumental Methods of Analysis - IMA-2021" of open-access journal **MDPI Applied Sciences** (Impact factor 2.679). All submissions through the conference will receive a 20% discount for article processing charges.

Selected works related to chemical and biosensors, multiphysics phenomena simulation, field analysis—mobile analytical instruments, robotics and automation, miniaturized analytical systems (lab-on-a-chip), micro and nanofluidics, multiscale modelling and characterization of materials and processes, design and optimization of devices and systems with numerical approaches/techniques will be published in the Special Issue related to IMA-2021 of journal **MDPI Micromachines** (Impact factor 2.891). All submissions through the conference will receive a 500 CHF discount for article processing charges.

Selected works related to the application of analytical techniques for sustainable water treatment will be published in the Special Issue "Sustainable Approaches in Water Treatment Practices, Presenting Minimum Environmental Impact "of journal **MDPI Water** (Impact factor 3.103).

CERTIFICATE OF ATTENDANCE

All registered participants are eligible to receive a certification of attendance at the end of the conference. The certificate will be sent electronically to online participants.
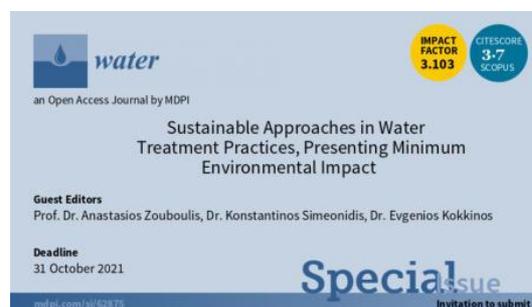
PROGRAM AT A GLANCE

Monday 20 September 2021

	09.30-10.00	Greetings
Session A1-Plenary	10.00-12.00	Virus spread analytics, Drug research
	12.00-12.15	Break
Session A2	12.15-14.00	Aerosol metrology, Advanced X-Ray techniques I
	14.00-14.30	Break
Session A3	14.30-16.15	Aerosol metrology, Advanced X-Ray techniques II
	16.15-16.30	Break
Session A4	16.30-18.30	Spectroscopy and applications

Tuesday 21 September 2021

Session B1-Plenary	09.30-11.30	Trends in analytical techniques
	11.30-11.45	Break
Session B2	11.45-12.30	Chromatography, Mass spectrometry I
	12.30-14.00	Virtual exhibition
	14.00-14.30	Break
Session B3	14.30-16.15	Chromatography, Mass spectrometry II
	16.15-16.30	Break
Session B4	16.30-18.15	Chromatography, Speciation, Bioanalytics, Bioseparation
	18.15-20.00	Poster session I

Wednesday 22 September 2021

Session C1	09.30-11.30	Food analysis I
	11.30-11.45	Break
Session C2	11.45-12.45	Food analysis II
	13.00-14.00	Poster session II
	14.00-14.45	Break
Session C3	14.45-16.15	Chemistry associations
	16.15-16.30	Break
Session C4	16.30-18.30	Environmental analysis, Chromatography, Mass spectrometry
	18.30-18.45	Break
	18.45-20.00	Poster session III

Thursday 23 September 2021

Session D1	09.30-11.30	Materials analysis
	11.30-11.45	Break
Session D2	11.45-13.45	Electrochemistry, Sensors, Archaeometry
	13.45-14.30	Break
Session D3	14.30-15.30	Mobile analytical instruments, Robotics and Automation, Chemometrics
	15.30-16.15	Closing Ceremony-Awards

Monday 20 September 2021

SESSION A1

Plenary Lectures

Virus spread analytics | Drug research

Chair: M. Mitrakas, M. Ochsenkühn, F. Tsopelas

Covid-19: the world's first environmental disease - integrative analytical and computational perspectives

D.A. Sarigiannis

[1]Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[2]University Institute of Advanced Study IUSS, 27100 Pavia, Italy

E-mail: sarigiannis@auth.gr

COVID-19 challenges the way the global human community addresses public health protection from infectious diseases and the factors that determine their prevalence. The environmental dimension of the disease spread and of the associated health risk is an essential part of the pandemic puzzle. Ambient air temperature, total humidity and UV-B radiation flux are key parameters determining the potency of the virus to "dock" itself onto the human lung epithelium and thus influence the effective intrinsic transmissibility of different SARS-CoV-2 variants. Prior exposure to immunotoxicants, such as perfluorinated compounds, has been elated to negatively influence the ability of the human immune system to respond effectively to SARS-CoV-2 contamination. Alas, recent human biomonitoring studies have demonstrated that most Europeans are exposed to PFCs at non-negligible levels. Exposure to fine and ultra-fine particles through the air has been shown to increase human susceptibility to SARS-CoV-2, resulting in a 9% increase in mortality rate of vulnerable individuals for 1 ug/m^3 increase in ambient air PM2.5. Aerosol-based dispersion of the virus has been deemed the second most important pathway of exposure after direct exposure to biological droplets from sneezing or coughing. Our ability to detect and monitor continuously low levels of environmental pollution may well prove to be an essential element for an effective strategy countering the spread of COVID-19. Wastewater epidemiology has been a very important tool serving predictive models of viral spread dynamics in humans with valuable information regarding the overall viral load independently from the incidence rate of the disease in the community. Enhancement of the current state-of-the-art with metagenomics-based pathochip microarrays may significantly increase the applicability and reliability of waste and wastewater epidemiology accounting explicitly for the total pathogenic load affecting humans. When it comes to humans, two are the main methodological approaches for contamination detection: (i) detection of genetic material (or proteins) from SARS-CoV-2 in respiratory-tract specimens, or (ii) the detection of antibodies against the virus in blood samples. Detection of viral RNA is the only accepted diagnostics to confirm active infection. Analytical science is important to understand what is determined/detected in each type of assay and assess and explain the limitations of each approach (limits of detection and quantification, existence and practical implications of false positives or false negatives). Integration of the data gathered through these tools requires both a comprehensive framework for instrumental analysis that can cover efficiently all environmental compartments, human matrices and even waste and wastewater, and an intelligent multi-scalar dynamic computational platform that allows us to capture effectively the environmental and social dynamics that govern the disease spread and severity. The COVID-19 Risk Evaluation (CORE) system incorporates all these elements and, since the beginning of the pandemic in Europe, provides daily analyses of the pandemic dynamics and the time trends of the associated health risk. CORE allows long-term predictions and scenario analysis taking into account non-pharmacological interventions and pharmacological solutions, while considering the evolutionary pressure on the virus leading to an extended number of mutations that present partial immune escape features.

Wastewater-based Epidemiology during COVID-19 pandemic

N.S. Thomaidis

Laboratory of Analytical Chemistry, Department of Chemistry, National and Kapodistrian University of Athens, Panepistimioupolis Zografou, 15771 Athens, Greece

E-mail: ntho@chem.uoa.gr

December 2019 will be remembered as the springboard of Coronavirus Disease 2019 (COVID-19), which causes viral pneumonia and many other health effects. However, the measures taken to restrict the virus spread affected societies, economies and indirectly mental and physical health, habits and lifestyle. Wastewater-based epidemiology became a promising chemical tool for monitoring SARS-CoV-2 infection dynamics, as well as population's chemical exposure, public health, lifestyle and how the ongoing pandemic affect the aforementioned parameters [1, 2, 3]. In the present study, the effects of COVID-19 pandemic on population of Athens, employing WBE, are presented. During the first wave of the pandemic (March-April 2020), the analysis of wastewater revealed significant increase in surfactants (+196%) and biocides (+152%), while important decrease was observed for tobacco and industrial chemicals (-33% and -52% respectively). For illicit drugs, and especially for stimulants such as amphetamine, methamphetamine and cocaine, an increasing trend was observed from March 2020 to March 2021. In addition, during March 2020, when there was not effective treatment for SARS-CoV-2, antiviral and antibiotics demonstrated significant increases (+170% and 57%), while NSAIDs use was decreased (-27%). Oxazepam, one of the most common benzodiazepine, reached peak use levels during December 2020, while during January to March 2021 the use levels remained high. On 7th of November, when the 2nd total lockdown was announced, SARS-CoV-2 viral load showed a stable trend until the end of January when extremely high viral loads were observed for the first time in Athens. For the following three months (February to May), the observed increase was resulted from SARS-CoV-2 alpha-variant, as revealed by NGS analysis of RNA extracts. After the end of the restrictions and lockdown (May to June) SARS-CoV-2 viral load demonstrated a decreasing trend during June, while the prevalence of delta-variant and the absence of restriction measures resulted in a fourth wave with high viral load in wastewater in July and August 2021. The time-trend analysis of viral load in wastewater of Athens were correlated with infections dynamics and epidemiological variables, demonstrating the potential of WBE to be used as an early warning system of the pandemic dynamics.

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Biomimetic chromatography to predict *in vivo* distribution of drug discovery compounds

K. Valko

UCL, School of Pharmacy, London, UK, Bio-Mimetic Chromatography Ltd, Stevenage Herts, UK

E-mail: k.valko@ucl.ac.uk

Chromatography is a powerful analytical separation technique which can also be used to measure compound interactions with stationary phases. Biomimetic chromatography is designed to mimic the biological partition processes of compounds by using specific protein and phospholipid stationary phases. The carefully standardized gradient chromatographic retention times obtained on biomimetic stationary phases are used in models of the *in vivo* distribution of compounds [1]. The volume of distribution, tissue binding, and various toxicity parameters can therefore be estimated from the measured biomimetic chromatographic properties of early drug discovery compounds. The non-specific binding of compounds can be predicted by the sum, of the albumin and phospholipid binding that also showed an excellent inverse correlation with the drug efficiency [2]. The HPLC based estimation of drug efficiency together with the in vitro potency can be used to estimate the dose [3] that is required to achieve the desired free concentration of the drug at the site of action. In this way organic chemists are able to use this data at early stages of the lead optimisation process. In this way, the drug design and discovery processes can be accelerated with reduced costs. The models have been successfully used in over 20 drug discovery programs ranging from CNS to respiratory targets. More recently, the applicability of the method has been investigated for potential peptide therapeutics where the traditional methods for lipophilicity determination and protein binding could not be used.
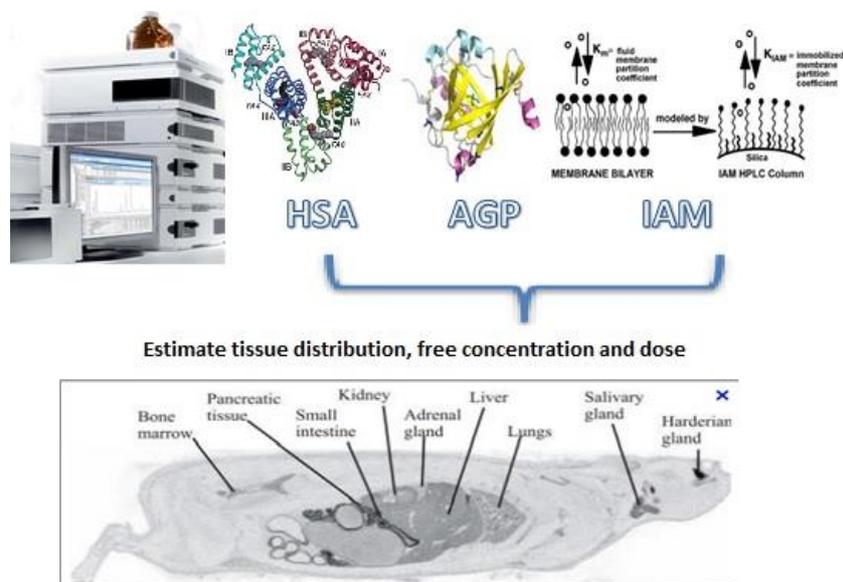
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Magnetoliposomes for efficient AC-Magnetic Field Remote Controlled Drug Release

M.E. Fortes Brollo, M.C. Concepción Serrano, <u>M.d.P. Puerto Morales</u>

Instituto de Ciencia de Materiales de Madrid, ICMM/CSIC, Sor Juana Inés de la Cruz 3, 28049 Madrid, Spain
E-mail: puerto@icmm.csic.es

Magnetic hyperthermia is an interesting alternative for some kind of cancer difficult to remove, such as brain tumours, or especially resistant to other therapies, such as triple-negative breast cancer [1]. However, there are still a limited number of approved nanoformulations and clinical trials due to some drawbacks of this therapy, i.e. lack of effective accumulation of nanoparticles in the tumour after intravenous injection, alteration of heating due to agglomeration of nanoparticles in lysosomes inside the cell, and uncertainty and lack of control of the temperature at the nanoscale.

Here we will show that by combining liposomes, magnetic nanoparticles and a drug, we were able to prepare a nanocarrier (< 200 nm) for directing drugs to specific tissues or organs, avoiding side effects [2]. It is possible to control the spatial distribution of the nanoparticles in the liposome by controlling the nanoparticle surface charge (Fig.1). Magnetoliposomes accumulate in the cell cytoplasm reaching 100% labelling efficiency and maintaining cell viability above 80%. Finally, by adjusting the field conditions, drug delivery can be triggered by overcoming the melt transition temperature of the lipid bilayer by remote control using an alternating magnetic field.
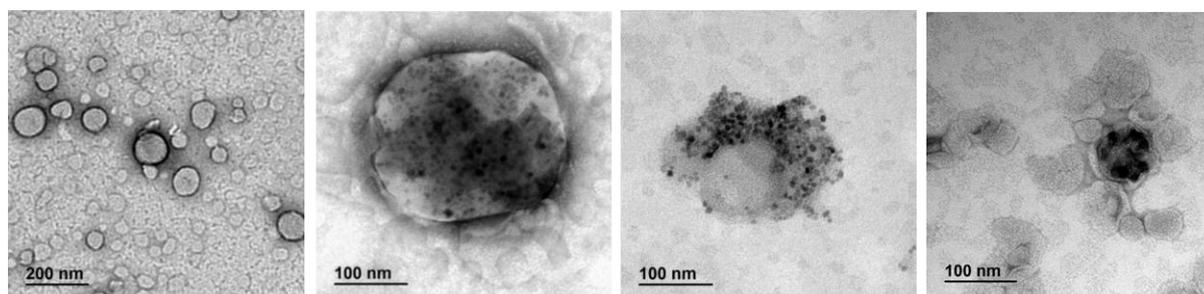

Figure 1. From the left to the right: TEM images of liposomes and magnetoliposomes having magnetic nanoparticles inside the lipid bilayer (coated with oleic acid), attached to the liposome surface (coated with dimercaptosuccinic acid) and encapsulated inside its aqueous volume (coated with aminopropylsilane)

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Monday 20 September 2021

SESSION A2

Aerosol metrology | Advanced X-ray techniques I

Chair: D. Eichert, K. Eleftheriadis

Traceable analysis of aerosols and advanced materials at the nano- and microscales by X-ray spectrometry

B. Beckhoff, K. Bzheumikhova, P. Hönicke, Y. Kayser, C. Stadelhoff, R. Unterumsberger, A. Wählisch, N. Wauschkuhn, J. Weser, and C. Zech

Physikalisch-Technische Bundesanstalt (PTB), Abbestraße 2-12, 10587 Berlin, Germany

E-mail: Burkhard.Beckhoff@PTB.de

The development of new materials and the assessment of aerosols require the correlation of the materials' functionality or particle toxicity with their chemical and physical properties. To probe these properties, analytical methods that are both sensitive and selective at the nano- and microscales are required. The reliability of most analytical methods is based on the availability of reference materials or calibration samples, the spatial elemental composition of which is as similar as possible to the matrix of the specimens of interest. However, there is a drastic lack of certified reference materials in particular at the nanoscale. PTB addresses this challenge by means of a bottom-up X-ray analytical method where all instrumental and experimental parameters are determined with known contributions to the uncertainty of the analytical results. This first-principle based approach does not require any reference materials but a complete characterization of the analytical instruments' characteristics and, in addition, of the X-ray fundamental parameters related to the elements composing the sample. X-ray spectrometric methods allow for the variation of the analytical sensitivity, discrimination capability, and information depth needed to effectively reveal the spatial, elemental, and chemical specimen parameters of interest. Examples of aerosols characterization, interfacial speciation, elemental depth profiling, as well as layer composition and thickness characterizations in advanced materials will be given. Recent instrumental achievements provide access to beam profiles in the nanometer range to qualify heterogeneous materials as well as towards the operando speciation of battery materials. X-ray spectrometry (XRS) under grazing incidence or detection is capable to reveal analytical and dimensional information from layered 2D and 3D systems as well as aerosols or particles deposited on flat substrate surfaces. The SI traceable XRS approach of PTB allows for the qualification of calibration samples for surface contamination as well as for nano- and micro-scaled layers to be employed in laboratory XRF instruments for the establishment of chemical traceability related to quality management purposes.

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Element Mass Concentrations in Ambient Aerosols, a Comparison of Results from Filter Sampling / ICP-MS and Cascade Impactor Sampling / Mobile Total Reflection X-ray Fluorescence Spectroscopy

S. Seeger[1], J. Osan[2], O. Czömpöly[2], A. Gross[3], H. Stosnach[3], L. Stabile[4], M. Ochsenkuehn-Petropoulou[5], L. A. Tsakanika[5], T. Lymperopoulou[5], S. Goddard[6], M. Fiebig[7], F. Gaie-Levrel[8], Y. Kayser[9], B. Beckhoff[9]

[1]Bundesanstalt für Materialforschung und -Prüfung BAM, D-12203 Berlin, Germany
[2]Centre for Energy Research (EK), H-1121 Budapest, Hungary
[3]Bruker Nano GmbH, D-12489 Berlin, Germany
[4]Department of Civil and Mechanical Engineering, University of Cassino and Southern Lazio UNICAS, 03043 Cassino, Italy
[5]Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering, National Technical University Athens NTUA, Iroon Polytechniou 9, 15773 Athens, Greece
[6]National Physical Laboratory, NPL, Teddington TW11 0LW, UK
[7]Norwegian Institute for Air Research NILU, N-2027 Kjeller, Norway
[8]Laboratoire National de Métrologie et d'Essais, 75724 LNE Paris, France
[9]Physikalisch-Technische Bundesanstalt, PTB, D-38116 Braunschweig, Germany

E-mail: stefan.seeger@bam.de

Quantitative chemical analysis of airborne particulate matter (PM) is vital for the understanding of health effects in indoor and outdoor environments and required by EU air quality regulations. Typically, airborne particles are sampled on filters, followed by lab-based analysis, e.g., with inductively coupled plasma mass spectrometry (ICP-MS). Within the EURAMET EMPIR AEROMET project [1,2], cascade impactor aerosol sampling was combined with on-site total reflection X-ray fluorescence (TXRF) spectroscopy. The study aimed at a proof of principles for this new mobile and on-size tool for the quantification of aerosol element compositions and element mass concentrations within short time intervals of less than 12 h. In a field campaign the method's technical feasibility could be demonstrated [3]. The TXRF results were traced back to a stationary, reference-free XRS setup in the laboratory of the German national metrology institute PTB at the BESSY II electron storage ring in Berlin, Germany. Simultaneous PM10-filter sampling, followed by standardized lab-based analysis, allowed for a comparison of the field campaign data of both methods. As Fig. 1 shows, the correspondence between PM10 filter sampling and ICP-MS, and on the other hand, cascade impactor sampling and TXRF is quite encouraging. However, for some of the analysed elements, e.g. V and Pb, the observed deviations are higher than expected and this highlights the fact, that spectral deconvolution strategies for TXRF on cascade impactor samples still need some improvement.
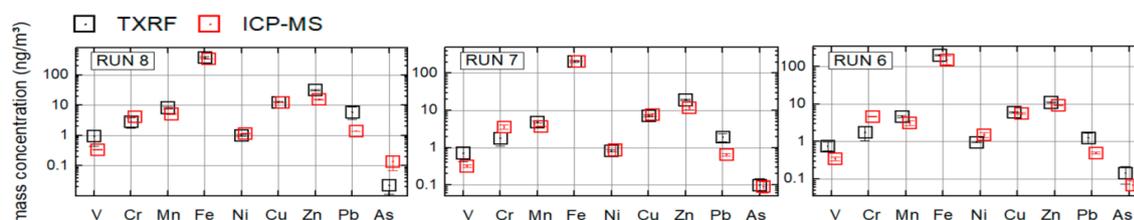

Figure 1. Comparison of element mass concentrations in the PM10 fraction during three runs of a field campaign. Absolute errors determined by respective spectroscopy software.

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X-ray fluorescence and absorption methods for identifying sources of urban aerosol pollution episodes

J. Osán[1], O. Czömpöly[1], V. Groma[1], E. Börcsök[1] and S. Pollastri[2]

[1]Environmental Physics Department, Centre for Energy Research, Konkoly-Thege M. út 29-33., H-1121 Budapest, Hungary

[2]Elettra Sincrotrone Trieste, Strada Statale 14 - km 163,5, I-34149 Basovizza, Trieste, Italy

E-mail: osan.janos@ek-cer.hu

European countries have made progress in reducing particulate air pollution in recent decades being concerned about their heath effect. Especially submicron particles can be detrimental due to their ability to reach alveolar region of the lung. Ultrafine particles (d < 100 nm) can even enter directly to the blood circulatory system. Although the number concentration and size distribution of particulates can be readily investigated by scanning mobility particle sizers, the chemical composition and speciation of particles are of high importance. The size distribution of elemental concentrations has a time variation dependent on sources and meteorological conditions. Both natural and anthropogenic sources can be strongly variable in time. Identifying unique pollution episodes is a difficult task hence the presence of emitted particles could be only a couple of hours.

Size fractionated urban aerosol samples were collected using a May-type cascade impactor onto 7 stages, covering a 70 nm – 9 µm diameter range. Sampling onto silicon wafers enables application of multiple non-destructive analytical methods like total-reflection X-ray fluorescence (TXRF), X-ray absorption near edge structure (XANES) or even scanning electron microscopy (SEM). Using a low-power, laboratory TXRF spectrometer, 0.1 ng/m^3 detection limit could be reached for transition metals at individual impactor stages from 4 m^3 of air sampled [1]. By combination of cascade impactor sampling and TXRF analysis, the particle size dependence of concentrations was obtained for both major (S, K, Ca, Fe) and trace elements (Cu, Zn, Br, Pb). Based on elemental size distribution determined by TXRF pollution episodes were identified. Samples with elevated Cu, Br concentrations were subjected to XANES measurements at the XRF beamline of Elettra (Trieste, Italy). The XANES spectra were analyzed through linear combination fitting of spectra from reference compounds.

With the combination of TXRF and XANES, Cu present in the samples of most episodes studied could be successfully linked to exhaust and non-exhaust type traffic related sources. For a unique short episode with co-existing elevated Cu and Br in the fine fraction, the ratio of organic/inorganic Br species could be determined. Both Cu and Br XANES results underlined that this pollution episode was caused by a local anthropogenic source.

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NPL Studies into dental surgery airborne particulates for COVID-19 mitigation

J.T. Tompkins, D.M. Butterfield, A.S. Brown

Air Quality and Aerosol Metrology Group, National Physical Laboratory, Teddington, TW11 0LW, UK

Email: jordan.tompkins@npl.co.uk

The National Physical Laboratory (NPL) has conducted two studies into the measurement of aerosols produced by dental Aerosol Generating Procedures (AGPs) at a MyDentist dental surgery. The first study was a feasibility study in July 2020 using an Optical Particle Counter (OPC) in a surgery where AGPs were performed on real patients. This attempted to recreate the conditions found in a dental surgery (including both high speed and 3-in-1 AGPs), in line with the UK's Covid-19 guidance at the time. This included the mitigations in use in dental surgeries across the UK: pumped ventilation, an open window, high-bore suction in the mouth and four handed dentistry.
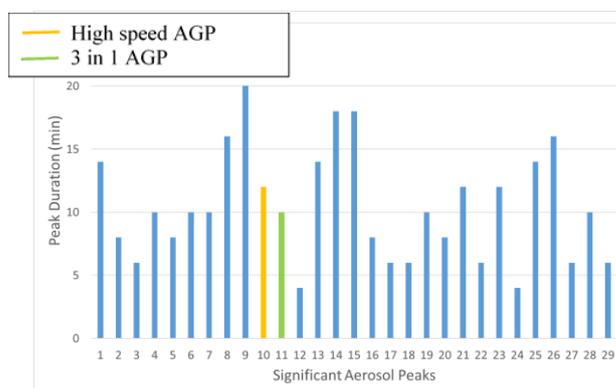
Figure 1. Significant aerosol peaks from AGPs in the first study

The first study suggested that in this specific mitigated clinical environment:
- OPC particle number concentrations in the size range 0.2-10 μm were driven by non-AGP background effects.
- The impact from the range of 41 AGPs on particle number concentrations was limited (Figure 1).
- The duration of significant increased particle number concentration events measured by the OPC did not exceed 20 minutes (Figure 1). The mean event duration time was approximately 10 minutes.
A follow-on study was conducted in February 2021. This study was much larger, with five different OPC instruments in two differently sized rooms with AGPs performed on both dental patients and a surgical manikin. The second study observations supported the first:
- Manikin tests showed a significant increase in OPC-measured particle mass concentrations when the surgery was unventilated (air filtration off, window closed).
- The real patient study found little impact on in-surgery particle concentrations from the range of 19 AGPs where listed mitigations were used. This was true for both particle mass concentrations and particle number concentrations.
- The study highlighted the difference between OPC-measured particle number concentration and particle mass concentration measurements, in relation to aerosol concentrations in the room.
We present the results from both studies and examine the effectiveness of using OPCs in dental studies. We also highlight the importance of the OPC calibration standardisation work in projects such as the AEROMET II EMPIR project.

X-ray Photoelectron Spectroscopy and Scanning Auger Microscopy and Spectroscopy: Theory and Applications

D. Karfaridis, G. Vourlias

AMDe lab, Department of Physics, Aristotle University of Thessaloniki, 54124, Greece
E-mail: dkarfari@physics.auth.gr

X-ray photoelectron spectroscopy (XPS) is an analytical method for material identification, structural and chemical state analysis. The analysis is performed when the sample's surface is irradiated with soft X-rays to ionize atoms and release photoelectrons from the nucleus. The kinetic energy of the escaped photoelectrons is limited by the depth that can be obtained, giving the technique a high surface sensitivity by limiting the sampling depth to a few nanometers (~ 5 nm). Photoelectrons are collected and analyzed by the instrument giving spectra of the emitted intensity as a function of the kinetic or beam energy of the electrons. Since each element has a unique set of binding energies, this technique can be used to determine the quality of elements. Also, the peak regions in the nominal bond energy can quantify the concentration of the respective elements. Small changes in the binding energy (chemical shifts) provide vital data on the chemical state of the sample, enabling chemical analysis of the surface (short-range) without destroying the sample. In addition, Auger electron scanning electron microscopy combines scanning electron microscopy (SEM) and Auger electron spectroscopy (AES) techniques. An electron beam scans the surface, and the collected electrons are energetically analyzed to detect Auger peaks. The intensity of the Auger peaks as a function of the position of the incident electron beam allows us to visualize (spatially) the element from the matching peaks. The depth of information of the technique corresponds to the depth of information offered by Auger electron spectroscopy which typically means the first 2 or 3 atomic layers of the surface.
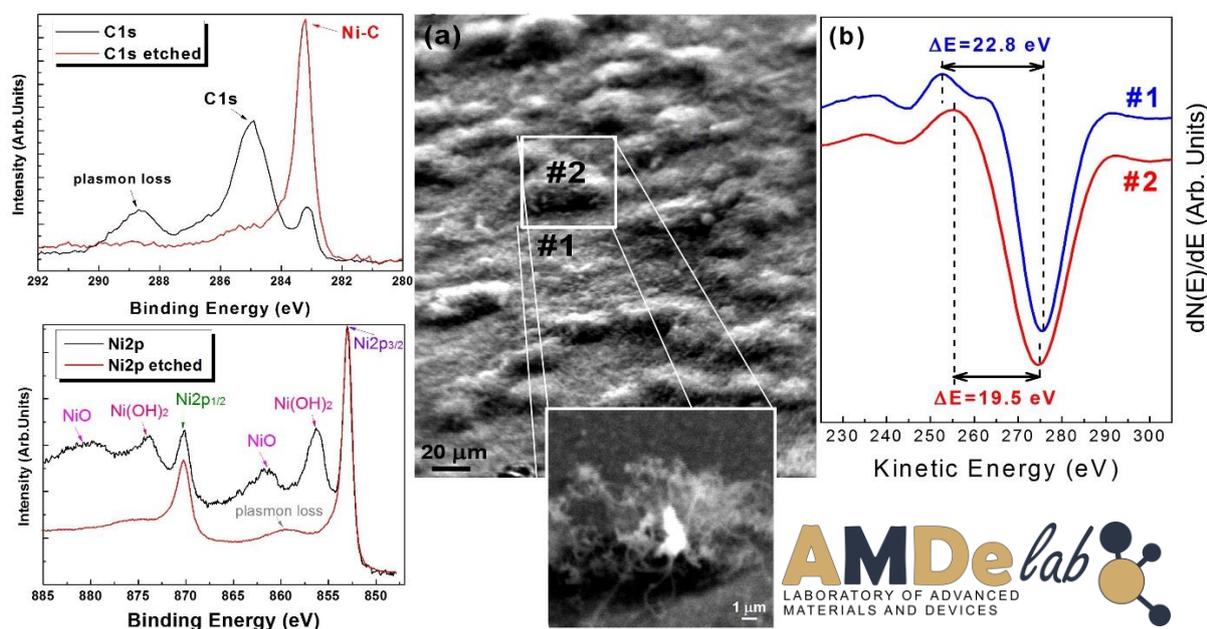
HR XPS measurements of nanocomposite catalysis of DLC: Ni for the growth of CNTs before and after the Ar Ion etching of the surface (at left). (a) FEG AES measurements and quality tests of CNTs on two different spots as shown in the SAM images. Spot #1 indicates an excellent quality of CNTs because of the nano-scale Ni particles as a substrate for the growth of CNTs. Spot #2, on the other hand, shows bad quality CNTs. (b) AES spectra from the two spots.

Technological Advantages in Photoelectron Spectroscopy: Near Ambient Pressure (NAP-XPS) Instrumentation and Applications

L. Socaciu-Siebert, P. Dietrich and A. Thissen

SPECS Surface Nano Analysis GmbH, Voltastr. 5, 13355 Berlin, Germany

E-mail: liana.socaciu-siebert@specs.com

Over the last decades Near Ambient Pressure (NAP) – XPS has demonstrated its promising potential in a wide variety of applications. Starting from operando studies of surface reactions in catalysis, the applications soon have been enhanced towards studies of processes at liquid surfaces. Since more than 15 years, the need for basic studies of the fundamental chemistry at solid-liquid interfaces has attracted growing interest. Dip-and-pull experiments at synchrotrons finally demonstrated, that in-situ and operando XPS in electrochemical experiments can be realized, mainly using synchrotron beams, significantly contributing to the basic understanding of modern energy converting or storing devices, like batteries, fuel cells, etc.

The development of pure laboratory NAP-XPS systems by using small spot and high photon flux density monochromatic X-ray sources together with the development of optimized sample environments, like Peltier coolers and operando liquid cells opens the possibility for the preparation and analysis of a multitude of liquid samples or solid-liquid interfaces on a reliable daily base. Furthermore, the combination of various analysis methods (NAP-SPM, IRRAS, NAP-HAXPES etc.) allows for gaining insights into the fundamental processes that take place during a chemical reaction. Most sophisticated experiments so far have been operando electrochemistry in a classical three-electrode setup.

Existing solutions for NAP-XPS will be presented, with an emphasis on latest developments of instruments and material analysis methods. Examples and results will be shown as well as future perspective of applications and scientific contributions of routine operando NAP-XPS.

Monday 20 September 2021

SESSION A3

Aerosol metrology | Advanced X-ray techniques II

Chair: B. Beckhoff, S. Seeger

The role of aerosol science on understanding and preventing SARS-CoV-2 transmission in the community

K. Eleftheriadis

NCSR "Demokritos", 15310 Ag. Paraskevi, Athens, Greece

E-mail: elefther@ipta.demokritos.gr

The worldwide spread of the SARS-CoV-2 virus has quickly become a pandemic partly due to the fast and unconstrained mechanism or mechanisms of transmission. Infection is usually present in the upper respiratory tract, which is typically the initial site of infection and source of replication for transmission for influenza viruses [1]. Despite the well-known potential for airborne spread of coronavirus disease (COVID-19) the mechanical properties, airborne behaviour, transport and capture as inhaled micro-nano particles by human subjects present in the same area as the host, has not been adequately recognized. However, there is well documented evidence for the significant potential for inhalation exposure to viruses in microscopic respiratory droplets (aerosol) at short to medium distances (up to several meters, or room scale). Before the current pandemic airborne infectious disease transmission was found not only to occur by coughing and sneezing, but also during normal speech, which also yields large quantities of particles that are too small to see by eye, but are large enough to carry a variety of communicable respiratory pathogens [2].

This work will demonstrate examples of the aerosol properties governing residence time and transmission in indoor space and explore the use of preventive measures to mitigate this route of airborne transmission. Studies have demonstrated beyond any reasonable doubt that viruses are released during exhalation, talking, and coughing in aerosol particles small enough to remain aloft in air and pose a risk of exposure at distances beyond 1–2 m from an infected individual

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Total reflection X-ray fluorescence reference materials for cascade impactor air quality monitoring systems

L. Vigna[1], M. Gottschalk[2], E. Cara[3], F. Ferrarese-Lupi[3], A. Verna[1], S. L. Marasso[1,4], S. Seeger[2], L. Boarino[3], N. De Leo[3], F. C. Pirri[1,5] and M. Cocuzza[1,4]

[1]Politecnico di Torino, Torino, 10129, Italy
[2]Bundesanstalt für Materialforschung und -prüfung (BAM), Berlin, 12489, Germany
[3]Istituto Nazionale di Ricerca Metrologica (INRiM), Torino, 10135, Italy
[4]CNR-IMEM, Parma, 43124, Italy
[5]Istituto Italiano di Tecnologia, Center for Sustainable Future Technologies, Torino, 10144, Italy

E-mail: lorenzo.vigna@polito.it

The development of traceable methodologies to measure particulate matter concentration and monitor air quality in polluted areas of Europe is an important challenge to be addressed in order to reduce health and environmental backlashes. The use of cascade impactor sampling methods coupled with total reflection X-ray fluorescence (TXRF) spectroscopy for elemental mass concentration measurements is an accurate way with fast response and low level of uncertainty [1]. Certified reference materials are being developed to mimic the cascade impactors patterns with known quantities and distribution of target elements.

To this goal, a new method has been developed to obtain flexible, reusable and low-cost parylene C shadow masks by using photolithographic steps. After the fabrication, the obtained micro stencils are applied to 30 mm acrylic substrates in order to evaporate different metals and thus replicating the Dekati patterns.

Another type of reference samples can be obtained by directing the self-assembly of block copolymers (BCPs) nanotemplates inside pre-lithographed Dekati patterns and by infiltrating the BCPs with metallic oxide. The absolute quantification of infiltrated materials by means of reference-free GIXRF allows tuning the process for the required metallic mass deposition.
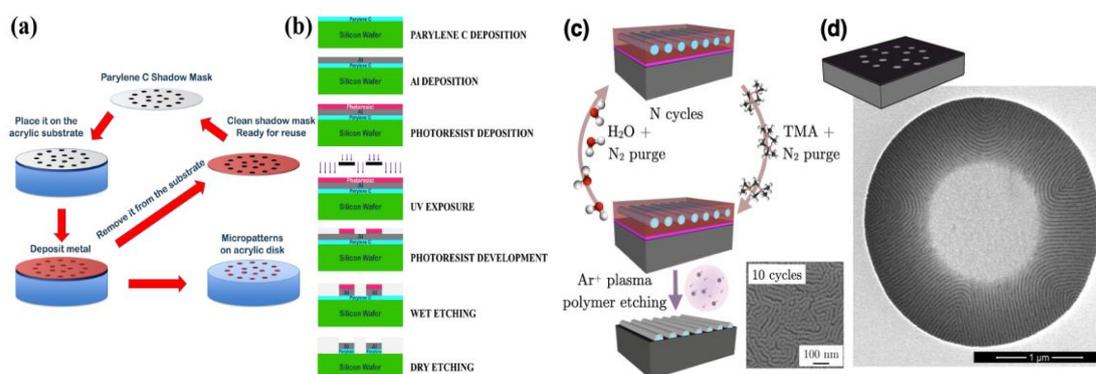

Figure 1. (a) Sequence of steps for micropatterning using reusable parylene C shadow masks and (b) sketch of the fabrication steps used to obtain the parylene C stencils. (c) Scheme of the sequential infiltration synthesis of BCPs cylindrical nanotemplates and SEM image of the resulting metallic oxide replica. (d) Microtemplate on silicon used to direct BCPs' self-assembly in the Dekati pattern.

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Experimental measurement protocol of aerosol deposition relating to a bend in a large ventilation network with rectangular ducts

D. Costa[1, 2], J. Malet[1] and E. Géhin[2]

[1]Institut de Radioprotection et de Sûreté Nucléaire (IRSN), PSN-RES, SCA, LEMAC, Gif-sur-Yvette, 91192, France
[2]Univ Paris Est Creteil, CERTES, F-94000 Creteil, France

E-mail: delphine.costa-upec@irsn.fr

Aerosol deposition in ventilation plays a key role in particulate pollution transfers inside buildings, but has been mostly studied in ventilation systems by applying models developed on the basis of experimental results obtained mainly on aerosol sampling lines. The size of these lines differs greatly with the size of industrial ventilation ducts, whose cross-sections are larger than 50 centimeters. Besides, in the literature, few experimental studies focused on singularities, like bends, mostly at a medium scale with circular or square cross-sections. By applying these models, aerosol deposition measurements are affected by a scale effect and possibly by airflow patterns specific to rectangular ducts due to secondary flows.

In this study, we focus on aerosol deposition relating to a bend, in rectangular ducts (600 x 400 mm²) of the DIESE research facility, a new ventilation network of industrial size built at IRSN. We present the corresponding experimental protocol and aerosol deposition velocity measurements.

Experimentally, we need to generate enough aerosols to compensate the dilution due to high airflows. Furthermore, monodisperse aerosol generators are generally lab-scale devices, leading to detection challenges: we need to come to a compromise and find a measurement technique with a low enough limit of detection. Consequently, we used a fluorescent aerosol and verified a) the aerosol size distribution and the injected mass flow; b) the repeatability of the volume concentration in the bulk, sampled on filters and c) the repeatability of aerosol deposition measurements, carried out by wiping wall surfaces and completed by fluorescence spectroscopy analyses. Additionally, we carried out aerodynamic measurements, such as velocity measurements with hot wire anemometry and PIV.

Inter-laboratory comparison of ED-XRF/PIXE analytical techniques in the elemental analysis of filter-deposited multi-elemental certified reference materials representative of ambient particulate matter

M. Gini[1], M.I. Manousakas[1,2,12], V. Kantarelou[2], A.G. Karydas[2], M. Chiari[3], A. Migliori[4], N. Civici[5], B. Veleva[6], K. Šega[7], L. Samek[8], C. Samara[9], Z. Kertesz[10], J. Osan[11] and K. Eleftheriadis[1]

[1]ERL, Institute of Nuclear & Radiological Sciences & Technology, Energy & Safety, N.C.S.R. "Demokritos", 15310 Athens, Greece
[2]Institute of Nuclear and Particle Physics, N.C.S.R. "Demokritos", 15310 Athens, Greece
[3]National Institute of Nuclear Physics (INFN), 50019 Sesto Fiorentino (FI), Italy
[4]Nuclear Science and Instrumentation Laboratory, Physics Section, International Atomic Energyy Agency, 2444 Seibersdorf, Austria
[5]Institute of Applied Nuclear Physics, University of Tirana, Tirana, Albania.
[6]National Institute of Meteorology and Hydrology, 1784 Sofia, Bulgaria
[7]Institute for Medical Research and Occupational Health, Ksaverska cesta 2, p.p. 291, 10001 Zagreb, Croatia
[8]AGH University of Science and Technology, Faculty of Physics and Applied Computer Science, Krakow, Poland
[9]Department of Chemistry, Environmental Pollution Control Laboratory, Aristotle University of Thessaloniki, University Campus, 54124 Thessaloniki, Greece
[10]Institute for Nuclear Research (Atomki), H-4026 Debrecen, Bem tér 18/c, Hungary
[11]Centre for Energy Research, Konkoly-Thege Miklós út 29-33., H-1121 Budapest, Hungary
[12]Laboratory of Atmospheric Chemistry, Paul Scherrer Institute (PSI), Villigen, Switzerland

E-mail: gini@ipta.demokritos.gr

The quantification of the elemental concentration of ambient particulate matter is a challenging task because the observed elemental loadings are not well above the detection limit for most analytical techniques. Although non-destructive nuclear techniques are widely used for the chemical characterization of ambient aerosol, only one multi-element standard reference filter material that mimics ambient aerosol composition has become recently available in the market. To ensure accuracy, reliability and comparability of instruments performance, multiple reference materials with different elemental mass loadings are necessary. In this study, an intercomparison exercise was performed to evaluate the measurement uncertainty and instruments performance using multi-element dust standard reference samples deposited on PTFE filters. The filter samples, produced by means of dust dispersion, were tested in terms of homogeneity, reproducibility and long-term stability. Eight laboratories participated in the exercise. The evaluation of the results reported by the participants was performed by using two sets of reference values: a) the concentrations reported by the Expert Laboratory, b) the robust average concentrations reported by all participants. Most of the reported on the certificate of analysis elements were efficiently detected in the sample loadings prepared as representative for atmospheric samples by the Expert Laboratory. The average absolute relative difference between the reported and the reference values ranged between 0.1 % (Ti) and 33.7 % (Cr) (CRM-2584). The participants efficiently detected most of the elements except from the elements with atomic number lower than 16 (i.e. P, Al, Mg). The average absolute percentage difference between the participants results and the assigned value as derived by the expert laboratory was 17.5 ± 18.1 % (CRM-2583; Cr, Pb excluded) and 16.7 ± 16.7 % (CRM-2584; Cr, P excluded).

Evaluation of Bone and Joint Quality in Mice Employing X-ray Microtomography

P. Giannoutsou[1], A. Kaspiris[2], P. Kastana[2], C. Kontoyannis[1,3], E. Papadimitriou[2] and M. Orkoula[1]

[1]Laboratory of Instrumental Analysis, Department of Pharmacy, University of Patras, Greece
[2]Laboratory of Molecular Pharmacology, Department of Pharmacy, University of Patras, Greece
[3]Institute of Chemical Engineering Sciences, FORTH, Greece
E-mail: malbie@upatras.gr

The quality of femur bone, lumbar vertebra and hip joint of mice, with aging, was evaluated in the present study. Male and female mice, of a wide range of ages (7 to 80 weeks), strain Black6, were scanned using X-ray microtomography. Parameters such as mineral density, were calculated to assess bone quality.

The parameter values as well as the images recorded reveal that bony and cartilage tissues remain in healthy condition until the age of 1 year. From this point on, decay takes place in bone (osteoporosis) and joints (osteoarthritis). Mineral density of femur and vertebra trabecular bone decreases, while the free space between bone rods increases. The spongy architecture is maintained with wider gaps. Trabecular bone becomes less.

The cartilage covering the femoral head in hip joint mineralizes. It converts to a thick bony layer, losing the elastic properties of the chondral tissue. Osteoarthritis is installed. On the other hand, subchondral bone remains dense, until the age of 80 weeks.
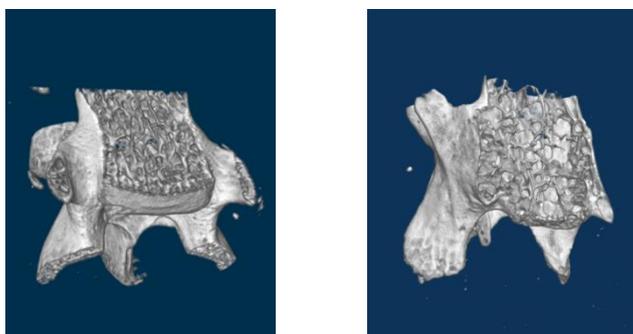
Picture 1. X-ray microtomographic section of a 7 weeks (left) and 80 weeks (right) lumbar vertebra. Trabecular bone is declining.
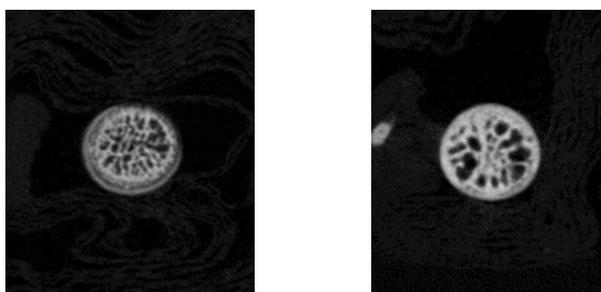
Picture 2. Left: 7 weeks (healthy) femoral head section. Outer cartilagenous layer is evident. Right: 80 weeks (osteoarthritic) femoral head section. Cartilage has completely mineralized.

Aethalometer multiple scattering correction in an Arctic environment

S. Vratolis, K. Eleftheriadis

ERL, Institute of Nuclear & Radiological Sciences & Technology, Energy & Safety, National Centre for Scientific Research "Demokritos", 15310 Ag. Paraskevi, Attiki, Greece.

E-mail: vratolis@ipta.demokritos.gr

In this work, measurements of aerosol optical properties at Zeppelin station, near Alesund, Svalbard are reported. The station is located at 78.907 N^o and 11.889 E^o, at 474 m above sea level (a.s.l) in the untouched Arctic environment, far away from substantial contamination sources and is therefore an ideal place for monitoring global atmospheric gasses and long-transported contaminants. Spectral absorption properties of atmospheric aerosol are continuously monitored at the station since 1998. Here we report absorption data from AE-31 and MAAP for the period between July 2015 and December 2017. Observations of the aerosol light absorption at 637 nm from the Multiangle Absorption Photometer (MAAP, Model 5012 Thermo Scientific) were used as reference. The instrument is described in detail in [1].
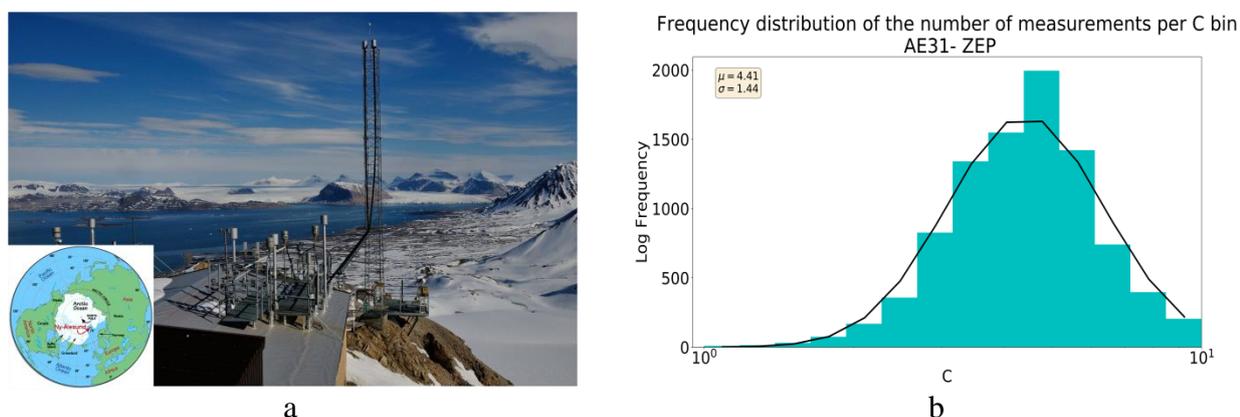
Figure 1. a: Zeppelin Station, b: Lognormal frequency distribution of the multiple scattering correction factor for the AE31.

The multiple scattering correction factor C for the Aethalometer is derived from the equation:

$$C = b_{ATN} (637nm)/(b_{absref} (637nm) * R@637nm)$$

b_{ATN} is the aerosol attenuation coefficient measured by the Aethalometer, b_{absref} is the aerosol absorption coefficient measured by the MAAP and R is the filter loading effect correction factor [2]. In Figure 1b we present the histogram of C values for the AE31 aethalometer and its lognormal fit.

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IMA-2021

Monday 20 September 2021

SESSION A4

Spectroscopy and applications

Chair: B. Michalke, E. Chatzitheodoridis

Timelines in Atomic Spectrometry

G.M. Hieftje

Indiana University, 980 Cape Marco Drive, #1205, Marco Island, FL 34145, USA

E-mail: hieftje@indiana.edu

Compared to fields such as art, philosophy and mathematics, atomic spectrometry is a very young discipline. The foundations of atomic spectrophysics and atomic spectrochemistry can be traced back to such scientific giants as Newton, Bunsen, and Kirchhoff. However, it can be argued that many aspects of current practice in *analytical* atomic spectrometry were introduced in the 19th, 20th and 21st centuries. Such developments include modern ways of producing and detecting atomic optical and mass spectra, the introduction of alternative sources for generating neutral atoms and atomic ions, advanced understanding of fundamental events that affect analytical figures of merit, an almost continuous improvement of such figures of merit, the melding of atomic and molecular spectrometry, and the application of atomic spectrometric methods to an almost unfathomable range of contemporary problems. However, even the most up-to-date methods, applications, and instrumentation for atomic spectrometry exhibit shortcomings. In this presentation, some of these limitations will be outlined and possible means to overcome them offered. These considerations will in turn suggest future directions that research in atomic spectrometry might profitably take.

Analytical and structural perspectives of NMR in natural products research: From mixture analysis to in-cell investigations

I.P. Gerothanassis

Section of Organic Chemistry and Biochemistry, Department of Chemistry, University of Ioannina, Ioannina, Greece

E-mail: igeroth@uoi.gr

A critical overview of recent analytical developments of NMR spectroscopy in natural products will be provided with emphasis in the following applications:

(i) chemical analysis of extracts without isolation or derivatization steps [1], 'in situ' direct monitoring of dynamic changes of metabolites [2], investigation of enzymatic reaction products [3], real time biotransformation monitoring [4], and metabolomics analysis [5].

(ii) "in-cell" NMR studies of the apoptotic activity of flavonoids [6] and polyunsaturated fatty acids with the Bcl-2 family of proteins and

(iii) quantum chemical calculations of high-resolution structures of analytes in solution based on NMR chemical shifts [7].

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Optimising the Workflow for Microplastics Analysis using FTIR Microscopy

I. Robertson

Perkin Elmer Limited, Seer Green, Bucks, UK

E-mail: Ian.robertson@perkinelmer.com

Analysis of environmental samples containing microplastics is essential to determine their prevalence and their impact. A range of analytical techniques have been applied to the analysis of microplastics. Of the techniques adopted, infrared (IR) spectroscopy, and more specifically IR microscopy, has established itself as a primary analytical technique for the detection and identification of microplastics.

The microplastics analysis workflow for IR microscopy consists of several steps involved in getting from the raw sample to answers, including the initial sampling through to data analysis. The steps involved may be different depending on the type of initial sample and the amount of sample cleanup required to prepare the sample for infrared (IR) analysis. This paper describes the different types of environmental samples, the sample collection methods, the range of different sample cleanup methods, and then deals more specifically with the best ways to optimise sample filtration for measurement by IR microscopy.

The principles of IR microscopy and the different sample measurement modes will be described, comparing and contrasting each type. IR microscopy and imaging experiments can generate significant quantities of data that need to be analysed to get the required information. The different methods for extracting data and information will be explained and suggestions made for best practice.

# Introduction to FTIR techniques for materials characterization

L. Malletzidou, T. Zorba, G. Vourlias

Advanced Materials and Devices Laboratory, School of Physics, Aristotle University of Thessaloniki, GR-54124, Thessaloniki, Greece

E-mail: labrinim@auth.gr

Fourier transform infrared spectroscopy (FTIR) has been established as one of the fundamental techniques for materials molecular characterization [1]. Using infrared irradiation -from Near IR up to Far IR-, information can be acquired from both organic and inorganic matter, with applications varying from determination and observation during synthesis, up to recording of ageing processes.

Among its well-known techniques of examination (transmittance for powders or thin films, and specular reflectance for materials with polished surface), attenuated total reflection (ATR) technique is ideal for examination of liquid, strongly absorbing and thick samples, while sample preparation is unnecessary and the specimens can be retrieved after examination.

The combination of an FTIR spectrometer with an infrared microscope (micro-FTIR) -equipped with single element MCT detectors- has the advantage that analysis can be performed in the micrometric scale, for both transmittance and reflectance modes, even with the use of a micro-ATR accessory. Over the last years, micro-FTIR spectroscopy has been equipped with focal plane array (FPA) detectors, evolving infrared microanalysis even more [2]. FPA detectors, consisting of thousands of individual detectors, provide simultaneously thousands of individual spectra -each one for every individual detector-, which leads to FTIR imaging with spatial and spectral dimensions, in contrast to FTIR mapping using single element detectors, in terms of measurement time and spatial resolution [3]. Thus, FTIR imaging emerges in applications where chemical imaging is needed, such as clinical and biomedical applications, polymers and microplastics, industry and more [2–5].
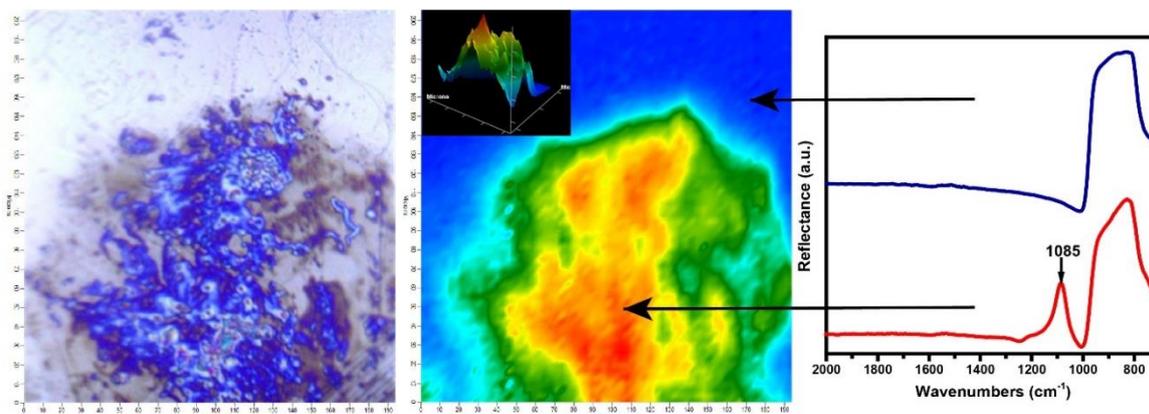
Microscopic image of a SiC surface including unknown superficial spots (left). FPA chemical images using integrated area under Si-O band (centre). Corresponding FTIR spectra of the SiC surface and the SiO_2 spots, the latter as an oxidation product (right).

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IMA-2021

Further developments in battery-operated microplasmas

L. Quadeer[1], D. Cebula[1], V. Karanassios[1,2]

[1]Department of Chemistry, University of Waterloo, Waterloo, Ontario, Canada
[2]Waterloo Institute of Nanotechnology, University of Waterloo, Waterloo, Ontario, Canada

E-mail: vkaranassios@uwaterloo.ca

Plasmas, and in particular, Inductively Coupled Plasmas (ICPs) are widely used in elemental analysis. In an ICP, the plasma is formed on top of a torch. In an ICP, its torch consumes a typical of about 20 L/min of Argon gas and requires about 1.25 kW of electrical power. We have been developing and characterizing miniaturized version of plasmas, called microplasmas. In contrast to laboratory-scale ICPs, our microplasmas can be operated from a battery (requiring about 10 W of electrical power) and a flow-rate of about 250 mL/min of Argon gas, thus they be used *on-site* (i.e., in the field). They can also be fabricated using a variety of inexpensive technologies (e.g., 3D printing) and materials (e.g., flexible polymeric substrates). An example of a microplasma is shown in the figure below [1]. A microplasma has been arbitrarily defined as one with one critical dimension (e.g., length, or width, or height, or radius), around 1000 μm or less [2, 3].
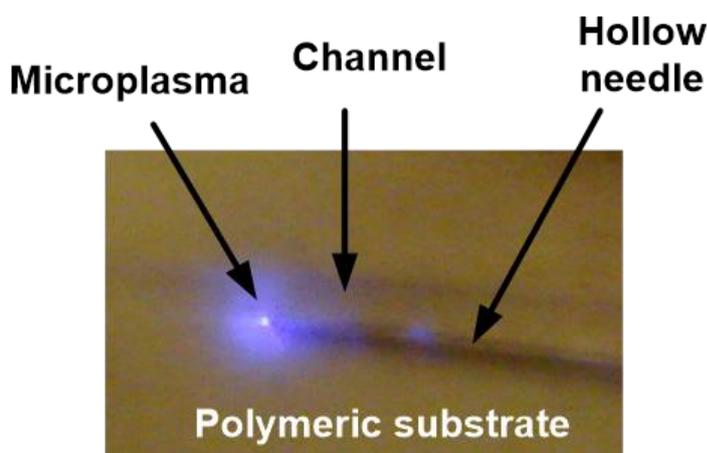

Figure 1. The microplasma (above) is formed at the end of a needle (with an ID=110 □m). The needle is offset from the microchannel for illustration purposes. The channel is covered using a flexible, polymeric substrate (e.g., 0.3 mm poly carbonate plate).

In this presentation, recent developments in portable, battery-operated microplasma spectrometry will be described in detail.

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ATR – FTIR as a Tool for Osteoarthritis Disease Diagnosis

<u>N. Pradakis</u>[1,2], N. Vordos[1,2], D.A. Gkika[3,4], M. Koffa[5], K. Tilkeridis[6]

[1]Physics Department, International Hellenic University, 65404, Kavala, Greece
[2]Biophysics Laboratory, Physics Department, International Hellenic University, 65404, Kavala, Greece
[3]Chemistry Department, International Hellenic University, 65404, Kavala, Greece
[4]Hephaestus Laboratory, International Hellenic University, 65404, Kavala, Greece
[5]Department of Molecular Biology and Genetics, Democritus University of Thrace, Alexandroupolis, Greece
[6]Orthopaedic Department, Medical School, University General Hospital of Alexandroupolis, Democritus University of Thrace, St. Niarhos 1, Dragana, 68100 Alexandroupolis, Greece

Email: npradaki@physics.auth.gr

Osteoarthritis (OA) constitutes one of the major factors contributing to a low-quality life among elders. It is the main cause of disability and limited mobility, especially for people aged 65 and older and can be described as an abnormal musculoskeletal condition which affects more than 300 million people on a global scale. Pain is the basic symptom resulting in a low-quality life. Currently, diagnosis is feasible through different means, such as radiographic analysis, ultrasonography, conventional examination, as well as lab analysis without, however, a straight-forward-quantitative global scale categorization based on the degree of damage. ATR-FTIR is an excellent method for analyzing biological samples such as tissues, cells or fluids. Even though this method has been previously applied for a variety of biological materials, there are only few papers employing this method for the analysis of synovial fluid (SF) to diagnose OA. The purpose of this work is to present a sufficient protocol about the analysis of synovial fluids aiming to identify potential biomarkers of OA. To do so, a review was performed to compile sample collection information and present the spectrum analysis protocols. The research period lasted between the 24th of March and the 12th of June 2021. A bibliographic search was performed on the following databases: Scopus, PubMed and Google Scholar, using the keywords "FTIR", "ATR", "synovial", "fluid" "osteoarthritis", "analysis" (and combinations thereof such as "synovial fluid" AND "analysis" AND "FTIR"). Different databases were used because they have different search algorithms and hence different results are presented. From the total of 729 papers checked, only a total of 35 were relevant to this work, out of which many were counted multiple times. The analysis of the relevant literature showed significant differences in both the temperature and the storage time of the samples. Immediate consequence is the large variation in the peaks of the spectra, as well as the conclusions drawn. The best protocol, according to the results of this work, corresponds to a 4 °C storage temperature for a 24 hours period before the analysis after joint aspiration. The temperature and storage were selected in terms of SF component quality preservation. In the future, more than 100 samples of synovial fluid will be analyzed, to investigate the existence of possible biomarkers.

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Tuesday 21 September 2021

SESSION B1

Plenary Lectures

Trends in analytical techniques

Chair: N. Thomaidis, F. Tsopelas

Instrumental analysis for metals in biology: ultratraces, isotopes, species and images

R. Lobinski

[1]Institute of Analytical and Physical Chemistry for the Environment and Materials, (IPREM-UMR 5254), CNRS, E2S UPPA, 64000 Pau, France

[2]Chair of Analytical Chemistry, Warsaw University of Technology, 00-664 Warszawa, Poland

E-mail: ryszard.lobinski@univ-pau.fr

The concentrations of the individual metals in biological systems are tightly controlled by the physiology and biochemistry of an organism. Biologically mediated processes lead to the isotopic fractionation and sub-cellular distribution. The metal-induced gene expression results in the synthesis of a myriad of biological ligands on the proteome and metabolome levels implied in the uptake, trafficking, accumulation and excretion of trace elements. There is increasing evidence that biochemical and physiological processes can contribute to the fractionation of mineral elements on the isotopic level, as it has been well established for oxygen, nitrogen or sulfur. Consequently, information on the metal concentrations, speciation, isotopic composition and distribution is essential for the understanding of the role and fate of metals in a biological system.

The status of a mineral element in a biological sample can be defined by: (i) concentration (possibly isotopic composition), (ii) chemical speciation and (iii) localization. The control of all the three is essential for the cellular metal homeostasis. Therefore, there is a need for analytical methods able to provide the necessary information for all the elements, for the maximum of species and with a maximum spatial resolution at an adequate time scale.

The lecture discusses the state of the art of the development of such a versatile "toolbox", its recent advances and evolution trends. Particular attention will be given to global approaches aimed at a simultaneous comprehensive speciation analysis of one or several elements (metallomics [1]). Recent trends in ultratrace analysis by ICP MS, precise isotope ratio measurements by multicollector ICP MS, high resolution (Fourier Transform) electrospray mass spectrometry, and nanoSIMS imaging will be highlighted.

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Reviewing determination methods for "Free Cu" and Fe(II)/Fe(III) ratio and development of a new and fast speciation techniques for reliable measurements of "free Cu" and Fe(II)Fe(III) ratio in various samples

B. Michalke

Helmholtz Center Munich – German research Center for Environmental Health, Research Unit
Analytical BioGeoChemistry
Ingolstädter Landstr. 1, 85764 Neuherberg, Germany

E-mail: bernhard.michalke@helmholtz-muenchen.de

Trace elements (or trace metals) are minerals present in living tissues in small amounts. Some of them are known to be essential, others may be essential aside from the remaining elements, which are considered to be nonessential. Trace elements function primarily as catalysts in enzyme systems; some metallic ions, such as iron and copper, participate in oxidation-reduction reactions in energy metabolism. Iron, as a constituent of hemoglobin and myoglobin, also plays a vital role in the transport of oxygen.

However, since copper and iron can show different oxidation states – i.e. they can be redox-active – both elements are probably causative in pathologic disease mechanisms of an extended set of severe illnesses. Such diseases include (for iron) forms of cancers, like prostate cancer and melanoma, as well as ferroptosis, which is a newly discovered peculiar form of non-apoptotic cell death and (for copper) Menkes and Wilson´s disease. Menkes Disease is a congenital metabolic disorder based on a copper metabolic disfunction. Wilson's disease is a rare inherited disorder that causes copper to accumulate in your liver, brain and other vital organs. For understanding pathogenesis mechanism and diagnosis purposes it is of vital importance to know the quantitative level either of free copper or quantitatively the Fe(II) and Fe(III) concentrations to quantify the cellular Fe(II)Fe(III) ratio.

Several methods have been published providing different sensitivity, selectivity and time-spending/saving procedures, for both, free copper determination and iron redox state.

This contribution summarizes shortly those methods, spots light on strength and limitations of each method.

For Cu, the „difference" method is the common clinical method for determining extractable Cu levels in serum, but also an IP-SCX-ICP-sf-MS method is reviewed, which is providing differentiation of Cu(I) and Cu(II), but also (Fe(II) vs. Fe(III). Molecular sieves and measurement of filtrate by ICP-MS or a newly published simple LP-SCX-ICP-MS method will be shortly evaluated. Finally, this lecture introduces a new SCX-ICP-MS or ICP-OES based determination method for free Cu determination, avoiding identified limitations from previous methods. This new approach allows also parallel monitoring of sulfur-compounds (mainly proteins), by that way proofing the free Cu signal being not co-eluting (not bound to) with S-ligands (proteins).

For iron redox speciation semi-quantitative dyes only for Fe(II) are in use. The Fe(II)/(III) ratio thus cannot be estimated from that approach. For overcoming this lack, we developed an innovative CE-ICP-MS technique that allows the simultaneous quantification of redox-active Fe(II) and redox-inactive Fe(III) ions, aside from total iron.. Via this method, we uncovered that amyloid precursor protein suppression results in a robust accumulation of Fe(II)-driven lipid peroxidation, sensitizing cancer cells to cell death by ferroptosis. This method was further modified to quantitatively distinguish ferritin-bound iron from other iron species, enabling to calculate the Fe-load of ferritin in the respective sample. Sequentially, in a second analytical run the cellular GSH:GSSG ratio in the same cell lysate is assessed.

As a proof of principle, the application to several different cell lysate types is shown as well as the respective changes in Fe(II)/(III) ratio due to cellular knock-downs or iron regulators.

Temperature Gradient Gas Chromatography (TGGC) – Pushing Separation Speed and Separation Efficiency at the Same Time

B. Klampfl, R.D. Müller, <u>E. Rosenberg</u>

Vienna University of Technology, Institute of Chemical Technologies and Analytics, Getreidemarkt 9/164
A-1060 Vienna, Austria

E-mail: erosen@mail.zserv.tuwien.ac.at

The "magic triangle" is well known in chromatography, representing the dilemma that only two of the three parameters separation speed, efficiency and sample capacity, can be optimized at the same time, however, at the expense of the third parameter [1].

Although the development of new stationary phase materials in HPLC (such as monolithic columns and columns with core-shell materials) seems to have alleviated the consequences of this dogma, it still holds true for all forms of chromatography in use nowadays.

In gas chromatography isothermal operation will produce the best separation for a complex mixture, however, at the expense of excessively long separation times and consequently loss of sensitivity. For this reason, temperature programmed separations are typically used, as these allow to optimize the capacity factor and hence retention and to improve thus resolution even for a sample of constituents differing widely in their volatility and thus elution temperature.

Normal temperature-programmed separation is characterized by a temperature increase which is changing with time (from low to higher temperatures), but is not changing in space, i.e., along the length of the separation column.

In contrast, systems for gas chromatographic separation have been introduced by Boeker et al. [2] and also earlier by Contreras et al. [3] which make use of a negative thermal gradient. This gradient is stationary in time, however non-stationary in space which means that the temperature decreases along the column from the hot (injector) end to the cold (detector) end. This setup shall be termed temperature gradient gas chromatography (TGGC). While this setup appears counter-intuitive, it leads to a peak focusing as the analytes are moving into a colder part of the column in which their travelling velocity is reduced. This results in very narrow peaks – hence a good chromatographic resolution –, and as very short columns (ca. 2 m) are typically used, also in very fast separations.

Since its beginning, there has been an intensive discussion whether or not TGGC is capable of improving resolution beyond what is achievable in isothermal GC [4,5]. We will present and explain the opposed viewpoints, and arrive at a conclusion which considers the arguments from both sides. Moreover, we will present different options to experimentally realize TGGC from completed and ongoing work in our laboratory [6,7].

Acknowledgment

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Quo Vadis Electroanalytical Chemistry

J. Barek

Charles University, Faculty of Science, Department of Analytical Chemistry, UNESCO Laboratory of Environmental Electrochemistry, Albertov 6, CZ 128 43 Prague 2, Czech Republic

E-mail: barek@natur.cuni.cz

In spite of well-known advantages of modern electroanalytical methods (low cost, easy automation, portability resulting in easy on site measurements and compatibility with "green analytical chemistry"), these methods are not so frequently used in practical laboratories as they would deserve. In our UNESCIO laboratory of environmental electrochemistry we are trying to improve the situation by developing new sensors/biosensors applicable for the determination of biologically active compounds (drugs, plant hormones, pesticides, biomarkers, etc.). Our effort will be illustrated by newly developed electroanalytical methods based on i) new electrode materials, e.g. boron doped diamond [1], silver solid amalgam [2], carbon film [3], ii) electrodes modified with molecularly imprinted polymers [4], iii) combinations of electroanalytical and separation methods [5,6,7], iv) measurements in flowing systems [8].

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Tuesday 21 September 2021

SESSION B2

Chromatography | Mass spectrometry

Chair: E. Rosenberg, D. Lambropoulou

Challenges in cannabis testing

J. P. Hopkins[2], F. Aimone[1], A. Dalmia[3], E. Cudjoe[4], T. Astill[4], J. Jalali[5], J.P. Weisenseel[3], F. Qin[4], M. Murphy[6], T. Ruthenberg[7]

[1] Perkin Elmer Viale dell'Innovazione, 3, 20125 Milano MI, Italy
[2] Malva S.A. 13 Ilission street, N. Kifissia 14564, Greece
[3] PerkinElmer, Inc., Shelton, CT
[4] PerkinElmer, Inc., Woodbridge, ON, Canada
[5] PerkinElmer, Inc., San Jose, CA
[6] SC Labs, Tigard, OR
[7] SC Labs, Santa Cruz, CA

E-mail: John.Hopkins@Malva-SA.com

Cannabis (*Cannabis* Sativa) is a plant of great commercial interest, but its complex resinous parts cause the analysis significant technical challenges. The plant has been known for its medicinal properties since ancient times[1], but owing to its psychotropic effects, it has been banned in most countries around the World to control its potential as a drug of abuse. Subsequently, several States have changed their laws allowing the medical usage of cannabis and/or cannabinoids for conditions such as Lennox-Gastaut syndrome and Dravet syndrome[2], which cause severe and frequent seizures which are often not manageable with traditional anti-epileptic medications, but also for chronic pain[3], terminal cancer[4] and multiple sclerosis[5].

National regulatory drug agencies have not yet agreed the testing required for cannabis, but generally, testing controls gross contamination, pesticides, mycotoxins, microbial contamination, residual solvents as well as cannabinoid contents and ratios. Often, recreational cannabis is also tested for its terpene profile which can reflect a user's subjective enjoyment, but these terpene fractions are of interest to medical researchers as well.

The analysis of cannabis uses chromatographic techniques to separate isobaric components prior to analysis by mass spectrometry. Cannabis matrix is hugely complex – much more so than fruit and vegetables, for example, which also require similar testing for consumer safety. A particular challenge is the rapid contamination and resulting decline in sensitivity of mass spectrometers.

We describe a mass spectrometer which does not suffer from sensitivity declines, as it does not use any voltages in the source and ion-transfer regions, instead using a novel laminar gas flow, combined with heated surfaces in the source and ion transfer regions to prevent contamination and resultant sensitivity loss. We present real-World data of the applicability of this technology in the analysis of very crude cannabis extracts.

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Green Microwave-Assisted Cannabis Extraction:
from Lab to Industry

<u>V.L. Tsamili</u>, A. Berberidis, A. Kaloyannis

ALS Analytical Laboratory Systems SA, Thessalias 6 str., Alimos 17456, Attica, Greece

E-mail: lina.tsamili@alssa.gr, info@alssa.gr

Cannabaceae plants *Cannabis sativa L.* (Cannabis), are rich in terpenes. Some terpenes of cannabis are relatively well known for their potential in biomedicine and have been used in traditional medicine for centuries, while others are yet to be studied in detail. Terpenes' medicinal properties are supported by numerous in vitro, animal and clinical trials and show anti-inflammatory, antioxidant, analgesic, anticonvulsive, antidepressant, anxiolytic, anticancer, antitumor, neuroprotective, anti-mutagenic, anti-allergic, antibiotic and anti-diabetic attributes, among others. Because of the very low toxicity, these terpenes have been proven safe and well-tolerated and are already widely used as food additives and in cosmetic products. [1]

Technical advances over the years have allowed the development of green extraction techniques, such as Microwave-Assisted Extraction (MAE). Microwaves (MW), a mature technique for extraction-process intensification, is attracting great amounts of attention in academic-research and industrial-application fields for its technological advantages, while global market needs suitable technologies for the smart recovery of bioactive Cannabis metabolites [2]. Well-established and innovative extraction protocols in addition to chromatographic separations, are reviewed highlighting their respective advantages, as Gas Chromatography (GC) techniques are also reported for accurate identification and quantification of terpenes in complex cannabis matrices [3].

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Overcoming analytical challenges in trace metal analysis by ICP-MS: A multi quadrupole perspective

H. Ernstberger[1], E. Kroukamp[2]

[1]PerkinElmer, Viale dell'Innovazione 3, 20126 Milano, Italy
[2]PerkinElmer, 501 Rowntree Dairy Rd, Woodbridge, ON L4L 8H1, Canada

E-mail: helmut.ernstberger@perkinelmer.com

The need for effective interference removal has been a major driver for innovation in quadrupole ICP-MS over the last 25 years or so. Various implementations of collision and reaction technology were augmented with triple quad instrumentation (ICP-MS/MS) for this purpose. Most recently, a multi-quadrupole instrument was made available [1]. This presentation reviews the distinguishing features of multi-quad analysis vs. triple quad analysis, supported by performance data. It is concluded that the use of multiple quadrupoles in ICP-MS allows the highest level of control of interferences, with ensuing benefits for accuracy, background equivalent concentrations, and detection limits. Supplementary features attributable to the multiple quadrupole technology, such as the possibility to extend the dynamic range in this instrumental implementation will also be discussed.

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Tuesday 21 September 2021

SESSION B3

Chromatography | Mass spectrometry II

Chair: R. Lobinski, A. Hiskia

Photocatalysis of diclofenac using metal-organic framework-based oxidant systems: kinetics, transformation products elucidation, and *in silico* toxicity

A. Kontogiannis[1], E. Evgenidou[1,2], C. Nannou[1,2], D. Bikiaris[3] and D. Lambropoulou[1,2]

[1]Laboratory of Environmental Pollution Control, Department of Chemistry, Aristotle University of Thessaloniki, GR-541 24 Thessaloniki, Greece
[2]Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Thessaloniki GR-57001, Greece
[3]Laboratory of Polymer Chemistry and Technology, Department of Chemistry, Aristotle University of Thessaloniki, Thessaloniki, GR-541 24, Greece

E-mail: dlambro@chem.auth.gr

In the present study, the photocatalytic degradation of Diclofenac under solar simulated light (SSL) using Basolite F300®, a MOF (Metal-organic framework) material, as catalyst was investigated. The process was enhanced by the addition of oxidants [1], [2] (H_2O_2 or $K_2S_2O_8$) since Fenton-like reactions appear to take place [3]. The identification of transformation products (TPs) was performed on experiments conducted on the optimum – as derived from the kinetics experiments- parameters affecting the photocatalytic procedure (i.e. type of oxidant, oxidant concentration) using LC–Orbitrap MS/MS instrumentation. According to the identified TPs, parent compound transformation followed several transformation routes including hydroxylation, decarboxylation, ring formation, combined with subsequent oxidation and/or ring opening. Lastly, predictions of acute and chronic toxicity on each of the detected TPs on fish, daphnid and green algae were performed *in silico* using ECOSAR v2.0 with a view to risk assessment of the photocatalytic process and interpretation of the *in vivo* toxicity results on daphnia magna. Although many of the identified TPs appear to be more toxic than the parent compound the overall toxicity of the solution is significantly reduced at the end of the process.
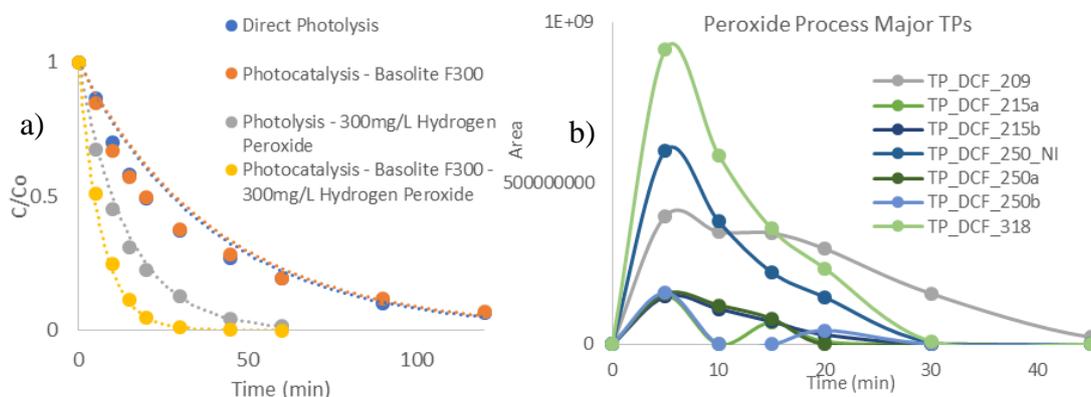

Figure 1. Kinetic profiles of (a) diclofenac under different photocatalytic systems, (b) formed TPs under SSL- MOF-H_2O_2 system

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Stop Searching - Start Looking. A breakthrough in chromatography technologies

R. Rozenich

Waters Corporation, Austria

E-mail: m.kontogiannis@hellamco.gr

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Development and validation of a novel analytical approach for the simultaneous speciation analysis of mercury and selenium in fish by HPLC-ICP-MS following enzymatic extraction

M. Ribeiro[1,2,3] , A. Leufroy[1], T. Guérin[4], I. Castanheira[2], J. Silva[3], P. Jitaru[1]

[1] ANSES, Laboratory for Food Safety, F-94700 Maisons-Alfort, France

[2] Departamento de Alimentação e Nutrição, Instituto Nacional de Saúde Doutor Ricardo Jorge, INSA IP, Avenida Padre Cruz, 1649-016, Lisboa, Portugal

[3] Centro de Química Estrutural, Instituto Superior Técnico, Universidade de Lisboa, Av. Rovisco Pais, 1049-001, Lisboa, Portugal

[4] ANSES, Directorate of Strategy and Programs, F-94700 Maisons-Alfort, France

E-mail: petru.jitaru@anses.fr

Mercury (Hg), via its organometallic species, methylmercury (MeHg) is one of the most toxic pollutants on a global scale. MeHg bioconcentrates in the marine environment and hence its presence in human food and particularly in fishery products is a matter of concern as seafood consumption is the major route of human exposure to MeHg. Several authors have hypothesized that selenium (Se) species may reduce the bioaccumulation of MeHg in fish and/or increase its elimination from fish or human body but data are very scarce to fully confirm this hypothesis. It appears also that the ability of Se to provide protective effect against Hg toxicity is species dependent. The poor knowledge of MeHg behaviour in fish & fishery products in relation with Se exposure is explained partially by the challenges related to the analytical chemistry, especially the lack of truly accurate method capable to perform simultaneous speciation analysis of several Hg and Se species in complex biological matrices such as the fish.

This study aims at the development and validation of a new analytical method for the simultaneous speciation analysis of Se and Hg by HPLC–ICP-MS. For this purpose, four Se species such as Se(IV), Se(VI), selenomethionine (SeMet) and selenocysteine (SeCys) and two Hg species, namly Hg^{2+} and MeHg were separated by an anion-cation exchange HPLC hyphenated to inductively coupled plasma-mass spectrometry (ICP-MS). The species were extracted simultaneously by enzymatic hydrolysis in a closed microwave system.

The method was validated by means of the accuracy profile approach by carrying out 3 measurement series in duplicate on different days over a timespan of 3 weeks.

These analytical performances allow the application of the method to simultaneous speciation analysis of Se and Hg in fish within the framework of a project aiming to better understand the antagonistic relationship of these elements.

IMA-2021

The coupling of CE and ICP-MS/MS as a novel tool for investigation of the gold nanoparticle-cisplatin based drug targeted delivery systems formation

A. Wróblewska[1], A. Milewska[1], M. Drozd[2], M. Matczuk[1]

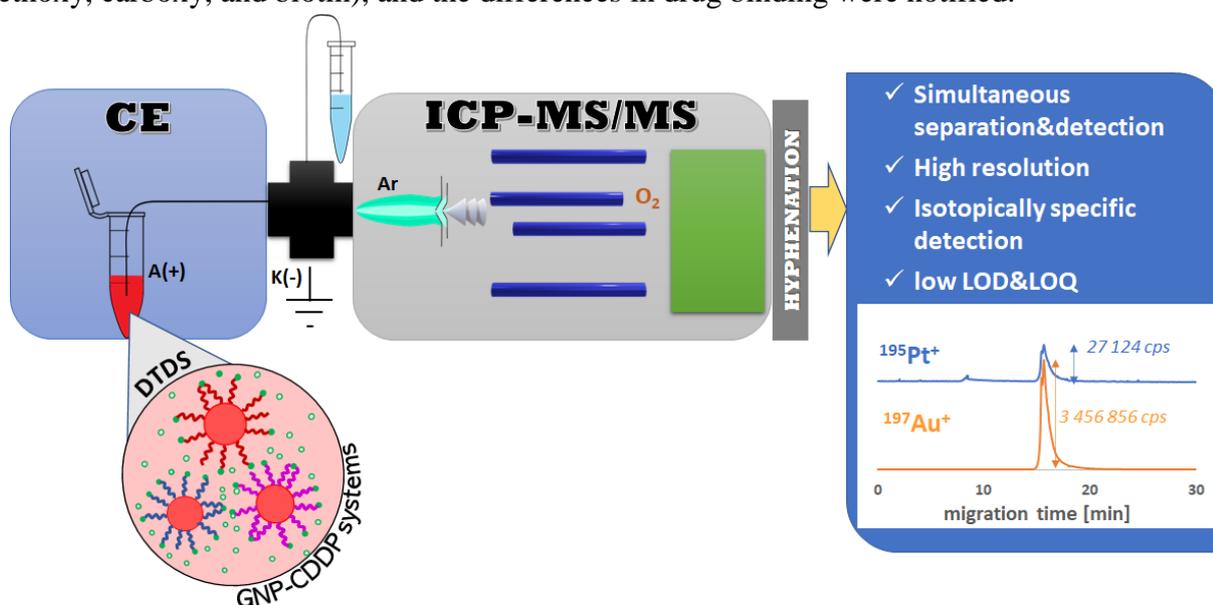[1]Chair of Analytical Chemistry
[2]Chair of Medical Biotechnology
Faculty of Chemistry
Warsaw University of Technology
Noakowskiego St. 3, 00-664 Warsaw, Poland

E-mail: awroblewska@ch.pw.edu.pl

Despite the wide application in classical chemotherapy, cisplatin (CDDP) does not exhibit selectivity towards cancerous cells, contributing to severe side effects and intrinsic or acquired resistance of various cancer types. The gold nanoparticle-based (GNPs) drug targeted delivery systems (DTDS) were proposed to overcome the mentioned issues. In trials of such systems (composed from GNPs and CDDP), the optimization of efficient synthetic procedure to obtain the stable in (simulated) human serum conditions conjugates between system constituents is a crucial aspect. In these DTDS, the drug form and the type of nanocarrier surface modification play a significant role. Moreover, it is essential to select analytical methods dedicated to the quantitative analysis of DTDS formation and stability profile. Most often are applied the microscopic and spectrophotometric techniques, which do not guarantee the DTDS determination in the reaction mixture. A promising solution is using the hyphenated techniques composed by the coupling of the separation and detection modules, enabling the simultaneous analysis and accurate identification of sample constituents on the trace level. Herein, we present the research concerning the employment of CE-ICP-MS/MS (capillary electrophoresis - inductively coupled plasma tandem mass spectrometry) to examine the influence of GNPs surface modification type on the CDDP binding effectiveness and stability of formed DTDS. During investigations, the three types of GNPs were tested (GNPs with various functionalization groups: methoxy, carboxy, and biotin), and the differences in drug binding were notified.

Novel CE-ICP-MS/MS-based methodology for the study of Fe_3O_4 nanoparticle's interactions with human serum proteins under physiological environment

<u>J. Sikorski</u>, J. Kruszewska, A. Kamińska, M. Jarosz, M. Matczuk

Chair of Analytical Chemistry, Faculty of Chemistry, Warsaw University of Technology, 00-664 Warsaw, Poland

E-mail: jsikorski@ch.pw.edu.pl

Due to unique magnetic properties, low cytotoxicity, stability, biocompatibility, and biodegradability, superparamagnetic iron oxide nanoparticles (SPIONs) have been extensively researched for medical applications. In particular, their potential use in cancer diagnosis and therapy, e.g., as drug delivery agents, is of great interest. However, this requires the development of appropriate methods to study the behavior of nanoparticles under physiological conditions.

In this work, we present a novel capillary electrophoresis-inductively coupled plasma tandem mass spectrometry (CE-ICP-MS/MS) method to determine SPIONs that provide monitoring of their interactions with human serum proteins in the simulated physiological environment [1]. The separation of differently charged SPIONs was achieved employing capillary electrophoresis, while the use of tandem ICP-MS minimized polyatomic interferences and allowed the quantification of nanoparticles in a wide range of concentrations. The selection of oxygen as collision gas brought better results for simultaneous detection of sulfur (a marker of proteins) and iron (a marker of SPIONs), while the mass-shift approach was chosen to monitor the analytes due to the higher background noted for them using on-mass mode. In the separation method, the temperature of tray and cassette and pH of background electrolyte were adjusted to imitate physiological conditions. The rest of the parameters (type of background electrolyte, applied voltage, volume of sample) were optimized to obtain the best quality of signals and the recovery of analytes from the capillary.

The optimized method was used to investigate the SPIONs–albumin interactions after 30 min of incubation. The obtained results confirmed the formation of nanoparticle−proteins conjugates. Summarizing the presented methodology could be used in the future for the investigation of the iron oxide nanoparticles changes in biological samples.

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QSRR driven insight into retention in multimodal chromatography

B. Otašević, B. Svrkota, J. Krmar, A. Protić, M. Zečević

University of Belgrade – Faculty of pharmacy, Vojvode Stepe 450, 11221 Belgrade, Serbia

E-mail: biljana.otasevic@pharmacy.bg.ac.rs

Liquid chromatography system in which several separation mechanisms are integrated in the composition of a single column is called multimodal or mixed-mode chromatography (MMC). MMC systems are classified based on combined separation mechanisms as bimodal (RP/HILIC, RP/IEX, HILIC/IEX) and trimodal (different combinations of RP/HILIC/IEX). The main benefit of MMC lies in widening the spectra of properties of analytes that can be simultaneously chromatographed (nonpolar, polar, organic, inorganic, ionized and / or non-ionized analytes). In this way, it is possible to reduce the number of required analyses for one complex sample compared to unimodal chromatographic systems. For that reason, the popularity of MMC has been growing fast in recent years. However, in line with this achievement, MMC is characterized by large number of intermolecular interactions governing separations which are related to the properties of the analyte (charge and polarity) and chromatographic conditions (ionic strength and the pH of the aqueous phase and the content of the organic solvent) [1]. In order to get insight into relative contribution of aforementioned factors to retention of selected group of analytes, preferred chemometric approach is Quantitative Structure Retention Relationship (QSRR) study. The QSRR models relate the physical-chemical properties of analytes reflected by assigned molecular descriptors with their retention behaviour in predefined experimental space described by the range of chromatographic conditions (instrumental and mobile phase composition related factors). Apart from its general purpose to assist in the characterization of observed chromatographic system, the reliable predictions of retention behaviour of so-called system blind analytes (analytes of known chemical structure but not subjected to experimentations) can also be derived from a QSRR model. In such way, the development of MMC based analytical method can be rationalized by saving time and other resources [2].

This research demonstrates the QSRR study performed on 31 pharmaceuticals covering wide range of polarities, acid-base properties and divergent retention in RP/WCX system (Thermo Acclaim Mixed Mode WCX-1 3 μm, 2.1x150 mm column). This system was subjected to variations of the mobile phase composition (30-50% (v/v) of acetonitrile; 3.8-5.6 pH value and 20-40 mM ionic strength of acetic buffer) and column temperature (30–38 °C) according to the plan of central composite design of experiments. The machine learning algorithm based on Artificial Neural Network was used for relating these independent variables to cube root transformed retention factors of analytes as observed responses. The network comprising of 11-7-1 topology was trained through 1200 cycles with learning rate set at 0.3 and momentum set at 0.5. Cross validation and external validation were used to prove good statistical performances of built model (Root Mean Square Error values 0.131 and 0.147 and Squared Correlation vales 0.963 and 0.944, respectively). According to the weighting scheme used, volume fraction of acetonitrile, pH of aqueous phase and descriptors related to hydrophobicity and molecule size demonstrated the greatest impact towards retention in MMC.

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Identification of transformation products derived from pharmaceutically active compounds in surface waters by liquid chromatography–high-resolution mass spectrometry

K. Anagnostopoulou[1,2], C. Nannou[1,2], E. Evgenidou[1,2], D. Lambropoulou[1,2]

[1]Laboratory of Environmental Pollution Control, Department of Chemistry, Aristotle University of Thessaloniki, Thessaloniki, GR-541 24, Greece
[2]Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center, Thessaloniki, 57001, Greece

E-mail: dlambro@chem.auth.gr

Surface water quality depends on a variety of parameters, including emerging contaminants (ECs). The interest in ECs has focused on pharmaceutically active compounds (PhACs) among others, since they continuously end up to the environment through human excretion, direct disposal of expired/ unused formulations in sewers or household waste etc. Therefore, PhACs result in the discharge of WWTPs, which are the main route for their entry into the water ecosystems. Once entered aquatic environment, pharmaceuticals are subjected to biotic (aerobic/anaerobic degradation by microorganisms) and abiotic processes (hydrolysis, photolysis, etc.). Hence, the study of their fate by investigating their transformation products (TPs) is necessary to evaluate their potential on aquatic ecosystems. High-resolution mass spectrometry (HRMS) coupled with liquid chromatography (LC) is a promising trend in environmental analysis, since "suspect" compounds such as the expected TPs can be identified, despite not only the limited information but also the unavailability reference standards. HRMS has been proved promising for untargeted or suspect profiling of TPs in environmental matrices since it greatly expands the ability to capture the broad spectrum of potential TPs. The collected data from suspect and non-target screening of TPs can be used either prospectively or retrospectively.

The aim of the present study was to identify TPs of PhACs in surface waters, by means of LC–HRMS, by the cutting-edge Orbitrap analyzer, after applying a generic sample preparation protocol with solid phase extraction (SPE). Initially, a dynamic list of suspect TPs was constructed, based (a) on the occurrence frequency of the parent drugs in surface waters, (b) the prescription/consumption rates of the drugs, and (c) the available bibliographic data on the occurrence of transformation products of PhACs. The list comprised the expected molecular formula, the exact mass of the pseudomolecular ion (calculated automatically), the ESI mode (positive/ negative) and the expected fragment ions, when available. The criteria for the identification of suspect compounds were (a) peak intensity ($<10^4$), (b) isotopic profile (70%), (c) S/N>10, (d) presence of at least one fragment. The processing of the results revealed more than 70 TPs of PhACs. Notably, in some cases the relative ion abundance corresponding to the TPs was greater than the parent compound, underlining the importance to study the TPs along with the parent compounds. To expand the suspect screening to a wide-scope non-target analysis, the raw data were also processed with the Compound Discoverer 3.2 software, and all the detected TPs were also identified. Finally, an *in-silico* approach based on the ECOSAR was exploited to the assess the ecotoxicity induced by TPs.

Tuesday 21 September 2021

SESSION B4

Chromatography | Speciation | Bioanalytics | Bioseparation

Chair: K. Valko, I. Panderi

On-line Hyphenation of CE with MC-ICP-MS for Species-specific Isotopic Analysis of Sulfur

S. Faßbender[1], K. Rodiouchkina[2], F. Vanhaecke[2], B. Meermann[1]

[1]Federal Institute for Materials Research and Testing (BAM), Division 1.1 – Inorganic Trace Analysis
Richard-Willstätter-Straße 11, 12489 Berlin - Germany
[2]Atomic & Mass Spectrometry – A&MS research unit Department of Chemistry, Ghent University, Campus Sterre, Krijgslaan 281 – S12, 9000 Ghent, Belgium

E-mail: bjoern.meermann@bam.de

In many scientific fields, isotopic analysis can offer valuable information, e.g., for tracing the origin of food products, environmental contaminants, forensic and archaeological samples (provenance determination), for age determination of minerals (geochronological dating) or for elucidating chemical processes. Up to date, typically bulk analysis is aimed at measuring the isotopic composition of the entire elemental content of the sample. However, the analyte element is usually present under the form of different species.[1] Thus, separating species of interest from one another and from matrix components prior to isotope ratio measurements can provide species-specific isotopic information,[2,3] which could be used for tracing the origin of environmental pollutants and elucidation of (environmental) speciation. Using on-line hyphenations of separation techniques with multicollector-ICP-MS (MC-ICP-MS) can save time and effort and enables the analysis of different species during a single measurement.

In this work, we developed an on-line hyphenation of CE with multicollector-ICP-MS (CE/MC-ICP-MS) for isotopic analysis of sulfur species using a multiple-injection approach for instrumental mass bias correction by standard-sample bracketing. With this method, the isotopic composition of sulfur in sulfate originating from river water could be analyzed without sample preparation.[4] The results were compared to data from off-line analysis of the same samples to ensure accuracy. The precision of the results of the on-line measurements was promising regarding the differentiation of the river systems by the isotopic signature of river water sulfate. The great potential of this method is based on the versatility of the applied separation technique, not only in the environmental field but also for, e.g., biomolecules, as sulfur is the only covalently bound constituent of proteins that can be analyzed by MC-ICP-MS.

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Challenges and innovations for trace level quantification of bioactive compounds in biological matrices using tandem mass spectrometry

<u>L. Barreiros</u>[1,2], S. R. Fernandes[1,2], S. Machado[1], E. M. P. Silva[1,3] and M. A. Segundo[1]

[1]LAQV, REQUIMTE, Department of Chemical Sciences, Faculty of Pharmacy, University of Porto, Porto, Portugal
[2]School of Health, Polytechnic Institute of Porto, Porto, Portugal
[3]CESPU, Institute of Research and Advanced Training in Health Sciences and Technologies, Gandra, Portugal

E-mail: lbarreiros@ff.up.pt

The development of analytical methods amenable to deal with real-life samples and to produce accurate results within a short time is a current demand of the modern world. During the last years, tandem mass spectrometry (MS/MS) based techniques have become the method of choice for the targeted analysis of trace levels of chemical compounds in complex matrices due to their inherent high sensitivity and selectivity. Most strategies rely on the coupling to a separative technique prior to MS/MS detection. In this work, the association HPLC-MS/MS for quantification of bioactive compounds in biological matrices will be critically discussed. The steps of sample preparation and analytical determination will be addressed. Moreover, the main analytical features of each developed method, including selectivity, accuracy, precision, limits of detection (LOD) and quantification (LOQ), stability, recovery and matrix effects will be highlighted.

The implementation of a methodology based on UHPLC-MS/MS for monitoring tranexamic acid (TXA) in human plasma samples at sub-microgram per milliliter levels was pursued [1]. Despite the recognition of TXA as an important antifibrinolytic agent, there is lack of pharmacokinetic and pharmacodynamic data, claiming for the development of accurate and sensitive quantification methods. In a different context, considering the continuous search for new treatment strategies for human immunodeficiency virus (HIV) infection, an HPLC-MS/MS method was developed for the simultaneous quantification of the antiretrovirals tenofovir and efavirenz. The target samples were mice tissue and fluid samples recovered from a pharmacokinetics study with nanoparticles [2]. Finally, BIBP 3226 is a selective neuropeptide Y Y1 receptor antagonist that has proved *in vitro* a positive impact in bone turnover, thus providing good perspectives towards its application as a pharmacological tool for bone regeneration. Having in mind the need to elucidate receptor-antagonist internalization mechanisms, the challenge was to develop a methodology based on HPLC-MS/MS that permitted to determine the low quantities of antagonist expected to be internalized by cells [3].

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Prediction of acute aquatic ecotoxicity of emerging contaminants using biomimetic chromatographic measurements

<u>C. Stergiopoulos</u>[1], F. Tsopelas[1], K. Valko[2], M. Ochsenkühn-Petropoulou[1]

[1]Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering, National Technical University of Athens, Greece

[2]Bio-Mimetic Chromatography Ltd., Stevenage, Herts, United Kingdom

E-mail: chrisxp3@hotmail.com

The presence of emerging contaminants in aquatic environments can have adverse effects on ecosystems and long-term consequences on human health [1]. Therefore, hazard assessment of compounds seems inevitable in order to evaluate their ecological risk, although no specific regulations seem to exist until nowadays [2]. In this work, the potential of biomimetic chromatography to predict ecotoxicological endpoints of organic compounds of emerging concern was investigated. For this purpose, retention factors of 36 structurally diverse drugs and 39 structurally diverse pesticides were measured on immobilized artificial membrane (IAM), immobilized human serum albumin (HSA) and immobilized alpha-1-acid-glycoprotein (AGP) stationary phases [3]. As ecotoxicological endpoints, half-maximal lethal concentration (LC_{50}) values of fish (rainbow trout, fathead minnow, bluegill sunfish and sheepshead minnow species) and half-maximal effective concentration (EC_{50} - immobilization) values of water flea (daphnia magna spp.) were considered [4]. Data was compiled from the US Environmental Protection Agency (EPA). For reasons of comparison, corresponding models were derived using calculated physicochemical properties. Models derived with biomimetic properties gave better statistics and superior predictive performance from those based on calculated physicochemical properties. Overall, biomimetic chromatography can be suggested as a promising method in order to screen or rank chemicals in regard to their ecotoxicological risk.

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Multiplex lateral flow assay for liquid biopsy applications

P. Kalligosfyri[1], S. Nikou[2], S. Karteri[3], H. Kalofonos[3], V. Bravou[2] and <u>D. Kalogianni</u>[1]

[1] Department of Chemistry, University of Patras, 26504 Rio, Patras, Greece
[2] Department of Medicine, University of Patras, 26504 Rio, Patras, Greece
[3] Division of Oncology, Department of Internal Medicine, University Hospital of Patras, 26504, Rio, Greece

E-mail: kalogian@upatras.gr

Liquid biopsy is a non-invasive alternative approach than that of conventional tissue biopsy for cancer diagnosis and real-time monitoring of the disease. Liquid biopsies target specific biomarkers, such as blood circulating tumour cells (CTC), circulating tumour DNA (ctDNA), circulating microRNAs and exosomes that are secreted by the tumours. Body fluid samples, such as blood sample, are used in liquid biopsies. The presence of circulating tumour DNA (ctDNA) or cell-free DNA (cfDNA) in blood circulation has been related to early cancer detection and disease monitoring for proper treatment. The detection and quantification of cfDNA is a great analytical challenge due to its low molecular weight, its extremely low abundance in blood circulation and the need for the discrimination of ctDNA that carries specific tumour-related gene mutations from normal cfDNA. So, the discrimination of ctDNA from normal cfDNA is based on the detection of specific gene mutations that are located only at the tumour DNA. In this project, a new rapid multiplex lateral flow assay was developed for the simultaneous detection of three major single-point mutations of the KRAS gene, related to colorectal cancer, in ctDNA, as well as the normal cfDNA. The multiplex test was initially optimized with synthetic DNA targets and cell lines that express both the normal and the mutated gene. The test was then successfully applied to tissue samples and blood serum samples for cfDNA detection. The proposed lateral flow assay offers a simple, cost-effective and rapid alternative of liquid biopsy that is completed within 15 min, while the detection is accomplished by bare eye. This lateral flow assay offers multiplex detection of single point mutations, while it is universal and can be applied in the future for the detection and identification of several different kinds of cell-free or circulating tumour DNA molecules.

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Study of scandium extraction from Greek bauxite residue by *Acetobacter Tropicalis*

K. Kiskira[1, *], Th. Lymperopoulou[2], L.A. Tsakanika[1], Ch. Pavlopoulos[3], K. Papadopoulou[3], K.M. Ochsenkühn[1], G. Lyberatos[3], M. Ochsenkühn-Petropoulou[1]

School of Chemical Engineering, National Technical University of Athens, Iroon Polytechniou 9, Zografou campus, 15773 Athens, Greece; [1] Laboratory of Inorganic and Analytical Chemistry;[2]Products and Operations Quality Control Laboratory ;[3]Laboratory of Organic Chemical Technology

E-mail: kirki.kis@gmail.com

Bauxite residue (BR) is an alkaline by-product generated from the Bayer process for alumina production. It is considered a hazardous material due to its volume (120 million tons of production annually) and alkalinity. The disposal of BR constitutes a complex global environmental problem [1]. BR can be used as a secondary source for valuable metals such as rare earth elements (REEs). Greek BR contains a high scandium (Sc) content of around 0.1 kg/ton that represents 90–92% of the economic value for all REEs [2]. Sc extraction from BR has been studied mainly by hydrometallurgical methods using different mineral acids such as nitric, hydrochloric, sulfuric acid under different leaching conditions [2-4]. Biotechnologies such as bioleaching can play an important role in metal recovery and they are considered as 'a green technology' with operational flexibility and low energy requirements [5,6]. The aim of this study was the investigation of the development of a bioleaching procedure using a pure culture of *Acetobacter tropicalis*, an acetic acid bacterium that produces high amount of organic acids, for Sc recovery from Greek BR and the optimization of the process by testing different parameters such as BR (S/L) ratio, biomass production, final pH, percentage recovery of Sc and the organic acids produced by the microorganism. A chemical analysis for Sc and Fe of the leachate solutions was conducted using ICP-OES, additionally, for Fe, AAS was used. The identification of organic acids was performed with HPLC. The number of bacterial cells was counted by a standard plate count (SPC) method. XRD analysis of BR before and after bioleaching was performed. The micro-morphology of the samples was also examined by SEM equipped with EDAX for semi-quantitative chemical analysis. The maximum extraction of Sc was 42% in a one-step bioleaching process at 1% S/L in 20 d. The main organic acids produced were acetic, oxalic and citric. The results indicate a synergistic effect of the different organic acids produced by the bacterium. Optimisation of the bioleaching process is in progress, using different microbial cultures such as *Aspergillus niger* and *Penicillium oxalicum* aiming to minimise the incubation time and maximise Sc recovery for up-scaling bioleaching for the Sc extraction from BR.

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Accumulation and biological role of zolendronate in rabbit intervertebral discs

A. Kouroumalis, <u>C. Christophoridis</u>, D. Kletsas

Laboratory of Cell Proliferation and Ageing, Institute of Biosciences and Applications, NCSR "Demokritos", Athens, Greece *e-mail: dkletsas@biodemokritos.gr

Low back pain is considered one of the major chronic disorders, caused mainly due to the degeneration of the intervertebral disc (IVD). IVDs lie between the vertebral bodies constituting the joints of the spine, and they consist of two distinct regions, the outer annulus fibrosus (AF) and an inner gelatinous core (the nucleus pulposus, NP). Biphosphonates, used for a long time in osteoporosis management, have been reported to be beneficial in the homeostasis of tissues resembling IVDs, such as articular cartilage. However, the effect of bisphosphonates, particularly zolendronate (ZOL), on the homeostasis of IVDs at the cellular and tissue level has not been addressed until now, as far as we know. The determination of ZOL in complex biological matrices presents several analytical challenges, thus, an efficient sample treatment procedure, in combination with effective chromatographic separation and highly sensitive detection are necessary.

The scope of this work was to develop and validate a SPE-liquid chromatography -tandem mass spectrometry (SPE-LC-MS/MS) analytical method, using on-cartridge derivatization, in order to evaluate the presence of ZOL in various tissues (rabbit blood plasma, skin and intervertebral disc, including AF and NP), and to study its effect on IVD cells.

Samples were acquired as follows: Plasma was collected from centrifuged rabbit blood. AF, NP and skin tissue dissected from rabbits. The tissues were chopped, enzymatically digested and then processed for ZOL characterization. For the determination of ZOL in blood plasma and tissues, on-cartridge SPE (Hypersep SAX) derivatization with trimethylsilyl diazomethane (TMS-DAM) was carried out. Ibandronate was used as an internal standard. The final extract was analyzed by LC-MS/MS. Method specificity, linearity, range of measurement, trueness, precision and limits of detection/quantification were assessed for validation purposes. The cytotoxic effect of ZOL on primary rabbit AF IVD was estimated by MTT and Neutral Red assay and intracellular ROS production with DCFHDA assay. Finally, extracellular matrix genes' expression was estimated with quantitative RT-PCR.

The analytical method presented excellent linearity ($R^2 > 0.990$), high recoveries (75-87%), increased repeatability (0.5-5.6%) and low LOD values (15pg/mg in the case of skin and IVDs) in all matrices studied. Plasma, skin and AF samples, presented a similar temporal variation pattern in the concentration of ZOL, with a peak at 30-60 min and then was quickly reduced approaching baseline levels. ZOL was not detected in NP tissues. Physiological concentrations of ZOL were non-cytotoxic and unable to increase ROS production in rabbit AF cells. Finally, ZOL provoked an anabolic phenotype in AF IVD cells.

Our data indicate that ZOL accumulates in the AF region of the IVD, and not to the core of the tissue, i.e. NP which is characterized by an absolute absence of vascularization. The presence of ZOL seems to have a positive (anabolic) effect on this tissue.

Machine leaning techniques for the correlation of retention factors of biomimmetic chromatography with pharmacokinetic properties

P. Danias, F. Tsopelas, A. Pappa

Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering, National Technical University of Athens, Iroon Polytechniou 9, 157 80 Athens, Greece

E-mail: pan.danias@hotmail.com

Biomimetic chromatography, involving Immobilized Artificial Membrane (IAM) and Immobilized Artificial Protein Chromatography as well as micellar chromatography, has gained considerable attention due to its ability to simulate biological processes, in particular for rapid screening of candidate drugs in the early stages of drug discovery. Biomimetic chromatographic indices can be used for the prediction of pharmacokinetic properties, such as volume of distribution, plasma protein binding (%PPB), Human Intestinal Absorption (HIA) and Blood Brain Barrier (BBB) Penetration [1]. In some cases, however, standard computational techniques like linear and non-linear regression seems to be insufficient to successfully correlate biomimetic indices with pharmacokinetic properties, such as volume of distribution and BBB penetration. Machine learning and Artificial Intelligence techniques have the potential to provide an insight of the underlying molecular mechanism and construct algorithms with superior predictive performance [2, 3].

In the present work several Machine Learning (ML), Deep Learning (DL) and Artificial Intelligence Techniques were used to construct models based on biomimetic chromatographic indices with certain pharmacokinetic properties. For this purpose, retention factors of over than 100 structurally- diverse drugs were considered. Additional physicochemical properties, such as positively and negatively charged molecular fractions, topological polar surface area and molecular weight were tested. Validation of the constructed models was carried out using appropriate test sets.

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Wednesday 22 September 2021

SESSION C1

Food analysis I

Chair: G. Theodoridis, C. Biliaderis

Verification of food authenticity claims: Consolidated strategies and new trends in data modelling

P. Oliveri

Department of Pharmacy (DIFAR), University of Genova – Viale Cembrano, 4 – I-16148 Genova – Italy

E-mail: oliveri@difar.unige.it

Verification of authenticity claims is a challenging analytical problem of interest to many fields, from cultural heritage studies to the control of trade products, including cloths, drugs and, above all, food. The question to be addressed – namely compliance of a given product with the declared claims – is qualitative and closely related to the one of quality control. In the scientific literature, most of the published papers answer such a question by means of discriminant classification methods, but it can be easily demonstrated that discriminant strategies are not appropriate and, in many practical situations, they may lead to incorrect predictions. In fact, all discriminant methods look for a delimiter between two – or more – classes, determined using a contribution from all of the classes considered. This means that all of the classes must be correctly defined and the samples included must be thoroughly representative of each class since they have a crucial influence on the decision rule to be derived. This is extremely important when the focus is on a single class like, for example, cases involving verification of an authenticity claim. In fact, in such a case, the discriminant approach would require the collection of two sets of training samples: one representative of the product to be characterised and a second representative of the entire production of the same product that does not comply with the given claim. Such a condition is rarely realisable in practice, and collected sets of non-compliant samples are often under-representative of the non-compliance possibilities. This inevitably leads to biased decision rules, the outcomes of which are heavily dependent on those samples included in the non-compliant set [1]. The most appropriate family of chemometric methods for addressing this type of problem goes by the name of class modelling [2]. Such methods perform verification of compliance with a specification by defining a multivariate enclosed class space, at a predetermined confidence level, for authentic samples of the class under investigation, enabling what is referred to as one-class classification. The first class modelling methods introduced into chemometrics were SIMCA (soft independent modelling of class analogy) [3] and UNEQ (unequal dispersed classes) [4]. Recently, strategies based on partial least squares (PLS) regression have been introduced, such as the partial least squares density modelling method (PLS-DM) [5]. Models built in such a way has the advantages of describing perfectly the compliant samples and being free from the distribution of non-compliant samples in the training set. Issues related to development, optimisation and validation of suitable class models for authenticity verification of food products will be critically analysed and discussed.

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High throughput analysis for reliable detection of EVOOs adulteration – A non-target metabolomics approach for geographical origin and varietal discrimination

I.F. Tzavellas [1,2], S.K. Drakopoulou [1,2], M.E. Dasenaki [1,2], N.S. Thomaidis [1,2]

[1] National and Kapodistrian University of Athens, Department of Chemistry,
[2]Laboratory of Analytical Chemistry, Panepistimiopolis Zografou, 15771 Athens, Greece

E-mail: ntho@chem.uoa.gr

Extra virgin olive oil (EVOO) has been acknowledged as an emblematic food of the Mediterranean diet, both for its distinctive taste, but foremost for its high nutritional value. The declaration of olive oil superiority, officially substantiated by EU health claim establishment regarding olive oil polyphenols (EU 432/2012), has laid the foundations for in-depth authenticity study in the field to reassure the uniqueness of the product [1].

Olive oil, due to its high financial impact on the global market, is frequently subjected to some kind of adulteration. Talking numbers, olive oil tops the food ingredient fraud ranking according to FDA, with one of the most frequent profit-driven fraudulent procedures being its partial substitution with other vegetable oils. However, special attention must also be paid to the assurance of variety and geographical origin in order to highlight the special qualifications of the product[2].

Serving this purpose, in this study a non-target screening workflow based on LC-QToF-MS analysis have been developed and optimized for the detection of geographical origin and varietal adulteration in EVOOs samples. Thus, two different adulteration studies have been designed and carried out, including the analysis of samples of different EVOO variety and geographical origin, as well as adulterated samples. The adulterated samples were constructed in the lab, in different adulteration ratios, ranging from 10 to 50%. Noteworthy differentiations in EVOOs profile were recorded, while potential authenticity markers of origin and variety for EVOOs discrimination have also been introduced. Data analysis and evaluation in non-target HRMS screening workflows was performed, significantly assisted using advanced data processing tools combined with supervised and unsupervised chemometric techniques.

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A multi-instrumental approach for the physicochemical characterization of a cotton honey-based spread produced by controlling compositional and processing parameters

V. Sereti[1], A. Lazaridou[1*], C. Tananaki[2], C. G. Biliaderis[1]

[1]Laboratory of Food Chemistry and Biochemistry, Department of Food Science and Technology, School of1Agriculture, Aristotle University of Thessaloniki, P.O. Box 235, Thessaloniki 54124, Greece
[2] Laboratory of Apiculture–Sericulture, Department of Horticulture and Viticulture, School of Agriculture, Aristotle University of Thessaloniki, Thessaloniki 54124, Greece

e-mail: athlazar@agro.auth.gr

Creamed (or set) honey is a type of spread produced by a controlled crystallization process resulting in a product with a large number of small, non-detectable by palate crystals, light color and a smooth and creamy texture, rendering it into an easily spreadable product like peanut butter[1,2]. Honey from cotton blossoms tends to crystallize rapidly resulting in a product with hard texture and low spreadability, even if it is processed via this controlled crystallization treatment. The aim of this study was to develop a cotton honey-based spread (CHS) with acceptable physical and textural properties by adding fructose (1-6%) and water (up to 18% final moisture) into the initial cotton honey as well as by controlling the temperature during crystallization, T_{cryst} (5-23°C x 20 days) and the conditioning (tempering) step as a follow-up process, T_{cond} (20-30°C x 10 days). Multi-instrumental analysis (rotational rheometry, differential scanning calorimetry, optical microscopy, colorimetry, texture analysis) and sensory (spreadability, mouthfeel, overall acceptability) evaluation were used for characterization of textural properties of the CHS preparations. The addition of fructose and water were found to be the main factors that affected significantly ($p<0.05$) the quality parameters of the product. More specifically, with increasing fructose and water contents the crystallinity index (i.e. Glucose/Water ratio) in CHS formulations decreased leading to significantly ($p<0.05$) lower values of apparent and complex viscosities, hardness, spreadability work and melting enthalpy of sugar microcrystals, as well as improved the organoleptic characteristics of this spread product. Moreover, with increasing T_{cryst}, only the lightness and the melting enthalpy values decreased, while the damping factor increased. Instead, the T_{cond} did not affect any of the above parameters ($p>0.05$). Additionally, the Glucose/Water ratio was found to correlate positively with all color, thermodynamic, texture analysis parameters and degree of crystallization of the final product, as well as with all the rheological parameters of the spreads, with exception of loss tangent. Overall, strong correlations were identified between the compositional, physicochemical attributes and the sensory characteristics of honey spread, implying that a multi-instrumental analysis of CHS preparations along with sensory evaluation can be a valuable holistic approach for understanding the macrostructure, evaluating the physical properties and quality attributes, as well as predicting the consumer preference of this honey-based product.

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Physical properties and quality evaluation of composite dough and bakery products using a multi-analytical instrumental approach

K. Kotsiou, A. Lazaridou, C.G. Biliaderis

Laboratory of Food Chemistry and Biochemistry, Department of Food Science and Technology, School of Agriculture, Aristotle University of Thessaloniki, P.O. Box 235, 54124, Thessaloniki, Greece

e-mail: athlazar@agro.auth.gr

Fortification of a staple food, rich in rapidly digestible carbohydrates, such as bread, with legumes, which are significant sources of proteins and dietary fibers, might be nutritionally beneficial for improving public health. Nevertheless, inclusion of legume flours into wheat dough formulations is challenging, since it modifies the rheological behavior of the dough [1-2], adversely affecting bread volume and crumb textural characteristics [1-3]. In this study, a multi-analytical instrumental approach was used for the characterization of the physical properties and quality evaluation of composite doughs consisting of wheat and legume flours and the resultant breads. Flours from yellow split pea (YSP) and chickpea (CP) were used, due to their high contents in protein and dietary fiber, to substitute wheat flour at 10-20% levels; moreover, roasting of the legume seeds before milling was employed as a quick and cost-effective pretreatment to eliminate the legume off-flavors and therefore to promote acceptability of the final products. Fundamental rotational rheometry revealed, that inclusion of 15 and 20% CP or 20% YSP flours in the dough formulations, significantly increased their storage modulus and zero shear viscosity, while reduced maximum creep strain and compliance, attributes that are related to strong and elastic doughs with a great resistance to flow and deformation. Empirical rheological testing with farinograph showed that YSP flour inclusion in the formulations at 15 and 20% levels resulted in doughs with significantly decreased stability, while CP flour inclusion at the same levels, improved doughs stability. Extensography showed that incorporation of both CP and YSP flours at high levels in the doughs negatively affected the resistance to extension/extensibility ratio, a factor that is strongly correlated to baking performance. However, the YSP – wheat composite doughs were able to recover their extensograph parameters upon repeated extension, followed by a resting period, thus making them suitable for bread formulations that require multiple cycles of kneading and resting, while the doughs fortified with CP became very resistant to extension after this processing stage. Regarding end-products, higher levels of added legume flours resulted in breads with low specific volumes and harder crumbs as shown by a laser-based scanner and a Texture Analyser, respectively. Monitoring, by Texture Profile Analysis, the changes in crumb hardness during bread storage showed that samples with 20% YSP flour exhibited higher staling rates, which could be partially attributed to the higher rate of amylopectin retrogradation of this formulation as revealed by Differential Scanning Calorimetry. The findings suggest that multi-instrumental analysis of composite doughs and breads is a very useful analytical toolbox for characterization of composite doughs and bakery products with improved nutritional characteristics and for the prediction of their shelf life and consumer acceptability scores.

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Discrimination of feta cheese from different Geographical region, packaging type and production date according to its chemical composition and volatile profile

M. Lygka, T. Massouras

Laboratory of Dairy Research, Agricultural University of Athens, Iera Odos 75, Votanikos, 11855 Athens, Greece

E-mail: lygka@hotmail.com

The present work aims to verify whether the chemical composition and the volatile profile could be considered as chemical biomarkers linked to geographical origin, packaging type, and production date of feta cheese. One hundred and ten (110) samples of feta cheeses aged between 3-12 months, packaged in wooden barrel or in tin container were collected from 9 different geographical regions (throughout mainland Greece and Lesvos). Samples were analyzed for chemical composition (fat, protein, moisture and salt) using a Foodscan apparatus. The volatile compounds were extracted by headspace Solid phase microextraction (HS-SPME) and analyzed using a gas chromatography-mass spectroscopy (GC / MS). The values of chemical parameters and the volatile compounds were treated for the geographical regions, packaging type and production day variables. A total of 84 volatile compounds were identified including 18 acids, 16 alcohols, 8 aldehydes, 14 esters, 20 ketones, 8 alkane and 2 alkene. The most abundant compounds were organic acids followed by alcohols, ketones and esters. Using Multifactor Anova and discriminant analysis, based on volatile profile of cheese samples achieved a classification rate of 85%, 94,83%, and 84,35% for geographical regions, packaging type and production day respectively. Regarding the chemical composition all samples were within specifications of PDO feta cheese (fat in dry matter >43% and humidity up to 56%). Statistically significant higher values of fat and protein content and lower moisture content were found in samples of feta cheese packed in wooden barrels compared those packed in tin container. None of the three variables significantly affected the chemical composition of feta cheese.

Thus, the volatile profile of feta cheese can be considered as a biomarker to differentiate feta cheese of a certain origin, packaging type and production date.

Application of spectroscopic techniques coupled with multivariate statistical analysis for the origin discrimination and quality control of agrofoods

<u>R. Kontzedaki</u>, M. Velegrakis

Institute of Electronic Structure and Laser, Foundation for Research and Technology-Hellas (IESL-FORTH), 700 13 Heraklion, Crete, Greece

E-mail: vele@iesl.forth.gr

Food quality control is often based on modern and sophisticated analytical methods and processes that are costly, time-consuming and require specialized personnel [1], [2], [3]. At the Institute of Electronic Structure and Laser of FORTH, we employ an alternative approach in order to investigate the rapid determination of the characteristic substances contained in food samples, with little or no processing. For this study, optical spectroscopic techniques [Absorption (UV-Vis-NIR), Fluorescence and Raman spectroscopy] are applied. In this way, we can record the optical spectrum that is the characteristic "fingerprint" and reflects the chemical composition of a sample. Different samples have a different fingerprint and the detected differences or similarities, coupled with multivariate statistical analysis methods (Principal Component Analysis–PCA, Partial Least Squares Discriminant Analysis PLS-DA and PLS-regression), are used for origin and quality control. In this work several results of these studies will be presented. Greek white and red wine varieties from different wineries were studied. Also, samples of Cretan olive oil were investigated, that were obtained from the Pancretan Olive Oil Competition and had been graded by a trained team of testers using human senses. The samples were discriminated based on their quality and crop type (organic and conventional cultivation).

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Determination of volatile compounds in Spirulina food supplements using HS-SPME – GC/MS

<u>A. Paraskevopoulou</u>, T. Kaloudis, A. Hiskia, T.M. Triantis[*]

Institute of Nanoscience and Nanotechnology, NCSR Demokritos, Athens, Greece

E-mail: t.triantis@inn.demokritos.gr

Cyanobacteria are photosynthetic microorganisms found in terrestrial and aquatic environments, capable of producing a large number of metabolites with great structural and functional diversity. *Arthrospira spp.*, commercially known as *Spirulina*, is a cyanobacteria species widely used as a food supplement because of its high amount of proteins, vitamins, pigments and fatty acids. In addition to these bioactive substances, *Spirulina* produces numerous volatile organic compounds (VOCs) which are important in determining the quality and nutraceutical properties of spirulina products, including their taste and aroma. Most of these secondary metabolites are not toxic, while some exhibit antimicrobial, anticancer or antibacterial activities [1] and have a wide range of applications in food, cosmetics and pharmaceutical industries [2].

This study focused on the analysis of the VOCs profile in twenty-two (22) commercial *Spirulina* food supplements from different geographical origins using headspace solid-phase microextraction (HS-SPME) combined with gas chromatography mass spectrometry (GC-MS). Volatile compounds were identified after mass-spectral deconvolution, matching of spectra and retention indices using the NIST MS library and AMDIS software.

As many as 229 volatile components were identified and categorized into various chemical classes, while specific compounds were detected in the majority of the analyzed samples, e.g. pyrazines, furans and furanones (3-acetyl-2,5-dimethylfuran; 2(4H)- benzofuranone-5,6,7,7a-tetrahydro-4,4,7a-trimethyl), esters, aldehydes (hexanal; benzaldehyde; 2-heptenal; 2-octenal; safranal; β-cyclocitral), alcohols (1-hexanol; 1-octen-3-ol), ketones (2-heptanone,6-methyl; cyclohexanone,2,2,6-trimethyl; α-ionone; β-ionone epoxide; 5,9-undecadien-2-one), saturated hydrocarbons (pentadecane; hexadecane and heptadecane), terpenes, alkenes, pyridines and benzenes. Identification of compounds was based on a high matching score (80-100%) combining a close similarity of their mass spectra and their retention index values with those of reference compounds present in the NIST MS Library. On the basis of total integrated peak areas of the GC-MS elution profile, the most abundant constituents of Spirulina VOCs were heptadecane (relative content 8% - 20,6%), β-ionone (2% - 6,56 %), pentadecane (1,5% - 5,95%), trans-β-ionone (1,5% - 3,36 %), hexanal (0,1% - 2,4%) and 1-octen-3-ol (0,1% - 2,22 %). Results show the potential of *Spirulina* as an important source of volatile compounds and its possible usage not only as a dietary supplement but also as a raw material for the isolation of these compounds and their further application in the food, pharmaceutical and cosmetics industries.

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SESSION C2

Food analysis II

Chair: P. Oliveri, C. Georgiou

Physicochemical and sensorial properties of wheat bread with sourdough made from starter culture isolated from a submerged chickpea fermentation extract

C. Nouska, M. Hatzikamari, C. G. Biliaderis, A. Lazaridou[*]

Department of Food Science and Technology, Aristotle University of Thessaloniki, Thessaloniki, 541 24, Greece

e-mail: athlazar@agro.auth.gr

During bread production, baker's yeast is used for the development of organoleptic characteristics and the leavening of this food product. Recently, consumers are turning into traditional products, leading the bakery industry to produce bread supplemented with sourdough. Sourdough can be also used as a leavening agent in breadmaking and is responsible for the characteristic flavor and the volume of bread due to the organic acid and gas production. Also, sourdough, may prolong shelf-life of bread by the production of bacteriocins, organic acids, amylolytic enzymes etc. [1,2]. In this work, a Greek traditional sourdough made from a starter culture isolated from a submerged chickpea fermentation extract was used as a leavening agent in wheat breads. The dominant microflora developed during submerged fermentation which was responsible for the leavening effect was found to be clostridia (7.04 ± 0.20 logcfu/g); similar findings have been reported in other studies [3,4]. Provided that clostridia isolated from chickpea sourdough have been found not to form any toxins [3], there was no concern for health hazard. Thus, 10 clostridia strains were isolated from chickpea sourdough and identified with molecular techniques (PCR) as *Clostridium perfringens*; one of these strains was used as starter culture for the production of sourdough from chickpea flour both in liquid and freeze-dried form before incorporated into wheat bread. Loaf specific volume increased with the addition of liquid ($2,64\pm0,05$ ml/g bread) and freeze-dried ($3,19\pm0,08$ ml/g bread) sourdough compared to control bread which was leavened by baker's yeast ($2,20\pm0,04$ ml/g bread). Crumb hardness and melting enthalpy of the retrograded amylopectin in sourdough supplemented breads during storage at $25^{o}C$ for 4 days were lower than control formulation as revealed by Texture Profile Analysis (TPA) and Differential Scanning Calorimetry (DSC), respectively, indicating a lower staling extent. Additionally, organic acids analysis conducted by Anion-exchange HPLC-DAD showed higher acetic acid content for sourdough breads compared to control, while butyric and isobutyric acids were produced only by the sourdough starter culture giving to these bread formulations their characteristic flavor; the latter was confirmed by the sensory analysis which was also showed higher crumb softness and overall acceptability rates for sourdough breads supporting the data from instrumental analysis. Overall, chickpea sourdough from *C. perfringens* starter culture isolated from a submerged chickpea fermentation extract can improve textural and sensorial properties of wheat breads and extend their shelf-life.

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Polystyrene biodegradation by mealworms larvae: Identification of degradation products and metabolic alteration using GC-MS

E.D. Tsochatzis [1], I. E. Berggreen[1], H. Gika[2,3,4], G. Theodoridis[2,3,4]

[1]Department of Food Science, Aarhus University, Agro Food Park 48. 8200 Aarhus N. Denmark
[2]FoodOmicsGR Research Infrastructure, AUTh node, GR 57001 Thessaloniki, Greece.
[3]Biomic AUTh, Center for Interdisciplinary Research and Innovation, GR 57001, Thessaloniki, Greece.
[4]Aristotle University of Thessaloniki, Thessaloniki, Greece.

E-mail: Emmanouil.Tsochatzis@food.au.dk

The present study investigated the biodegradation of polystyrene (PS) plastic by mealworm (*Tenebrio molitor*) on different diets followed by untargeted screening of larvae gut intestine tissue and frass (manure and feed residuals) to investigate the existence of polymer-generated organic residues. Three different diets, consisting of PS, rolled barley and water were tested. PS degradation rates ranged from 16 % to 23 % within 15 days, with no statistical differences in survival rates. The larvae fed with ad libitum barley:PS (20:1 w/w) and water (diet 2) had the highest growth rate and the highest PS consumption, along with barley:PS of 4:1 w/w (diet 1) (Figure 1).
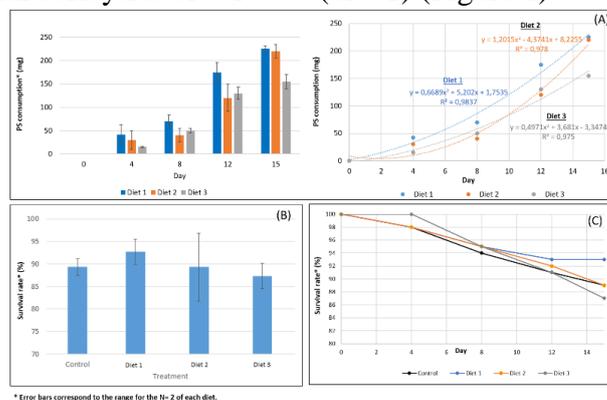
Figure 1. (A) PS consumption throughout the experiment; (B) final survival rates of *T. molitor* larvae) within the different applied feeding experiments, and (C) survival rate during the experiment. Diet 1 is barley:PS (4:1), Diet 2 is ad libitum barley:PS (20:1) + H_2O and Diet 3 is ad libitum barley:PS (20:1).

A GC-TOF-MS analysis revealed no contaminating substances in the gut intestine tissue, nor styrene or PS oligomers, whilst several bioactive compounds and traces of alkanes, mostly with small carbon chains, were present. Metabolomics analysis on the collected frass, either on the lipophilic ($CHCl_3$) or the polar fraction ($MeOH-H_2O$) was performed. Styrene and PS oligomers (dimers, trimers) were identified, though in a relatively low total amount, up to a total of 346.0 ng/mg. 2,4 di-tert butylphenol was identified in both frass and tissue, coming from the PS polymer (Non-intentionally added substances; NIAS) (Figure 2). It was shown that degrading and contaminating organic compounds occur at low levels, in both gut intestine and frass, during bio-degradation of PS.
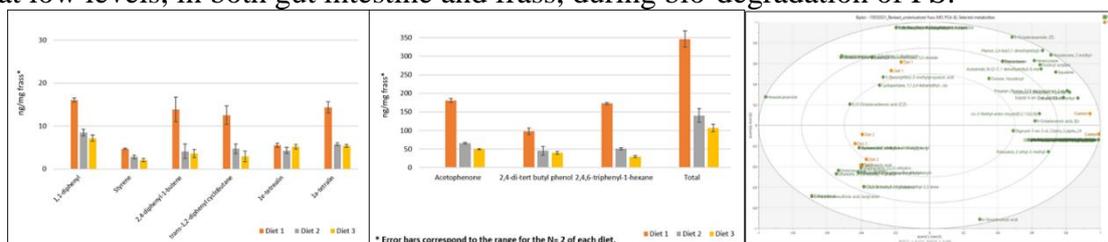

Figure 2. Mass fractions (ng/mg) of monomer (styrene) and PS oligomers in the frass from PS biodegradation by *Tenebrio molitor* larvae.

In the polar fraction of frass, bioactive molecules (fatty acids, amides) were identified, together with several hydrocarbons, mostly with longer carbon chains. The formation of these substances indicated enzymatic and biochemical activity in the larvae-gut intestine.

Elemental metabolomics for honey authentication

S.A. Drivelos[1], G. Danezis[1], M. Halagarda[2], S. Popek[2], C.A. Georgiou[1]

[1]Chemistry Laboratory, Department of Food Science and Human Nutrition, Agricultural University of Athens, 75 Iera Odos, 118 55 Athens, Greece
[2]Department of Food Product Quality, Cracow University of Economics, ul. Sienkiewicza 5, 30-033 Krakow, Poland

E-mail: cag@aua.gr

Elemental metabolomics concerns food safety and quality. Utilization of elemental profile promotes authenticity and leads to useful markers such as rare earth elements. Elemental profiles have proved reliable indicators for geographical origin determination in many cases and furter they could provide information on genetic origin. Processing methods are also related to elemental fingerprint. Omics technologies harness the impressive power of high resolution mass spectrometry (HRMS) [1]. Although REEs and precious metals have great potential for determination of geographical origin they have not been studied extensively. Increased interest in rare earths as authentication markers will grow stronger in the near future.

Trace and rare earth elements of 93 honeys of different botanical type, origin and production method have been studied through ICP-MS. Information obtained using rare earths and trace elements were utilized via chemometric evaluation for the classification of honey samples. Discriminant Analysis (DA) was successful for botanical type and geographical origin classification while Cluster Analysis (CA) was successful only for botanical type. Through Probabilistic Neural Network (PNN) analysis, 85.3% were correctly classified by the network according to their geographical origin and 73.3% according to production method, organic – non organic. A Partial Least Squares (PLS) model was constructed, giving a prediction accuracy of more than 95%. This study shows that elemental profile could discriminate honey according geographical and botanic origin. This is an exploratory study pointing out the value of a large scale sampling study. To explain the incapability of elemental profile to discriminate honey production method a study on the elemental content of floral nectars and their changes along pesticides use is required [2].

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Advanced and Comprehensive Screening Methodologies for the Monitoring of Intentionally & Non-Intentionally Added Substances in Food Contact Materials

C. Kanakaki, E. Mermingi, N. Thomaidis

Department of Chemistry, National and Kapodistrian University of Athens, Panepistimiopolis Zografou, 157 71 Athens, Greece

E-mail: ckanakaki@chem.uoa.gr

The packaging has become an indispensable element in the food manufacturing process, as it protects the food e.g. from light and microbial contaminations, improving the self-life and stability of products during storage. However, it has been found to represent a source of contamination itself through the migration of substances from the packaging into food. The identification of these substances is an issue that deserves great attention to assure food quality and consumer safety.

Aiming the comprehensive screening of intentionally and non-intentionally added substances (IAS & NIAS) in food contact materials (FCMs), GC-MS/MS and GC-APCI-QTOFMS methods were developed, for the target screening of volatile migrants. Furthermore, corresponding contaminants of low volatility, thermal instability and high polarity were identified by LC-MS/MS and LC-ESI-QTOFMS methods. Particularly the high resolution mass spectrometric (HRMS) methods applied were utilized for further suspect and non-target screening and retrospective analysis of the obtained FCMs. For these screening procedures the entire spectrum of food simulants was evaluated with different migration protocols being employed, depending on the intended use and origin of the analysed samples. Risk assessment of the identified migrating compounds was also performed using the corresponding legislation, online tools and in-house built software.

Application of the developed workflows into a large set of products and comprehensive screening of migrating chemical contaminants will promote the production of inert and consequently safe FCMs. Thus, this work will assess public health risks, by facing current and future challenges.

Wednesday 22 September 2021

SESSION C3

Chemistry associations

Chair: V. Samanidou, M. Ochsenkühn, M. Karayannis

FoodOmicsGR_RI: A Distributed National Research Infrastructure for the Comprehensive Characterisation of Foods

G Theodoridis

Department of Chemistry, Aristotle University of Thessaloniki, Thessaloniki, 54124, Greece

E-mail: gtheodor@chem.auth.gr

Foodomics is the omics field and the technology that combines food and nutrition sciences with advanced analytical techniques and bioinformatics, whereby a hypothesis-free approach is applied to characterize the food composition or food consumers' biofluids, addressing new challenges such as authentication, traceability, improvement of food produce, food quality and nutritional value. This holistic approach employs cutting-edge research infrastructure, where big data is produced generating new knowledge, highlighting previously unknown associations of biomolecules with the studied phenotype.

FoodOmicsGR_RI is a national research infrastructure aiming to perform and aid omics research in agri-food sector in the Greek research environment. This field is of high importance for the country and its agriculture because of the unique landscape Greece exhibits that result in a diverse and rich portfolio of Greek local products and foods. Therefore, the central scope of FoodOmicsGR_RI is to support the Greek agri-food sector by generating robust data on the composition and nutritional value of the local produce, thereby increasing product's position, market demand and provide a higher revenue for the producers.

The project comprises eight (8) Greek Universities and Research Centers. Analytical groups and food specialist groups from Universities of Thessaloniki, Athens, Crete, the Aegean, Ioannina, the Agricultural University of Athens, the International Hellenic University, and the Biomedical Research Center of the Academy of Athens comprise a team of 60 staff scientists and 40 newly recruited researchers from 20 scientific disciplines. BIOMIC AUTh, a new interdisciplinary laboratory of the Aristotle University Thessaloniki (http://biomic.web.auth.gr/), having long expertise in bio-analytical chemistry, is coordinating the project and is providing the current presentation. More details can be found on our website (http://foodomics.gr/).

EURACHEM Greece: promoting quality of analytical measurements, QA& QC in analytical laboratories.

M. Kostakis[1], V. Zonaras[2], H. Gika[3], A. Iakovakis[4], Christina Virgiliou[5], G. Theodoridis[5]

[1]Laboratory of Analytical Chemistry, Department of Chemistry, National and Kapodistrian University of Athens, Panepistimiopolis Zografou, 15771 Athens, Greece
[2]Laboratory of Pesticide Residues, Scientific Directorate of Pesticides' Control and Phytopharmacy, Benaki Phytopathological Institute, 14561 Kifissia, Attica, Greece

[3] Laboratory of Forensic medicine & Toxicology, School of Medicine, Aristotle University of Thessaloniki, Greece
[4] Veltia labs, Sindos 570 22 Thessaloniki

[5]Laboratory of Analytical Chemistry, Department of Chemistry, Aristotle University of Thessaloniki, Greece.

E-mail: gtheodor@chem.auth.gr
makostak@chem.uoa.gr

Eurachem is a network of national and other organizations in Europe that focuses on measurement quality and quality-related issues [1]. For more than 30 years, Eurachem organizes activities (workshops and training events) and provides documents (guidelines, information leaflets) in order to help laboratories, accreditation bodies, authorities, other stakeholders, practitioners, and researchers to ensure the quality of analytical measurement. Method validation, measurement uncertainty, metrological traceability, internal quality control and proficiency testing are the main cornerstones of Eurachem [1]. Moreover, upcoming analytical measurement issues, like sampling and conformity assessment, are also included in Eurachem working activities. In the direction to promote quality assurance in analytical measurement, all material (guidelines, leaflets, workshops' presentations) produced by Eurachem is available free of charge on the website (eurachem.org) of Eurachem.

The aim of the presentation is to inform the interested parties about Eurachem Greece, about the Eurachem's working groups, communicate their activities and discuss forthcoming issues and problems of the interest of Greek research ecosystem and subsequently bring them into the attention of the Eurachem.

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EuChemS-WP EiC: EuChemS Working Party on Ethics in Chemistry

A. Silvestru on behalf of EuChemS-WP EiC Steering Committee

Babeș-Bolyai University, Chemistry Department, Arany Janos 11, 400028 Cluj-Napoca, Romania

anca.silvestru@ubbcluj.ro

The WP EiC is a network of representatives of the EuChemS member societies with the goal to promote the understanding of the importance of ethical considerations in education and practice of pure and applied chemistry.

The WP EiC was founded shortly after 2010, when Hartmut Frank, Professor of environmental chemistry at the University of Bayreuth, together with other remarkable scientists, sustained by the Nobel laureates Richard Ernst and Roald Hoffmann, pointed out that the tremendous development of chemistry at the beginning of the 3rd millennium should be accompanied by an appropriate education and ethical approach in all chemistry-based activities, which might raise concerns regarding the use and the impact upon the environment and the society development. The group was subsequently enlarged by attracting other chemists and scientists and it was integrated as a Working Party in EuChemS.

The vision of WP EiC is based on the responsibilities of chemists on both internal and external level of their decisions and actions. The internal one takes into account a good scientific practice, research and professional integrity, while the external one includes topics such as the potential dual applications of chemical innovations, sustainability, risk assessment, as well as the role of specific expertise in science policy, innovation governance and political decisions.

The WP EiC is promoting ethical guidelines in science and education, cooperation and scientific exchanges between specialists from different countries and specific fields related to chemistry, encourages interaction and communication between young and experienced scientists, between academia and industry, thus fostering the scientific progress and innovation and aiming to better life, a healthy society, peace and a clean environment.

The efforts of the WP EiC to contribute to an ethical education of the young generations of chemists and scientists, our care of the ethical, social and cultural dimensions of practicing chemistry in the interest of the whole society is also reflected in the materials elaborated by several members of the group: Good Chemistry: Methodological, Ethical, and Social Dimensions, an online course (Jan Mehlich) and offered on the e-learning platform of EuChemS to students in EU, papers on topics at the intersection of chemistry and applied ethics (i.e. Angew. Chem. Int. Ed. 2011, 50, 8482 and Chem. Eur. J., 2017, 23, 1), conferences, etc.

The WP EiC gathers today members from 17 countries, representatives of national chemical societies or just members from academia and industry, interested for the future development of the society in times when chemistry should solve in an ethical way a lot of problems: resources economy, environment, health, weapons prohibition, etc. You are invited to join in a formal or informal way to our goals and care, by considering the importance of a responsible education, science, and specific activities in chemistry and associated fields.

For more information, please visit: https://www.euchems.eu/divisions/ethics-in-chemistry/ .

Wednesday 22 September 2021

SESSION C4

Environmental analysis | Chromatography | Mass spectrometry

Chair: A. Zouboulis, G. Hieftje

Development of an HRMS-based workflow to quantify PFAs in leachates and urban waters

L.-A Koronaiou[1,2], C. Nannou[1,2], D. Lambropoulou[1,2]

[1]Laboratory of Environmental Pollution Control, Department of Chemistry, Aristotle University of Thessaloniki, GR-541 24 Thessaloniki, Greece
[2]Centre for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center, Thessaloniki, 10th km Thessaloniki-Thermi Rd, GR 57001

E-mail: dlambro@chem.auth.gr

Per- and polyfluoroalkyl substances (PFASs) are a diverse group of synthetic organofluorine compounds that have been widely used in industrial applications and consumer products such as non-stick cookware, food packaging, fire-fighting foams, carpeting, apparels, and metal plating. PFASs are persistent in the environment, bioaccumulative in wildlife and humans and known to cause reproductive and developmental toxicity in laboratory animals and wildlife. Landfill leachate is the major pathway by PFAS exit the municipal solid waste landfills, where their concentrations vary over time and can be much greater than those found in sanitary wastewaters. On the other hand, especially PFOA and PFOS are poorly removed in the drinking water production chain by conventional purification processes but can be removed with active carbon filtration or by reverse osmosis. The present study comprises a comprehensive development of an analytical methodology aiming to trace and quantify a representative array of PFAs at the sub μg per liter scale, alleviating the potential sources of cross contamination as well. In order to achieve sufficiently low LOQs, optimized sample pre-treatment using solid phase extraction (SPE) was conducted. Extractions from water were performed using weak anionic exchange (WAX) SPE cartridges that were tested over HLB cartridges and prove to be more efficient for a large part of the target compounds. Isotopically-labelled standards were used for internal quantification. A high resolution (HR) Orbitrap Q Exactive Focus mass spectrometer (Thermo Fisher) equipped with a heated ESI source was used, acquiring spectra in negative ionization mode. The overall method was validated in terms of linearity, accuracy, precision, and method limits, while matrix-effect study demonstrated that the use of matrix-matched calibration is necessary to quantify with accuracy in such complex matrices. The method yielded acceptable recoveries (70%) for the vast majority of compounds, while the LOQs were equal or lower than ten ng/L. The application of the method in leachates as well as in WWTPs, revealed the occurrence of L-PFBS, PFHxA, and PFOA, while in WWTPs 6:2FtS, PFOSK, and PFOA were detected. The screening capabilities of this system, also allows the simultaneous or retrospective monitoring of other PFASs as well as the suspect screening of other emerging contaminants, using the raw data files obtained by the HRMS instrument.

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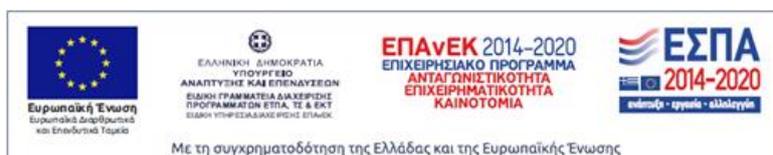
Combination of single algae-ICP-ToF-MS and multivariate statistics for (eco-) toxicological assessment

M. von der Au[1], O. Borovinskaya[2], C. Büchel[3], B. Meermann[1]

[1]Federal Institute for Materials Research and Testing (BAM) - Division 1.1 Inorganic Trace Analysis, Berlin, Germany
[2]TOFWERK AG, Uttigenstrasse 22, 3600 Thun, Switzerland
[3]Institute of Molecular Biosciences, University of Frankfurt, Max von Laue Str. 9, 60438 Frankfurt, Germany

E-mail: Marcus.von-der-au@bam.de

Diatoms are located at the bottom of the food chain. Thus, toxicological relevant metals taken up by diatoms can possibly accumulate within the food web and cause harmful effects. Diatoms are a common test system in ecotoxicology. Toxicological effects weaken the growth of algae which is by default investigated by means of fluorescence detection - diminished fluorescence compared to a non-exposed control group indicates an effect. On basis of the expose concentration as well as obtained fluorescence data potential threshold exceedance in e.g. surface waters is assessed.

However, this approach does not allow for the determination of "real" accumulated metal concentration in diatoms. Common approaches are based on microwave assisted digestion and elemental analysis via e.g. ICP-MS, ICP-OES or AAS. But, with regard to low absolute metal-content in algae this strategy is only feasible in case of availability of a high biomass.

To tackle this problem, alternative, complementary approaches are highly needed. Within the last years, sp-ICP-MS for nanoparticle as well as single cell analysis turned out as a powerful technique to analyze metal contents as well as size distributions on broad size range (nano- to low micrometer scale) [1]. But, common ICP-MS systems do not allow for multi-element detection within single particle/cell events [2, 3]. Thus, simultaneous MS detection devices are needed - just recently, ICP-ToF-MS experienced a revival [4].

Within our previous work, we developed an automated sample introduction system based on a HPLC system on-line with single particle-ICP-MS, which allowed for ionic background separation and single algae analysis [5]. However, for unambiguous tracing several fingerprint elements and multielement analysis in single algae (diatoms) is needed. Thus, we coupled our previous setup on-line to ICP-ToF-MS. Test diatom species were exposed to test substances (Zn) as well as nanoparticles (FeNPs).

The developed setup allowed for a fast, automated and multielement analysis in single diatoms. Furthermore, we combined our approach with multivariate data assessment - multielement detection of characteristic fingerprint elements allowed for an unambiguous diatom tracing. Clustering of diatoms according to metal exposure concentration levels was enabled. Our approach is a new potential tool in ecotoxicological testing.

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Relevant pesticide transformation products in surface waters using target and suspect screening with LC-HRMS

S.Petromelidou[1,2], C. Nannou[1,2], E. Evgenidou[1,2], D. Lambropoulou[1,2]

[1] Laboratory of Environmental Pollution Control Department of Chemistry, Aristotle University of Thessaloniki, Thessaloniki, GR–541 24, Greece
[2] Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Thessaloniki, 57001, Greece
E-mail: dlambro@chem.auth.gr

The intensification of agricultural production in recent decades has forced the rapid increase in pesticides use, which end up in the aquatic environment through processes such as surface runoff, rinsing, careless disposal of empty packaging, etc. Once entering the aquatic bodies, they pesticides undergo biotic and abiotic processes, resulting in their transformation into products that maybe sometimes are even more toxic than the parent compounds. As the determination of the parent compounds in the aqueous samples has been extensively studied, the scientific interest has shifted on their TPs, whose environmental fate is still under investigation. Additionally, their occurrence at relatively low concentrations and the absence of available analytical standards hampers their determination which turns to be a time consuming and laborious process. The presence of residues of pesticides and/or their transformation products (TPs) in surface water is directly related to water quality. Hence, to determine a worst-case impact of pesticide application in agriculture on surface water quality, the screening of TPs along with the parent compounds is desirable.

To this end, the aim of the present study was to utilize liquid chromatography– high-resolution mass spectrometry (LC– HRMS) for the identification of "suspect" TPs (suspect screening). Specifically, the aqueous samples were subjected to solid phase extraction (SPE) prior to analysis, so that a sufficient preconcentration factor be achieved. Afterwards, the analysis was performed by means of a Q Exactive Focus Orbitrap MS, operating in both positive and negative ionization mode. A dynamic list of suspect compounds was created, based on the pesticides that are used systematically and are frequently found in surface waters, according to recent literature. The *a priori* information included in the database was the molecular formula, the exact mass of the pseudomolecular ion, the electrospray ionization (positive/negative) and any available fragment ions from literature. The criteria for the identification of the suspect compounds were based on intensity, isotope pattern, retention time, and *in silico* fragmentation. This way, the suspect hits were prioritized and verified. After processing, 40 transformation products were identified, originating from 25 parent pesticides. The most common detected TPs were those of prometryn, penoxsulam, pyrimethanil, tebuconazole, thiabendazole and imazalil. Specifically, the TP of pyrimethanil with a molecular weight of 250.1186, was identified in all samples. The frequency of detection of imazalil TP with a molecular weight of 257.0243 was also high, since the TP was detected in 77% of the samples. Most TPs arose either from the process of hydrolysis, as almost all categories of pesticides are subject to hydrolysis, or photolysis since the samples studied were surface waters and are heavily affected by sunlight. Ultimately, to predict the acute and chronic toxicity of TPs, an *in-silico* approach based on ECOSAR was implemented, revealing increased chronic toxicity for the majority of the identified TPs, while none of them presented serious acute toxicity values.

Optimization of the sample preparation step by central composite design for the analysis of a wide array of emerging contaminants in wastewaters and tap water using LC-HRMS

A. Ofrydopoulou[1,2], C. Nannou[1,2], E. Evgenidou[1,2], D. Lambropoulou[1,2]

[1] Laboratory of Environmental Pollution Control, Department of Chemistry, Aristotle University of Thessaloniki, Greece
[2] Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Thessaloniki, 57001, Greece

E-mail: dlambro@chem.auth.gr

Multi-residue analysis is highly desirable for water quality control. To this end, a comprehensive workflow for the quantitative analysis of 172 emerging contaminants (pharmaceuticals and personal care products, illicit drugs, organophosphate flame retardants and perfluoroalkyl substances) has been developed for application to wastewater and tap water, based on solid phase extraction (SPE) and Orbitrap high resolution mass spectrometry (HRMS). Due to the large number of analytes with various physicochemical characteristics, the response surface methodology (RSM) employing a central composite design (CCD) and desirability function (DF) approach was exploited to optimize the sample preparation process, instead of the conventional single-factor analysis. The factors included in the design of experiments (DoE) were sample pH, eluent solvents composition and volume. Statistical analysis (ANOVA) proved the adequacy of the proposed model (2- factor interaction) as p-value < 0.05 followed by different diagnostic tests confirmed the good fitting. The best values to acquire DF close to 1 were pH 3.5, methanol/ethyl acetate ratio 87:13 and eluent volume 6 mL. The streamlined method was validated in terms of accuracy, linearity, method limits, reproducibility, and matrix effect. The proposed workflow combines sensitivity and robustness, with recoveries over 70%, method quantification limits <1 ng/L, and relative standard deviations <20% for most of the compounds. Slight matrix effect (ME) was observed except for OPFRs, for which strong ME was calculated. Method applicability was tested over wastewater collected from a municipal wastewater treatment plant in Thessaloniki (Greece), revealing the presence of 69 and 40 compounds in influents and effluents, respectively, at varying concentrations.
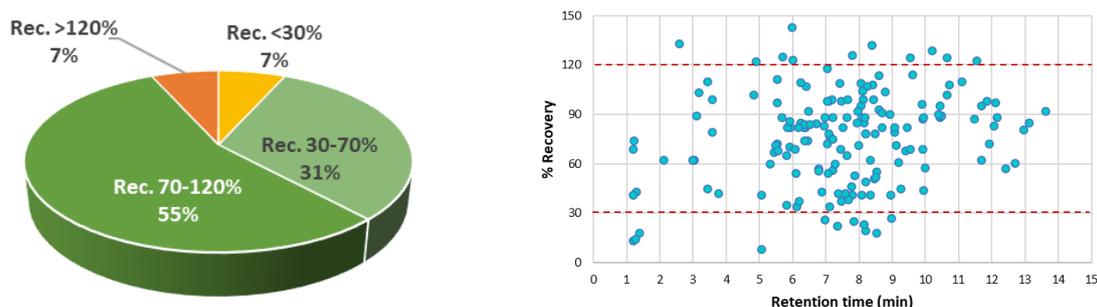
Fig. 1 (a) Percentage of pollutants according to their recoveries at a concentration level of 100 ng L^{-1} (tap water), (b) 2D mapping of the recovery percentages for all selected compounds over the 15-min LC run.

Occurrence of microplastics in wastewater treatment plant in the city of Thessaloniki

D. Kalaronis [1], N. M. Ainali [1,2], E. Evgenidou [1,3], M. Papageorgiou[4], A. Christodoulou[4], I. Lioumbas[4], G. Z. Kyzas[5] D. N. Bikiaris [2], D. A. Lambropoulou [1,3]

[1]Laboratory of Environmental Pollution Control, Department of Chemistry, Aristotle University of Thessaloniki, GR-541 24 Thessaloniki, Greece
[2]Laboratory of Polymer Chemistry and Technology, Department of Chemistry, Aristotle University of Thessaloniki, GR-541 24 Thessaloniki, Greece
[3]Center for Interdisciplinary Research and Innovation (CIRI-AUTH), GR-57001, Balkan Center, Thessaloniki, Greece
[4]Thessaloniki Water Supply and Sewage (EYATH) Co. S.A., GR-54622, Thessaloniki, Greece
[5]Department of Chemistry, International Hellenic University, GR-65404, Kavala, Greece

E-mail: dlambro@chem.auth.gr

The large-scale production of plastic products affects on environment since the huge amount of polymers waste is difficult to be managed properly and thus a significant quantity of them release to Ecosystems each year. As microplastics (MPs) can be defined the polymeric fragments with size < 5mm. Nowadays, wastewater treatment plants (WWTPs) are considered as one of the main pathways for MPs release in freshwaters. Although most of the WWTPs have effectively removed the microplastics (removal efficiency can range from 70 to 99%), the municipal WWTP is still an important source of microplastics in the environment. In this light, the present study is focused on MP contamination jn WWTP in the city of Thessaloniki which serves a population about 1,000,000 people in urban area of Thessaloniki. A systematic sampling was carried out from March 2021 to July 2021 with aim to identify and quantify MPs particles. The collected samples were concentrated into smaller volume through sieving with metal sieves (4-0.125 mm) and rinsing with ultrapure water. Then, concentrated samples were filtered through fiberglass filters (1.6 μm pore size) and an oxidation digestion was applied with 30 ml of 30% H_2O_2 for 3 days under continuous stirring to remove the organic matter. Subsequently, filters were dried in room temperature and examined through a stereoscopic microscope equipped with a digital camera for the mapping of filters with possible MPs particles. The next step included the application of a coupled chromatographic technique, using pyrolysis–gas chromatography / mass spectrometry (Py–GC/MS) for the identification well as quantification of MPs found in the real WWTP samples. Regarding our results, many polymeric fibers were identified and particles (including fragments and films) in different colors and shapes. According to the identification part, the majority of MPs particles were several types of polyethylene, polypropylene, polystyrene, and acrylics, while the polyethylene particles were quantified with a range value between 0.06-0.04 mg/m^3.

Untargeted analysis of cyanobacterial metabolites in lake Karaoun, Lebanon, by LC-HRMS with CyanoMetDB

S.-K. Zervou[1], N. A. Hammoud [1,2], S. Godin[2], T. Kaloudis[1], A. Hiskia[1], J. Szpunar[2], R. Lobinski[2]

[1] Laboratory of Photo-Catalytic Processes and Environmental Chemistry, Institute of Nanoscience & Nanotechnology, National Center for Scientific Research "Demokritos", 15310 Agia Paraskevi, Athens, Greece
[2] Institute of Analytical and Physical Chemistry for the Environment and Materials, (IPREM-UMR 5254), CNRS, E2S UPPA, 64000 Pau, France

E-mail: s.zervou@inn.demokritos.gr, ryszard.lobinski@univ-pau.fr

Cyanobacteria are photosynthetic prokaryotes that are naturally present in surface waters. Under favorable environmental conditions (temperature, nutrients, light), they can quickly multiply, forming cyanobacterial blooms. During the last decades, bloom events have been occurring with increasing frequency and severity worldwide, due to the climate change and increase in nutrients from anthropogenic activities. Cyanobacteria produce a huge number of secondary metabolites, including potent toxins, known as cyanotoxins and other bioactive peptides known as cyanopeptides. Due to the effects of these compounds on public health and on ecosystems, there is an urgent need to detect and identify known and unknown cyano-matabolites in surface waters. This is a great analytical challenge, since the number of analytical standards is extremely limited and fragmentation mass spectra are available publicly only for a few metabolites.

In the frame of the present study, a high resolution mass spectrometry (HRMS) method was developed to investigate the presence of cyano-metabolites in cyanobacterial bloom samples. Samples were collected from Lake Karaoun, Lebanon. Intracellular and extracellular metabolites were extracted with appropriate analytical protocols [1,2,3]. Analysis was performed by reverse phase LC coupled to an Orbitrap Fusion Lumos Tribrid mass spectrometer equipped with electrospray (ESI) ionization source operated in positive mode. Two fragmentation modes, the collision-induced dissociation (CID) and the higher-energy C-trap dissociation (HCD) were applied in order to obtain fragmentation spectra. Data treatment was carried out using Compound Discoverer software in combination to the new CyanoMetDB mass list [4] and related tools and databases for detection, identification and structural elucidation of the cyanobacterial metabolites. *In silico* fragmentation for confirmation of proposed structures was performed using Mass Frontier software and fragment ion search (FISh) scoring. Results indicate the presence of several congeners of microcystins and of some less studied cyanopeptide classes of anabaenopeptins, aeruginosins, cyanopeptolins, microginins and aerucyclamides proving the suitability of this approach for untargeted analysis of cyanobacterial metabolites in environmental samples.

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Hydrometallurgical recovery of scandium from bauxite residue by phosphoric acid leaching

L.A. Tsakanika[1], G. Panagiotatos[1], T. Lymperopoulou[2], E. Chatzitheodoridis[3], M.O.Petropoulou[1]

[1]Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering,; [2]Products and Operations Quality Control Laboratory; [3]Laboratory of Mineralogy, Petrology and Economic Geology, School of Mining and Metallurgical Engineering; NTUA, Iroon Polytechniou 9, Zografou campus, 15773 Athens, Greece

E-mail: veralyb@chemeng.ntua.gr,

Scandium (Sc) is a valuable element with important applications in modern and green technologies. In recent years, the European committee has classified it, as a critical raw material, due to its limited exploitable deposits and unstable supply chain. Hence, the recovery of scandium from secondary raw materials or industrial by products such as bauxite residue, is focused as an alternative promising solution. Bauxite residue (BR) also known as red mud, is the caustic and fine grain sized by-product after bauxite digestion for alumina's production with serious deposition problems and environmental impact. However, the waste is enriched in numerous valuable elements including Sc and rare earths (REEs). In Greece, BR contains significant amounts in REEs ($\sim$1kg REEs$\cdot$t^{-1}) with a high content in Sc ($\sim$100g Sc $\cdot$t^{-1}), stable for the last decades[1,2]. Thus, Greek by-product can be considered as a candidate scandium resource improving also its environmental fingerprint. Hydrometallurgical processes (direct leaching with acidic and alkaline solutions) are the most common techniques studied for Sc recovery[3]. In most cases Sc is extracted at the expense of selectivity especially with regard to iron.

The present study highlights the use of phosphoric acid as leaching agent for hydrometallurgical scandium recovery without any BR pretreatment. The selection of phosphoric acid was based on phosphorous species selectivity for Sc as well as its efficiency on other raw materials for Sc recovery or main elements removal[4,5]. The phosphoric leaching efficiency was evaluated according to Sc extraction yield and lower reagents' consumption. Various acid concentrations, solid to liquid ratios, final pH value, leaching temperatures and times were examined individually as well as in a comparative way aiming to the optimum conditions. The elemental analysis for Sc and main elements (Fe, Al, Si, Ti) in raw material and leachates solutions was performed by ICP-OES and AAS. The chemical, mineralogical analysis as well as the morphology and semiquant analysis of dry BR before and after leaching process was conducted using the XRF, XRD and SEM-EDAX techniques respectively. The characterization of the initial raw material was also studied by Raman analysis. According to the results, it was found that the optimum conditions for Sc recovery are the following: C_{acid}=5M, S/L=10%, T=25 ^{o}C, t=1h, with a scandium recovery about 50%.

Keywords: scandium, bauxite residue, leaching, phosphoric acid, ICP-OES, XRF, Raman

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Rare earth elements determination by inductively coupled plasma mass spectrometry after alkaline fusion

H.T. Mnculwane

Mintek, 200 Malibongwe Drive, Randburg, 2194

E-mail: hlengiwem@mintek.co.za

There is an increasing worldwide demand for rare earth elements (REEs) in new technology applications such as electronics, superconductors, lasers, computers, rechargeable hydride batteries, artificial diamonds, glass and ceramics, space applications, etc. Some of the world's most exotic and innovative technologies that play a significant role in our day-to-day life owe their success to rare earths. Although REEs are present in low concentrations in most minerals (over 190 minerals containing significant amounts of REEs), the primary resources of REEs are only three minerals, namely bastnaesite, monazite and xenotime. The determination of REEs in geological and metallurgical samples is extremely important because they are the source materials for all REEs products.

In order to improve the accuracy and precision of REEs and Y measurements by inductively coupled plasma - mass spectrometry (ICP-MS), the following conditions must be fulfilled: complete digestion of refractory phases and minerals in the samples, low procedural blanks, separation of interfering matrix and elimination of molecular and direct isobaric interferences on the mass of the analytes of interest. Conventional methods that use hotplate or high-pressure Parr Bombs and a mixture of HCL-HNO_3–HF-$HCLO_4$ [1,2] are effective but can take days to achieve complete dissolution of samples that contain highly refractory phases which is unacceptable in the mining and metallurgical industry. Such difficult samples necessitate high-temperature alkaline flux fusion because it offers a faster and more efficient alternative to acid digestion.

This paper presents an accurate, precise, rapid and reliable method for rare earth element analysis by ICP-MS with low detection limits, wide linear dynamic range and simple operation [3,4], which involves no sample pre-concentration, and is therefore, able to deliver data rapidly. The analytical performance of the improved method was evaluated in terms of limit of detection (LOD), limit of quantification (LOQ), linearity, accuracy and precision. The procedure was tested successfully on various REEs rock type certified reference materials and the results obtained were in reasonable agreement with published certificate values.

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Thursday 23 September 2021

SESSION D1

Materials analysis

Chair: M.P.Morales, K.Kordatos

The role of size and hyperthermia treatment on the distribution of magnetic nanoparticles for theranostic applications

F. Pinakidou, M. Katsikini, K. Simeonidis, E.C. Paloura, M. Agelakeris

School of Physics, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece

E-mail: fpina@physics.auth.gr

The spatial distribution and nanostructure of magnetite magnetic nanoparticles (MNPs) implemented in hyperthermia applications is addressed by combining SR-based μ-XRF elemental mapping with XAFS measurements. Their excellent behaviour towards tumour destruction is attributed to their increased heat conversion efficiency, a property that correlates directly to their size and also the structural characteristics of iron. We investigated a series of Fe_3O_4 MNPs with varying sizes (8 to 80 nm) prior and after hyperthermia treatment. As it is evident from the 2D μ-XRF maps shown in the insets in Fig. 1, the smallest MNPs (8 nm) are spread homogeneously with no preference for segregation, while upon increasing the size of the MNPs, the tendency of Fe for extended clustering becomes more prominent. In contrast, between MNPs of the same size, hyperthermia does not have a prominent effect on their distribution; only when the diameter is equal to 40 nm there exists a slight tendency for desegregation of the widely dispersed Fe-rich inclusions. These conclusions based on the μ-XRF 2D maps, can be quantitated by calculating the distribution of 15x15 $μm^2$ clusters of MNPs and are shown in Fig. 1. According to the Fe-K-XAFS analysis results, the majority of Fe^{+2} transform to Fe^{+3} while no pronounced modification in site occupancy is detected.
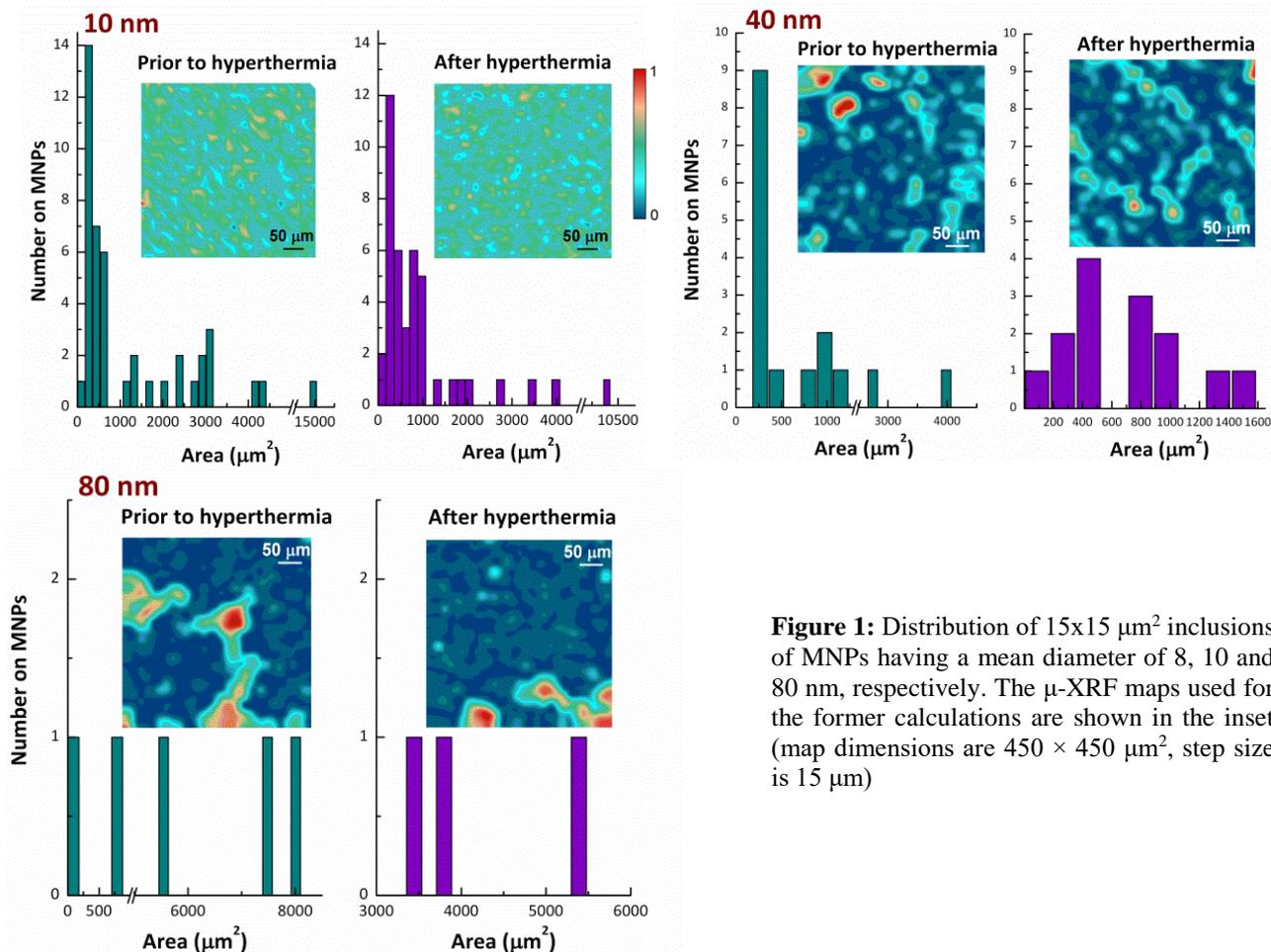

Figure 1: Distribution of 15x15 $μm^2$ inclusions of MNPs having a mean diameter of 8, 10 and 80 nm, respectively. The μ-XRF maps used for the former calculations are shown in the inset (map dimensions are $450 × 450\ μm^2$, step size is 15 μm)

Characterization of advanced technological material using XRD and SEM

E. Delli, E. Pavlidou, G. Vourlias

Laboratory of Advanced Materials and Devices, Department of Physics, Aristotle University of Thessaloniki, GR541 24 Thessaloniki, Greece

E-mail: evdelli@physics.auth.gr

Commercialization of conventional materials has greatly benefited several areas of everyday life and economy. However, the newly facing environmental challenges, the limited reliability and narrow life-cycle performance of old material systems as well as the constantly increased demand for advanced products reviled the necessity of establishing new technological materials with improved properties that can meet the manufacturing expectations. Developing of advanced technological materials requires experimental producing and thorough testing of the properties and performance of the new material to certify its usage and large scale production capability. In depth knowledge of the structural and chemical properties is considered the first step towards the fully understanding of a material and its usage. X-rays diffraction (XRD) and Scanning Electron Microscopy (SEM) are among the most used material characterization techniques enabling surface imaging, composition identification, structural quality and homogeneity investigation, failure analysis, etc. In this work we present the XRD and SEM results obtained for a wide range of advanced materials, such as polymer nanocomposites, thermoelectrics, anticorrosion coatings, biomaterials, minerals, etc. Results analysis indicated the importance of these two techniques in the quantification of the material properties and failure points making them a useful tool in the research process as well as in the commercial manufacture.
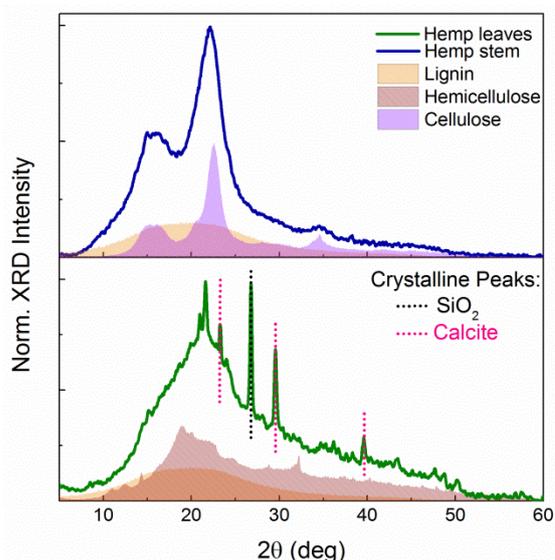

Figure 1. XRD diffraction patterns of feedstock obtained using the Rigaku Ultima Plus X-rays diffractometer placed in the AMDe Laboratory[1] of Physics Department of the Aristotle University of Thessaloniki.
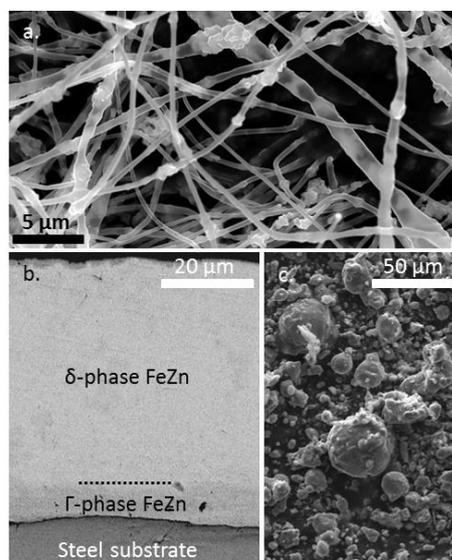

Figure 2. SEM images of a. bioactive polymeric-ceramic membranes b. anticorrosion coating (cross-section) and c. thermoelectric MnSi powder obtained using the JEOL JSM-7610F Plus Field-emission SEM recently installed in AMDe Laboratory[1].

[1] http://amdelab.physics.auth.gr/char-structure-en.htm

Chemically and thermally modified activated carbons for the removal of aromatic compounds

A. Bosacka, M. Wasilewska, D. Sternik, A. Derylo-Marczewska

Maria Curie-Skodowska University, Institute of Chemical Sciences, Maria Curie-Skoldowska Square 3, 20-031 Lublin, Poland

E-mail: alicja.bosacka@poczta.umcs.lublin.pl

Water is an important substance necessary for life of many organisms (people, animals, plants), however, industry development and globalization contribute to the increase in water pollution [1]. One of the hazardous substances are aromatic organic compounds which are difficult to decompose in water, therefore, water purification from these substances is a complex matter [2-3]. Activated carbons are commonly used sorbents for water purification because of unique textural and physicochemical properties: well-developed porous structure, chemical and thermal stability or mechanical strength [4]. The sorption effectiveness of activated carbons depends on many factors: textural properties and surface chemistry of carbons and also nature, molecular weight, dissociation capability, solubility (hydrophobicity) of adsorbates. Nevertheless, the new or improved materials are still in demand. In order to improve sorption efficiency towards aromatic substances, the carbon surface functionalization is needed [5]. The type and concentration of surface functional groups influence the bonding mechanism between the sorbents and solutes and therefore the sorption efficiency. The surface acidic or basic sites affect the charge, hydrophobicity and electron density of graphene layers of and therefore differentiate the output of sorbents towards sorbed substances [6].

In this work, commercially available granulated activated carbon (GAC) was chemically and thermally modified to diverse its surface chemistry. Textural, thermal, surface and adsorption properties were examined using several analytical techniques: low-temperature nitrogen sorption, X-ray photoelectron spectroscopy (XPS), thermal analysis coupled with mass spectrometry (MS) and Fourier transform infrared spectroscopy (FT-IR) and UV-Vis spectroscopy. Moreover, kinetic and equilibrium adsorption studies revealed the potential of applicability of studied activated carbons towards selected aromatic adsorbates.

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A novel method for production of molecularly imprinted polymers with cholesterol and other biologically important molecules

G. Becskereki[1]; M. P. Pešić[2]; M. D. Todorov[2]; G. Horvai[1]; T. Ž. Verbić[2], B. Tóth[1]

[1]Department of Inorganic and Analytical Chemistry, Budapest University of Technology and Economics, Szent Gellert ter 4, H-1111, Budapest, Hungary

[2]University of Belgrade – Faculty of Chemistry, Studentski trg 12-16, 11000, Belgrade, Serbia

E-mail: becske3@gmail.com

Molecular imprinting is a widely used technique to obtain selective adsorbents. The most commonly used method for making molecularly imprinted polymers (MIP) is the non-covalent method. Generally, the template interacts with the monomers in the prepolymerization mixture, creating secondary forces (van der Walls, ionic, hydrogen bonding) between the template and monomer. The mixture also contains porogen solvent to create larger surface and more pores. After polymerization, a highly crosslinked polymer is formed, with sites or cavities which are selective to the template and to similar molecules.

A novel method has been successfully tested for non-covalent imprinting [1]. The polymers were created in a simpler way than usual MIPs, skipping the step of the formation of prepolymerization complexes. The template is cholesterol, and no functional monomer is used. The prepolymerization mixture contains only an acrylic diester crosslinker and the porogen solvent. In this mixture ionic interactions are unlikely, and the porogen isopropanol prevents even hydrogen bonding between the template and the monomer. Despite of these apparently inadequate conditions, significant imprinting factor and improved selectivity were measured for cholesterol against some other steroids. Selective MIPs have been also created with this new method using biologically important template molecules such as ketoprofene, dibenzylamine and others. The obtained imprinting factor and selectivity improvement are remarkable even when compared to other non-covalent imprinting systems which are based on functional monomers.

The development of this new procedure and other studies have shown that selectivity of MIPs can be complicated [2]. Various experiments were carried out on a well-known, highly selective, theophylline imprinted polymer [3]. Results show, that selectivity of the MIPs might depend on the type of the analytical method where the MIP is used. This means that, the same MIP under the same condition can show different selectivity values in different experiments, for example in "radioligand binding assays", QCM sensors, electrochemical sensors or as chromatographic stationary phase.

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Polymer Inclusion Membranes as extraction device for isolation pharmaceuticals and dental components from water samples

K. Bielicka-Daszkiewicz, K. Poniedziałek , K. Lipińska

Institute of Chemical Technology and Engineering, Poznań University of Technology, Berdychowo 4, 60-965 Poznań, Poland

E-mail: Katarzyna.Bielicka-Daszkiewicz@put.poznan.pl

Polymer Inclusion Membranes (PIM) are popular in molecular transport in different solution. Most often they are used in transport and removing of ionic compounds, for example metal ions or organic ionic compounds. In this work membranes were used as extraction device for isolation organic compounds from water samples. Polymeric Inclusion Membranes (PIM) were composed of cellulose triacetate (CTA) as polymeric matrix, 2-NPOE was used as plasticizer and Aliquat-336 was used as carrier substance in different percentage. The influence of carrier and plasticizer presence on PIM's extraction efficiency was described. This extraction device was used for isolation ibuprofen, ascorbic acid and some compounds released from dental materials: camphor, dimethyl succinate (DBE-4), dimethyl glutarate (DBE-5), dimethyl adipate (DBE-6), camphor quinone (CQ), phenol, 2-hydroxyethyl metacrylate (HEMA), ethyl 4-(dimethyloamino) benzoate (EDMAB), 4-methoxyphenol (4-MP), triethylene glycol dimethacrylate (TEGDMA). These compounds were isolated from water samples. The best extraction efficiency of polymer membranes was achieved between 80-100% for three components membrane. The extraction ability of polymeric membranes was described and confirmed using Hansen Solubility Parameter determined for each analyte and membrane components. Determination of this parameter allows to describe the interaction between analytes and membrane, and conclude which membrane composition gives the best extraction properties.

The higher extraction recovery (near 100%) was achieved on three-component membrane composed of polymeric matrix, plasticizer and carrier. The results especially show the important role of carrier in the extraction process. This component allows proper adsorption and desorption of analytes. The Hansen Solubility Parameter was used for the description of mutual interaction between analytes and membranes components. Using this parameter the dispersive, hydrogen bonding and polar interaction were determined. Results show that this parameter is very suitable for description the interaction and allows to explain adsorption and desorption degree of individual analyte. This parameter also allows to predict the extraction properties of the extraction material in relation to a given analytes.

Micromagnetic analysis as a way to evaluate physical properties of magnetic nanoparticles in biomedical applications

N. Maniotis[1,2], D. Serantes[3]

[1] Physics Department, Aristotle University of Thessaloniki, Thessaloniki, 54124, Greece
[2] MagnaCharta, Center for Interdisciplinary Research and Innovation (CIRI-AUTH), 57001 Thessaloniki, Greece
[3] Applied Physics Department and IIT, Universidade de Santiago de Compostela, Compostela 15782, Spain

E-mail: nimaniot@physics.auth.gr

Magnetic particle hyperthermia (MPH) is a novel, minimally invasive, therapeutic modality, used as a cancer treatment, that employs a magnetic fluid (also termed ferrofluid) as the heating source. A magnetic fluid is a stable colloidal suspension of magnetic nanoparticles (MNPs) that can be injected directly into the tumor or delivered to the tumor via passive or active targeting upon intravenous administration. Once accumulated to the tumor area, MNPs are exposed to an external alternating magnetic field (AMF) that causes reversal of their magnetic moments, activating mechanisms of energy deposition in the form of heat [1]. The key measure used for characterizing the heating efficiency of nanoparticles in MPH is the specific loss power (SLP), which may be derived from the magnetic hysteresis loop area. In this work, the issue on whether dynamic magnetic properties of a poly-dispersed ferrofluid, modelled using micromagnetic simulations, can be extrapolated to analyse SLP values acquired at high frequencies of excitation fields is addressed. Micromagnetic finite difference simulations were performed by using OOMMF software package to obtain the dynamic hysteresis loops under a 24 kA/m AMF amplitude and for various frequencies (50-765 kHz). In OOMMF, finite difference method was used to find the solution of nonlinear Landau-Lifshitz Gilbert (LLG) equation which describes the precessional motion of nanoparticles magnetization when applying an effective magnetic field [2]. To create a system of randomly oriented nanoparticles having a certain volume fraction, 0.04%, that corresponds to a ferrofluid concentration of 2 mg/mL, we start with a perfect simple cubic lattice with a large lattice spacing so that the particle-particle distance is large enough to neglect dipolar interactions (non-interacting MNPs). System under study is a set of spherical magnetite nanoparticles of 30 nm with a lognormal size distribution. When the frequency of the applied magnetic field, f, was below ~300 kHz, SLP(f) followed an exponential increase trend. At this regime, the hysteresis area increases with f due to the increase of the phase delay in the magnetization response. After a critical f, MNPs magnetization reaches the maximum phase delay (maximum relaxation time of spins) and so any increase in f leaves unaffected the hysteresis area. To validate our approach, we employed a coupled electromagnetic-thermal model, based on COMSOL Multiphysics simulations that provides an accurate estimation of magnetic field and temperature distribution within the ferrofluid. The time-dependent temperature curves are obtained after 30 min. of MPH treatment for the same AMF amplitude used in OOMMF simulations (24 kA/m) and for two representative f values. One at low (200 kHz) and one at high (765 kHz) frequency regime. In each case, the MNPs volumetric power dissipation, imported in heat transfer model to find the temperature curves, is taken equal to the corresponding SLP(f) value. The numerical MPH curves were found in excellent agreement with the experimental data that observed under the same conditions.

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Thermal analysis techniques: Modern trends and applications

<u>E. Tarani</u>, K. Chrissafis

Laboratory of Advanced Materials & Devices, School of Physics, Aristotle University of Thessaloniki, GR-54 124, Thessaloniki, Greece

E-mail: etarani@physics.auth.gr

Thermal analysis (TA) plays an important part in our everyday life. Without thermal analysis, we would not have cars, PET bottles, wind power stations, or modern pharmaceutical products. TA refers to a set of procedures for determining the physical/chemical properties of a sample as a function of temperature or time during heating, cooling, or held at a constant temperature [1]. The TA techniques find strong applications in polymers, composite materials, metals, ceramics, foods, etc. In this study, it will be presented Differential Scanning Calorimetry (DSC) and Thermogravimetric Analysis (TGA), as well as the area in which TA techniques find strong applications. In detail, DSC is a thermal analysis technique that measures the energy difference in a sample and a reference material using a temperature-controlled programming [1]. A control system immediately increases the energy supplied to the sample or the reference depending on the type-reaction of the process, endothermic or exothermic. The equipment keeps both the sample and the reference at the same temperature. The DSC curve is presented in terms of heat flow versus temperature or time. Thermal transitions as a function of temperature and time provide quantitative and qualitative information on physical (and chemical) changes such as melting, crystallization, recrystallization, glass transition temperatures, cold crystallization, polymerization, degradation reactions, volatilization, or changes in heat capacity [1]. TGA is a technique in which the mass of the sample is measured isothermally or dynamic as a function of time and temperature, respectively, under a specific temperature program. The changes in sample weight (mass) can be caused by changes in chemical or physical properties. The most TGA instruments operate from ambient temperature to 1400 ºC with a purge gas which flows through the balance can create an inert atmosphere (N_2, Ar, He) or an oxidizing atmosphere (O_2, air). TGA is a technique for determining the thermal stability of solids and liquids, mass losses resulting from physical and chemical changes, quantitation of volatiles/moisture, vaporization, sublimation, deformulation/failure analysis, and residue/filler content [1].
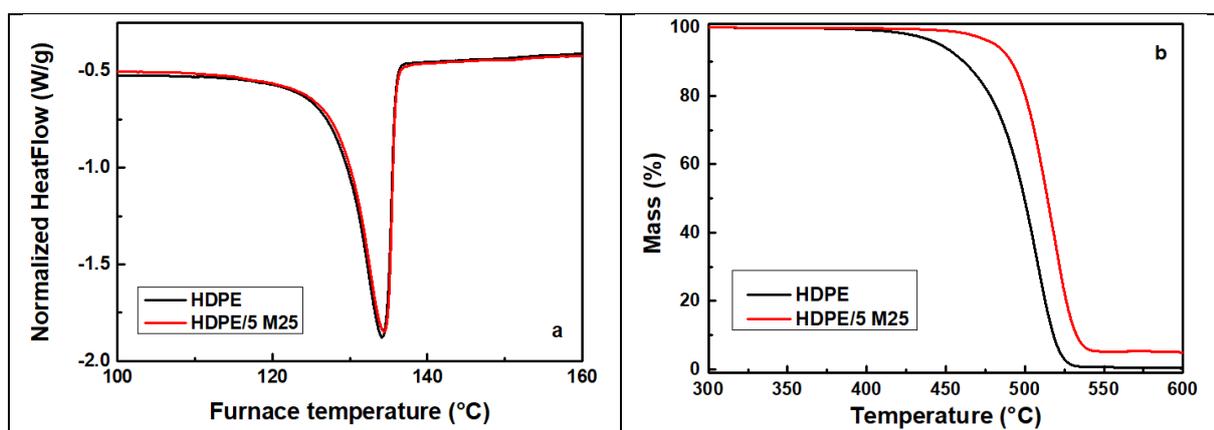

Figure 1. (a) DSC and (b) TGA thermograms for HDPE and its nanocomposite.

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Application of different analytical methods
for characterization of pharmaceutical materials

K. Milczewska, J. Maziarczyk, A. Borkowska

Poznan University of Technology, Institute of Chemical Technology and Engineering, Berdychowo 4, 60-965 Poznań, Poland

E-mail: Kasylda.Milczewska@put.poznan.pl

Many of the analytical methods can be used to characterization of pharmaceutical materials. Some of them were used to describe hybrid organic-inorganic materials applied in sustained drug delivery composed of Ibuprofen as a model drug, PEG 10000 as polymer and two different fillers (AEROSIL® 200V and AEROSIL® R816). Influence of the type of employed filler and the temperature of experiment on the surface properties and bulk interactions was estimated.

Inverse gas chromatography (IGC) has been applied to observe the interactions in that systems [1-3]. Flory-Huggins interaction parameters, the dispersive component of surface free energy and acid-base characteristic of the surface were used to assess the behaviour of the composites in terms of drug release. The temperature affects the magnitude of the interactions and thus the release of the drug from the hybrid matrix.

FTIR spectra were collected for all investigated materials and their components. Understanding the IR spectra can supply detailed information about the compound structure, as it is not possible for two compounds to have identical spectra [4].

The quantity of the released drug from the hybrid materials was measured using UV/VIS spectrophotometer [4].

The executed experiments allowed the estimation of the properties of prepared composites. Infrared tests confirmed the formation of hydrogen bond. Drug delivery profiles demonstrated explicitly that the use of a polymer in the composition decreases the initial high concentrations of the released drug. The type of the polymer has great influence on the cumulative percentage of the drug released.

Prepared materials present properties required in sustained drug release and may be successfully applied as drug delivery systems.

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SESSION D2

Electrochemistry | Sensors | Archaeometry

Chair: J. Barek, A. Economou

The use of polyaniline nanofibers for the construction of ion-selective electrodes with solid contact

K. Pietrzak[1], C. Wardak[1], S. Malinowski[2],M. Golonka[1]

[1]Maria Curie-Sklodowska University, Faculty of Chemistry, Institute of Chemical Sciences, Department of Analytical Chemistry, Maria Curie-Sklodowska Sq. 3, 20-031 Lublin, Poland
[2] Lublin University of Technology, Faculty of Civil Engineering and Architecture, Nadbystrzycka 40, 20-618 Lublin, Poland

E-mail: karolina.pietrzak@poczta.umcs.lublin.pl

Potentiometric methods characterized by low equipment costs, simplicity and speed of measurements as well as high selectivity and low limits of quantification with the simultaneous lack of necessity or minimal preparation of samples are very popular, especially in the study of environmental samples [1].

Currently, most of the research in this field is focused on the construction of electrodes with solid contact (SCISEs), in which the lack of an internal solution significantly facilitated the transport and storage of sensors, improved their mechanical resistance, and allowed their miniaturization and shape change [2].

The most important element of ion-selective electrodes is an ion-selective membrane containing a properly selected active ingredient (ionophore), which enables the sensor to work properly. Its qualitative and quantitative composition is optimized in order to obtain the best analytical parameters of the electrodes, however, its direct application to the surface of a solid electrode creates a blockage at the interface of adjacent phases. In order to obtain a stable and reversible electrode potential, it is necessary to use a transducer with electron-ion conductivity, which facilitates the flow of charge between the material of the electrode and the ion-selective membrane [3]. Carbon nanomaterials (graphene, fullerenes, carbon nanotubes or nanofibers) and various types of metal nanoparticles are used as such a transducers till now.

The construction and potentiometric tests of ion-selective electrodes with solid contact sensitive to nitrate (V) ions were described. As a solid contact, polyaniline nanofibers modified with chloride and nitrate ions were used. In order to study the influence of the presence of nanofibers number of types of electrodes were made, differing in their location and content, and then the analytical parameter values obtained for them were compared. First of all, the slope and linearity range of the calibration curves and their limits of detection were determined. Also their selectivity and stability as well as potential reversibility were examined.

It was found that polyaniline nanofibers based ion-selective electrodes were exhibited improved parameters compered to unmodified ones.

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Lead Film Electrode for the Determination of Ultratrace Aluminum by Adsorptive Stripping Voltammetry

J. Wasag[1], M. Grabarczyk[2]

[1]Department of Materials Engineering, Institute of Engineering and Technical Sciences, Faculty of Natural Sciences and Health, The John Paul II Catholic University of Lublin, 20-950 Lublin, Poland
[2]Department of Analytical Chemistry, Institute of Chemical Sciences, Faculty of Chemistry, Maria Curie-Sklodowska University, 20-031 Lublin, Poland

E-mail: joannawasag@kul.pl

Aluminum is one of the most abundant elements on earth (third after oxygen and silicon) and is present in the earth's crust at 7.5%. Does not exist in the free state. In its bound state, it is mostly in the form of aluminosilicates (aluminum salts of silicic acid), which are the main mass of the earth's crust [1]. Pure aluminum is used in the manufacture of many consumer products, for the production of electrical cables, which successfully used instead of copper. It is also used for smelting hard-melting metals from their oxides, as well as for the production of chemical equipment or coating with a protective layer of other metals. In the powder form as a protective paint, and for the production of explosives and organoaluminium materials. Aluminum is mainly used in the form of light alloys as an excellent construction material in all areas of industry [2]. In this way, aluminum can get into the environment, contaminating it, hence great emphasis is put on the monitoring of this element. Stripping voltammetry is a powerful technique for the determination of trace various metals ions. The adsorptive stripping voltammetry (AdSV) is preferred for the accumulation of the metal on the electrode surface. This is highly sensitive method because it is based on adsorptive pre-concentration of a complex of the determined metals ions. In literature data AdSV procedures for determination of trace aluminum concentration are described with the use of different working electrodes but mainly toxic mercury electrodes. In this work a procedure for the determination of aluminum was developed using lead film electrode (PbFE).

These method allowed to get very low detection limit and an excellent linear dynamic range. The cupferron was used as a complexing agent and PbFE as a working electrode. The obtained detection limit was 3.3×10^{-11} mol L^{-1} with accumulation time of 60 s. The linear calibration plot was obtained in the range from 1×10^{-10} mol L^{-1} to 2×10^{-7} mol L^{-1} and obey the equation $y = 0.156x + 0.084$, where y and x are the peak current (μA) and Al(III) concentration (nmol L^{-1}), respectively. The analytical application of this work was to enabling the determination of ultratrace Al(III) introduced into the environment as a result of corrosion of aluminum alloys. The solutions were tested after exposure to aluminum alloy AA2024 test panels. The influence of foreign ions on the measurements was also examined by selecting ions based on the composition of using aluminum alloy. It was dictated by the fact that they can accompany Al(III) being determined in the synthetic samples used in the measurements. The method seems to be promising to be used in the measurement of real samples.

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PCR-free Hepatitis B Virus determination by Electrogenerated Chemiluminescence

P. Nikolaou[1], S. Petralia[2], A. Zanut[1,3], G. Valenti[1], S. Conoci[4], L. Prodi[1], F. Paolucci[1]

[1] Department of Chemistry "Giacomo Ciamician", University of Bologna, Via Selmi 2 40216, Bologna, Italy.
[2] Dipartimento di Scienze del Farmaco e della Salute, University of Catania, Viale Andrea Doria, 6 - 95125 Catania, Italy
[3] Department of Chemical and Biomolecular Engineering, New York University, 6 Metrotech, Brooklyn, NY, 11201 USA.
[4] Dipartimento di Scienze Chimiche, Biologiche, Farmaceutiche ed Ambientali, University of Messina, Viale Ferdinando Stagno d'Alcontres, 31 - 98168 Messina, Italy

E-mail: Pavlos.nikolaou2@unibo.it

Electrochemiluminescence or Electrogenerated Chemiluminescence (ECL) is an electrochemical technique with a simple instrumentation which is based on a luminescent phenomenon, produced from an electron-transfer reaction taking place on the surface of the electrode and yielding light-emitting excited states. The most know luminophores/coreactant system in aqueous media is tris(2,2'-bipyridine) ruthenium(II) ($[Ru(bpy)_3]^{2+}$)/tri-n-propylamine with important analytical application in commercial assays for the detection of biomarkers. [1,2] According to World Health Organization (WHO) Hepatitis B Virus (HBV) is the main problem of the infection of 257 million people. [3] An ECL-based and also PCR-free sensor is developed for the determination of the double-standed DNA of HBV (ds-DNA). The ds-DNA of HBV has been immobilized and remained anchored between two monoclonal oligonucleotide chains probes (P1 and P2), thus the same time the triplex formation is created via Hoogstern H-bonds (Figure 1a). [4] The ECL-active molecule $[Ru(phen)_2dppz]^{2+}$ it can be remained intercalated in the ds-DNA according to the π-π^* stalking intercalation -dppz. [5] The ECL-based sensor is tested in a range from 0-10000 copies mL^{-1} obtaining a limit of detection of 2.7 copies mL^{-1} which is much more sensitive, comparing it with other techniques.
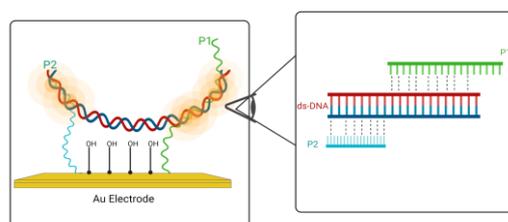

Figure 1: Immobilization of specific probes and triplex formation between ds-DNA of HBV and P1, P2 probes.

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Sensitive voltammetric approaches for the total isopropylmethylphenols quantification in plant material

G. Ziyatdinova, N. Chernousova

Analytical Chemistry Department, Kazan Federal University, Kremleyevskaya, 18, Kazan, 420008, Russian Federation

E-mail: Ziyatdinovag@mail.ru

Isopropylmethylphenols (carvacrol and thymol) are a natural phenolic monoterpenoids isolated from aromatic plants *Origanum vulgare*, *Thymus vulgaris*, *Lepidium flavum*, *Citrus aurantium bergamia* and etc. [1]. These plants are widely used as spices for cooking and can be considered as one of the sources of antioxidants in human diet. Furthermore, isopropylmethylphenols have antibacterial, antifungal, insecticidal, and anti-oxidative properties [2-4] promoting the wide use of these compounds in the cosmetic, food, and pharmaceutical industries. Thus, the control of isopropylphenols concentration in real samples is of practical interest. Isopropylmethylphenols are electrochemically active and detectable using electrochemical methods. Nevertheless, thymol and carvacrol are less studied in comparison to other natural phenolics. Being isomers, both compounds are oxidized at almost the same potentials. Therefore, total isopropylmethylphenols contents are usually quantified.

Sensitive voltammetric methods for the isopropylmethylphenols quantification have been developed using chemically modified electrodes. Multi-walled carbon nanotubes (MWNTs) and their layer-by-layer combination with electropolymerized thymolphthalein have been used as electrode surface modifier. Application of MWNTs significantly increases the effective surface area and conductivity. Polymeric film provides structural similarity to the analytes and the porous structure leading to the increase of the determination sensitivity.

The surface of the electrodes has been characterized with SEM, cyclic voltammetry, chronoamperometry and electrochemical impedance spectroscopy (EIS). SEM data confirm successful immobilization of the modifiers. According to the electrochemical data, the effective surface area of modified electrodes is significantly increased in comparison to GCE. The EIS data shows statistically significant decrease of the charge transfer resistance for the modified electrodes *vs*. GCE.

Thymol and carvacrol electrooxidation parameters on the electrodes developed have been found and corresponding oxidation schemes have been suggested. Differential pulse voltammetry has been applied for the quantification of isopropylmethylphenols. The analytical characteristics obtained are improved in comparison to the reported earlier for other modified electrodes. The selectivity of isopropylmethylphenols response in the presence of typical interferences and other natural phenolics is an important advantage of the methods developed. The analytical capabilities of the methods have been successfully shown on example of oregano and thyme spices. Validation to independent method has been performed.

Thus, the approaches developed are simple, reliable, cost-effective and can be applied for the routine analysis for the fast screening of the plant materials.

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Gas-phase biosensing and imaging of human body VOCs

K. Iitani[1], G. Zhang[2], K. Toma[1], T. Arakawa[1], <u>K. Mitsubayashi</u>[1,2]

[1]Institute of Biomaterials and Bioengineering, Tokyo Medical and Dental University (TMDU)
2-3-10 Kanda-Surugadai, Chiyoda-ku, Tokyo 101-0062, Japan
[2]Graduate School of Medical and Dental Sciences, Tokyo Medical and Dental University (TMDU)
2-310 Kanda-Surugadai, Chiyoda-ku, Tokyo 101-0062, Japan

E-mail: m.bdi@tmd.ac.jp

Gas-phase biosensors have been developed not only for human volatiles (acetone, methyl mercaptan, trimethylamine, ethanol, isopropanol, etc.) but also for residential harmful VOCs (formaldehyde, toluene, nicotine) causing sick-house syndrome, etc. In particular, the biof-luorometric (BF) gas sensors with an UV-LED excitation light and a PMT detector have indicated good sensitivity and gas-selectivity for target VOCs (formaldehyde, ethanol, acetaldehyde, acetone, isopropanol, etc.). In addition, the gas imaging system with enzyme immobilized mesh also has shown a spatio-temporal gas-imaging of human volatiles (i.e. ethanol, acetaldehyde, etc. after drinking). In this study, the gas phase biosensor (bio-sniffer) for breath acetone and the gas imaging system (sniff-cam) for skin ethanol in the gas phase will be demonstrated as unique non-invasive biosensing approaches.

The bio-sniffer for acetone vapor was constructed using a S-ADH (secondary alcohol dehydrogenase) reverse reaction by detecting the reducing of the NADH fluorescence intensity. The S-ADH immobilized sniffer was possible to monitor gaseous acetone (7 ppb to 20 ppm) continuously with a gas-selectivity based on the substrate specificity of S-ADH. The bio-sniffer allows us to measure the acetone concentration in exhaled breath from healthy subjects and diabetes (type I and II). The unique bio-fluorometric sniff-cam for gaseous ethanol was developed using alcohol dehydrogenase (ADH) immobilized mesh and the NADH imaging unit (UV-LED light array & highly sensitive camera), thus visualizing ethanol vapor not only in exhaled breath but also in transcutaneous gas after alcohol drinking.

The gas-phase biosensors would be useful for non-invasive measurement of some VOC biomarkers in the medical and health care fields.

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Application of WD-XRF, XRD, FT-IR, and XANES to Archaeological Pottery Samples

<u>**K.B. Dasari**</u>[1], **H. Cho**[2], **N. Lakshmana Das**[3], **T. Lee**[4], **Y.H. Yim**[2], **G.M. Sun**[1]

[1]Korea Atomic Energy Research Institute, Daejeon, 34057, Republic of Korea
[2]Korea Research Institute of Standards and Science, Daejeon, 34113, Republic of Korea
[3]GITAM (Deemed University), Visakhapatnam, 530045, India
[4]Korea Basic Science Institute, Seoul, 02841, Republic of Korea
(*Corresponding Author Email: dasari@kaeri.re.kr /gmsun@kaeri.re.kr)

Ancient cultural pottery is one of the important artifacts that reveal production technology and social changes which can be identified through characteristic mineral phases, chemical composition, and firing temperature of samples. In the present study, the elemental chemical composition and the percentages of iron phases (hematite, magnetite, and ferrous oxide) of pottery were determined using wavelength dispersive X-ray fluorescence (WD-XRF) and X-ray absorption near edge structure (XANES) techniques, respectively to understand the atmospheric conditions and color mechanisms along with approximate firing temperature using Fourier transform infrared spectroscopic (FT-IR). Five different ancient pottery samples belong to different age periods were selected for this study. FT-IR technique is applied to reveal the firing temperature of pottery samples, and FTIR spectrum of NP1 pottery is shown in Fig. 1. The bands observed at 3700, 1075 and 915 cm^{-1} and the presence of other minerals infers that the sample might have been fired around 500 °C. The Fe species in pottery was figured out by Fe XANES analysis. XANES experiment was carried at 7D XAFS beam line of Pohang accelerator laboratory (PAL) using 3 GeV X-ray energy. The Fe K edge (~7125 eV) XANES spectra were recorded in fluorescence mode using 7 elements Ge detector. Analyzed pottery varies in color from red to dark gray depending on the ratio of Fe^{2+}/Fe^{3+}. Fig. 2 shows that the Fe K-edge spectra of NP1 with red and NP2 with dark gray color of pottery. The predicted oxidation states of iron through XANES analysis are in good consistence with the actual pottery color. In addition, the minerals phases of pottery samples were determined by X-ray diffraction (XRD). The combined results from WD-XRF, FT-IR, XRD, and XANES indicate a promising approach for the investigation of manufacturing technology of ancient ceramics including firing temperature and provenance.
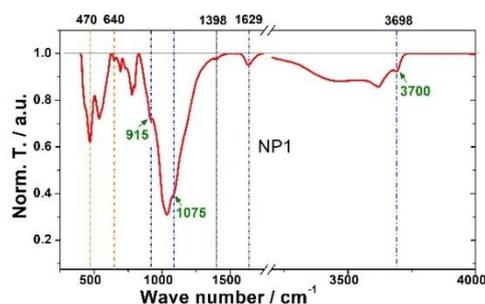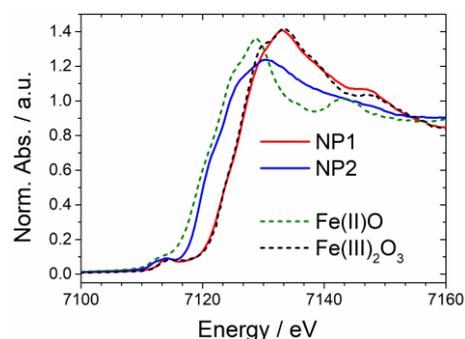

Fig. 1: FT-IR spectrum of ancient pottery

Fig. 2: Fe K-edge XANES spectra of potteries and reference compounds

Oil media on paper: The effect of linseed oil on the paper supports of works of art and printed material. A research matter of damage assessment.

P. Banou[1], K. Choulis[1], K. Tzia[2], D. Tsimogiannis[2], T. Karabotsos[1], L.A. Tsakanika[2], A. Alexopoulou[1]

[1] Department of Conservation of Antiquities and Works of Art, University of West Attica, Greece
[2] School of Chemical Engineering, National Technical University of Athens, Greece

E-mail: pbanou@yahoo.gr

Oil paintings and oil sketches on paper, prints, archival material and books present problems associated with the effect of oil medium on the paper support. Decrease of pH, discoloration, changes in opacity and embrittlement of the paper support are resulted by the absorption and diffusion of the oil medium, upon ageing. Although that research has indicated that oil mediums oxidise paper, the irregular occurrence of these problems and the variation on the extent of damage raise a complex matter of research with multi-parameter aspects. [1, 2, 3, 4]

This paper will present the methodology of a developing PhD research on the effect of linseed oil, as the principal oil medium in oil paints and printing inks, on paper. The changes caused by three types of linseed oil on the optical, morphological, mechanical and chemical properties of paper will be discussed, based on the results of research on mock-ups submitted to artificial ageing, in controlled conditions of relative humidity and temperature in airtight vessels. Research program involves colorimetry (CIELab* and diffused reflectance measurements), opacity, tensile strength and pH measurements, microscopy, SEM, non-destructive imaging testing, FTIR analysis and VOC analysis with GC-MS. The selection of the techniques used, the standards adopted, the conditions and the set-up of application will be discussed along with the results of the relevant pilot trials.

Figure 1 N. Gysis, Sewing studio (19th c.), oil sketch, National Gallery. Discoloration due to oil medium absorption is evident on the verso side.

Initial results indicate that comparative study of the results might provide secure hypothesis for the complex matter of damage of oil media on paper recorded by the conservators. However, the need of a non-destructive methodology for the evaluation of the condition of the works and the determination of stage of deterioration is regarded significant since they are associated with the preservation decision making for a wide range of cultural heritage objects. The application of non-destructive imaging techniques, digital microscopy, GC-MS analysis of VOC absorbed by a SPME needle and FTIR analysis will be discussed.

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Determining optimum irradiation conditions for analysis of heritage materials by Raman spectroscopy

A. Philippidis[1], D. Anglos[1,2]

[1]Institute of Electronic Structure and Laser, Foundation for Research and Technology-Hellas (IESL-FORTH), 700 13 Heraklion, Crete, Greece
[2]Department of Chemistry, University of Crete, 700 13 Heraklion, Crete, Greece

E-mail: filagg@iesl.forth.gr

Continuous advances in photonic technologies, have made available a broad range of spectroscopic methods based on intense radiation sources, which scientists have employed in various analytical investigations aiding archaeologists, historians and conservators in their studies and practice [1]. While most of these techniques are considered as non-destructive since no sampling is needed it is conceivable that during analysis photons or particles could potentially induce alterations on the analyzed material depending both on the nature of the substrate and the irradiation parameters. The literature considering radiation effects on Cultural Heritage (CH) objects is rather limited to date [2,3].

In this study, Raman systems with different laser excitation source, portable and laboratory, were employed for investigating samples of pigments both in pure form and in the presence of binding medium and varnish. The Raman spectrometers were used as irradiation and monitoring tool. The work concentrates on the effects of laser irradiation on inorganic pigments which are considered as common reference materials for accessing pigments discoloration effects during laser irradiation. Vermillion (α-HgS), Prussian blue ($Fe_4[Fe(CN)_6]_3$), red lead (Pb_3O_4), massicot (PbO) and cadmium pigments (CdS, CdS_xSe_y) are investigated in pure form (powder) and in different chemical environments. It is important to understand the physics and chemistry underlying light induced alterations for various paint materials. The study of different exposure parameters (laser power, exposure time, etc.) is crucial for the development of proper mitigation strategies and to establish safe irradiation boundaries and protocols for the use of Raman microscopy in art conservation.

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Thursday 23 September 2021

SESSION D3

Mobile analytical instruments | Robotics and Automation | Chemometrics

Chair: I.P. Gerothanassis, D. Anglos

Unifying Analytical Information during Multidisciplinary Research

E. Chatzitheodoridis[1], N. Tsakonas[1], E. Profitis[1], A. Thoma[1], N. Panagiotopoulos[2]

[1]National Technical University of Athens, School of Mining and Metallurgical Engineering
[2]European Space Agency, European Space Research & Technology Centre (ESTEC)

E-mail: eliasch@metal.ntua.gr

Analytical instruments of any kind (small devices, sensors, or integrated systems) still produce outputs of data that their content and structure are often proprietary. More importantly, this set of data does not contain adequate metadata for proper descriptions and comparisons. Especially in multidisciplinary sciences this is now a necessity for better describing and storing new knowledge. Contemporary efforts to standardise instrument data output and make it open access exist [1], but they are still not mature, therefore new developments are necessary. Having this necessity in mind, and within the M2M (Machine-to-Machine) framework [2] we propose to unify data that come out from instruments and methods (or algorithms) into a single object (or structure). This is not new in computer languages, however, the concept is new on handling instruments during research in the field and in the lab, where machines can perform integrated tasks that lead into the production of new knowledge, especially when machines have to operate and to communicate to each other autonomously.

We will present a new concept based also on a metalanguage, primarily to allow us to describe validated analytical data outputs from well-operating instruments that produce dynamically evolving data structures, capable of merging, exclusion, and processing, which scale up to complete pieces of knowledge. This is expected to significantly affect the way we publish new knowledge.

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Optimising the remote detection systems of alpha particle emitters using Monte Carlo simulations

C. Olaru[1], M. Zadehrafi[1], M.-R. Ioan[1], F. Krasniqi[2]

[1]Institutul Național pentru Fizică și Inginerie Nucleară Horia Hulubei (IFIN-HH), Str. Reactorului no.30, P.O.BOX MG-6, Bucharest - Magurele, Romania
[2]Physikalisch-Technische Bundesanstalt (PTB), Bundesallee 100, 38116 Braunschweig, Germany

E-mail: claudia.olaru@nipne.ro

In the context of a nuclear emergency event, such as transportation accidents, malfunction of nuclear installations or even terrorist attacks, the release of alpha-emitting radionuclides in the environment represents one of the most concerning radiological threat for human beings. This is due to alpha particles having large mass, high energy and high linear energy transfer, thus causing extreme damage to soft biological tissues if ingested or inhaled.

At present, the detection of alpha particles is done from short-distance using traditional detectors manipulated by hand, for the reason that alpha particles have a very short range in air. Therefore, in the case of a large-scale contamination, the emergency personnel would be exposed to high risk, and the measurements would be very laborious. To surmount the drawbacks of the currently used detectors, the possibility of remotely detecting alpha particle emitters in the environment is being studied and a novel optical detection system is currently under development and tested within the 19ENV02 RemoteALPHA EMPIR project [1]. One of the objectives of the project is to develop the prototype of a remote optical detection system for real-time radioluminescence mapping of alpha sources. To optimise the optical configurations of the system, Monte Carlo simulation toolkits are used for modelling the distribution of alpha particles in air and emitting volume of the resulting radioluminescence photons.

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Monitoring the response of plants and vegetation under stress via imaging spectroscopy

K. Hatzigiannakis[1], A. Vardis[2], M. Konstantinou[1,2], A. Sarris[3,4], D. Anglos[1,2]

[1] Institute of Electronic Structure and Laser, Foundation for Research and Technology-Hellas (IESL-FORTH), P.O. Box 1385, GR 711 10 , Heraklion, Crete, Greece

[2]Department of Chemistry, University of Crete, P.O. Box 2208, GR-71003 Heraklion, Crete, Greece

[3]Archaeological Research Unit (ARU), Department of History and Archaeology, University of Cyprus P.O. Box 20537, 1678 Nicosia, Cyprus

[4]Institute of Mediterranean Studies, Foundation for Research and Technology-Hellas (IMS-FORTH), Nik. Foka 130, Rethymno, Crete, Greece

E-mail: kostas@iesl.forth.gr

The use of spectroscopic sensors for the study of flora is fast advancing as it facilitates researchers to obtain valuable information about plants and vegetation, their local terrain and the environment they grow in as well as their health status since the presence of moisture and nutrients affects directly their spectroscopic response. Imaging spectroscopy plays a key-role in such studies since it allows whole field acquisition and stand-off performance. Its principle is based on the collection of the light backscattered from the plant or the plant field across sequential narrow spectral bands in the visible and the near infrared and this enables the extraction of a reflectance spectrum. Exploiting compact equipment and remote sensing schemes, spectral imaging has been widely applied in various scales ranging from land based imagers for fast small scale studies to airborne and even satellite multispectral sensors for broad area examination.

In the present study the response of leaves of various plant species in different physical conditions, from healthy to under stress due to lack of moisture, have been studied in the laboratory by means of a custom made spectral imaging camera (IRIS II). The same samples were also examined on a high spectral resolution UV-Vis-NIR spectrophotometer and a portable spot analysis diffuse reflectance spectrometer in order to correlate the data from the three instruments and certify the reliability of the acquired imaging data. Finally, results form an outdoors campaign employing the spectral imaging system and the portable spectrometer will be presented and the potential of using imaging spectroscopy for field measurements will be discussed particularly in the context of monitoring areas with shallow buried archaeological structures.

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Adaptation of Primary Source NIR Calibrations to Unlabeled Target Samples by Local Adaptive Fusion Regression and Null Augmented Regression

R.C. Spiers, <u>J.H. Kalivas</u>

Department of Chemistry, Idaho State University, Pocatello, USA

E-mail: kalijohn@isu.edu.

Multivariate calibration models using spectral data such as near infrared (NIR) spectra form accurate predictions by estimating a relationship between sample NIR spectra and analyte concentration. Complicating the use of a calibration model for prediction in real-time analysis situations is that sample spectra depend on measurement and instrument conditions such as temperature, instrument drift, manufacturer, etc., sample composition (analyte and other species amounts), and the physicochemical sample matrix effects from inter- and intra-molecular interactions. Thus, model performance degrades when target conditions are different from the original source calibration conditions. Unfortunately, this situation is too common. Presented are two methods that ensure the natural hidden variables making up target samples are being appropriately modeled to provide accurate predictions. Both processes permit on demand modeling.

One approach is considered local modeling where traditionally a subset of samples is selected from a large reference analyte library (spanning a substantial diversity of matrix effects) to form a linear model specific to predict analyte content of a target sample. However, this approach with current local modeling methods often fails because it is wrongly assumed that selecting spectrally similar samples yields calibration samples matched with respect to all hidden matrix effects. Rather than forming one local model, the presented approach, termed local adaptive fusion regression (LAFR), forms hundreds of linear local models. Thus, LAFR considers local modeling as a classification problem where each target sample is classified into one of the hundreds of linear calibration sets according to respective hidden matrix effects. There are four parts to LAFR: (I) library searching by a unique fusion approach to decimate a large library into a reasonably-sized library spectrally similar to the target sample, (II) breaking up the smaller library into linear clusters (calibration sets) of distinctive hidden matrix effects using our indicator of system uniqueness (ISU) hybrid fusion algorithm, (III) classification of a target sample into a calibration set by another ISU hybrid fusion process using over a hundred similarity measures that are extended up to thousands using our novel cross-modeling procedure, and (IV) prediction of the target sample analyte content by the selected calibration set. Results from multiple NIR datasets are presented demonstrating effective characterization of calibration and target sample matrix effects by LAFR and great improvement over global models in difficult big data type soil libraries with nearly 100,000 samples.

The second approach, termed null augmented regression (NAR), involves model updating (transfer learning) that forms a new model orthogonal to the spectral matrix effect differences between source and the unlabeled target samples to be predicted. An impediment to real world application of all model updating methods is model selection, especially in the case of adapting a model to new unlabeled samples. Presented is an automatic model selection process based on evaluating model diversity and prediction similarity (MDPS). Results for several NIR data sets are presented showing that MDPS selects reliable updated NAR models that either outperform or rival prediction errors from total recalibrations that necessitate laboratory analysis of samples to obtain the corresponding calibration sample reference values.

Tuesday 21 September 2021

POSTER SESSION 1

Aerosol Metrology | Advanced X-Ray Techniques Chromatography-Mass | Environment | Food/Proteomics | Sample SpectometryArchaeometry

Chair: D. Lambropoulou, B. Beckhoff

Magnetic nanoparticles an indicator of health risks related to anthropogenic airborne particulate matter

C. Martinez-Boubeta[1], K. Simeonidis[2], D. Sarigiannis[2], M. Kermenidou[2], Ll. Balcells[3]

[1] Ecoresources P.C., Giannitson-Santaroza Str. 15-17, 54627 Thessaloniki, Greece
[2] Department of Chemical Engineering, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[3] Institut de Cieencia de Materials de Barcelona, CSIC, 08193 Bellaterra, Spain

E-mail: cboubeta@gmail.com

A large body of literature has consistently shown an association between particulate matter (PM) pollution and rises in the numbers of deaths from cardiopulmonary disease, especially among older people and those with comorbidities [1]. Very recently, the possibility of airborne magnetic nanoparticles to enter the brain directly by crossing the olfactory unit has been demonstrated suggesting a potential hazard for the health of the aging human due to urban air pollution [2]. This work combines magnetic and microscopic chemical methods in order to provide insights into the impact of daily and seasonal life patterns on atmospheric pollution level in Thessaloniki, the second largest city of Greece. The amount of magnetically-responding dust collected in pumped-air filters was at least 0.5 % wt., although it shows seasonal maxima during autumn months (0.8 % wt.) when an increase in commuting occurs, reaching levels 50% higher in downtown than in suburban areas [3]. In combination with high-resolution transmission electron imaging and elemental analysis, it was possible to conclude that Fe_3O_4 and similar ferrites -some of them including heavy metals- with nuclei sizes of approximately 15 nm, are the dominant magnetic phases. Importantly, nasal cytologic samples collected from residents of these areas showed exactly the same magnetic behavior, thus, verifying the critical role of nanosized magnetic particles into the evaluation of air pollution threats. In this framework, we end by exploring whether seasonal variations of PM has positive causal effects on coronavirus mortality. Our results underscore that it is possible, if not likely, that pollution nanoparticles Granger causes COVID-19 death rates ($p < 0.05$).

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Application of X-ray Absorption Fine Structure Spectroscopy for the study of human nails

I Karagiannis[1], M Katsikini[1], F Pinakidou[1], E.C. Paloura[1], K Manika[2], D Papakosta[2]

[1]School of Physics, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece

[2]Pulmonary Department, Aristotle University of Thessaloniki, "G. Papanikolaou" Hospital, Exochi, Thessaloniki, Greece

E-mail: iorkarag@physics.auth.gr

Extended X-ray Absorption Fine Structure (EXAFS) spectroscopy is applied for the study of human nails. The studied nails were obtained from healthy donors and donors affected by lung cancer hospitalized in the University Pulmonary Clinic of the G. Papanikolaou hospital after their consent. Human nails contain in small amounts essential elements, e.g. Fe and Zn, whose excess or deficiency from the normal values has been related with disorders or diseases [1]. In order to determine the bonding geometry of Zn atoms in the nails, EXAFS spectra were recorded at the Zn-K-edge, at the BESSY-II storage ring of the Helmholtz Zentrum Berlin. Zn is found tetrahedrally coordinated with S and N or S and O atoms. Fitting of the spectra in the first nearest neighboring shell, in the k- and R-space, is shown in Fig. 1. Due to the small difference in the atomic number of N and O, the fitting yielded similar results and we could not discriminate differences due to bonding of Zn with either S and N or S and O. However, in both cases it was found that Zn is bonded to O or N atoms belonging to an aminoacid (most probably glutamic acid or histidine, respectively) contained in keratin and to S found in cysteine that is abundant in keratin. Therefore, Zn is boned to keratin playing a structural role in human nails. As it is evident from the Fourier Transforms (Fig. 1), variations are observed in the second nearest neighboring shell (3-4 Å). Simulation of the spectra using specific amino acids bonded to Zn is in progress. Finally, considerable variations in the number of S atoms bonded to Zn were found among the samples, which however are not related to the health condition of the donor.
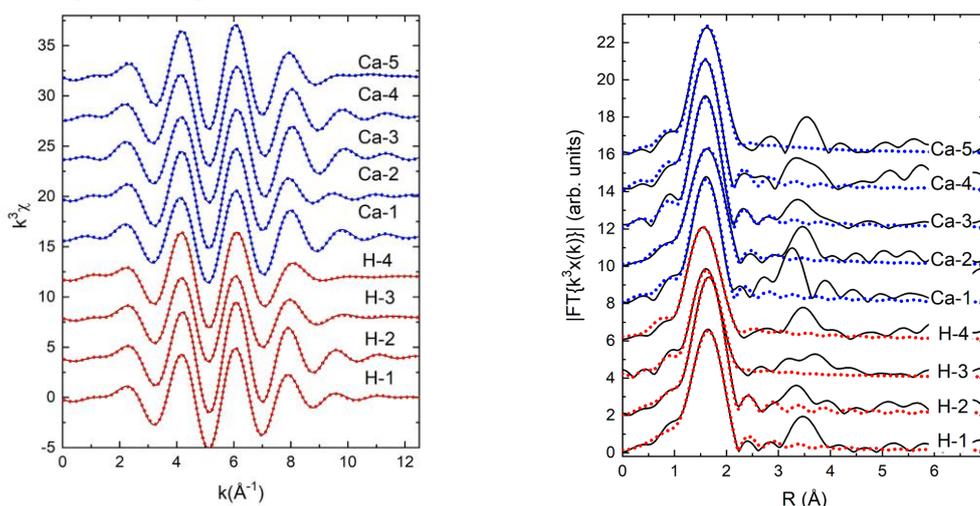

Figure 1: Fitting of Zn-K-edge filtered $\chi(k)$ spectra (left) and their Fourier Transforms (right) using $Zn - N\ and\ Zn - S$ photoelectron scattering paths. The experimental curve and the fitting are shown in solid and dotted lines, respectively. H-i and Ca-i correspond to nails from healthy donors and donors affected by lung cancer, respectively.

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X-Ray Diffraction analysis for semi-quantitative identification of important mineral phases in mining wastes by-products valorisation

E. Pagona[1], K. Kalaitzidou[1], X. Ntampou[1], E. Tzamos[2,3,4], K. Simeonidis[4,5], A. Zouboulis[2], M. Mitrakas[1]

[1] Department of Chemical Engineering, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[2] Department of Chemistry, School of Sciences, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[3] R&D Department, North Aegean Slops SA, 54627 Thessaloniki, Greece
[4] EcoResources PC, 54627 Thessaloniki, Greece
[5] Department of Physics, Aristotle University of Thessaloniki, 54124, Thessaloniki, Greece

e-mail: mmitraka@auth.gr

For the valorisation of the mining by-products/wastes of the Gerakini magnesite mine (Grecian Magnesite SA) into refractory products, X-Ray Diffraction analysis coupled with chemical analysis was applied, resulting to the semi-quantitative identification of the major mineral phases present and the qualitative evaluation of representative samples. The examined samples were selected from piles of mining by-products, which have been produced and stacked during the many years of operation of the magnesite enrichment plant. Samples are mainly ultramafic rocks (namely dunite), which have been gradually altered, due to serpentinization and other alteration processes and therefore, they have been partially or completely transformed into serpentinites, with a consequent degradation of their refractory properties. Aiming to "reverse" the serpentinization process and therefore upgrade the quality of these wastes, representative samples from the existing old, stocked piles of the mining wastes were handpicked. Samples were thermally treated at temperatures of 650° C, 850° C, 1300° C and 1600° C and afterwards studied for the determination of their initial and after-treatment mineral phases content by the application of X-Ray Diffraction analysis. As expected, XRD results revealed the qualitative and semi-quantitative identification of the major mineral phases of the samples, which are both considered important factors for the investigation of geochemical and thermodynamic parameters that influence the process of alteration (mainly through serpentinization), aiming to develop an appropriate thermal treatment process for the conversion of these mining wastes into useful refractory products with added value. The results of the samples treated at 1300° C showed that there is a re-crystallization process, favouring the deformation of olivine and the further formation of pyroxenes, due to the excess of silicon content, which is considered as a negative outcome, because of the better refractory properties of olivine (forsterite), comparing to that of pyroxenes (enstatite, clinoenstatite). The mineralogical wt.% content of the samples in -desired- olivine (forsterite) was found to be higher after thermal treatment at 1600° C, when compared to: 1) the content of samples in pyroxenes (clinoenstatite) at the same treatment temperature and 2) the content of samples in olivine after thermal treatment at 1300° C. However, the quantification of mineral phases after the thermal treatment at 1600° C is rather misleading, due to the presence of high glassy/amorphous phases at this temperature, because of the melting of pyroxenes (enstatite) at 1557° C. In conclusion, the application of X-Ray Diffraction analysis can provide valuable information and is considered as a particular important tool for evaluating the mineralogical phases, aiming to upgrade the refractory characteristics and find alternative uses for mining/mineral wastes under the application of circular economy concept.

Monitoring Fracture Healing in Mice Tibia Using Raman Spectroscopy and X-ray Microtomography

P. Giannoutsou[1], A. Karavalasi[1], E. Maggiorou[1], A. Kaspiris[2], E. Mourkogianni[2], C. Kontoyannis[1,3], E. Papadimitriou[2], M. Orkoula[1]

[1]Laboratory of Instrumental Analysis, Department of Pharmacy, University of Patras, Greece
[2]Laboratory of Molecular Pharmacology, Department of Pharmacy, University of Patras, Greece
[3]Institute of Chemical Engineering Sciences, FORTH, Greece

E-mail: malbie@upatras.gr

The study aims to the observation, with time, of the healing process of a fracture induced in tibia bone. For this purpose, mice of two different strains (Black6 and sv129/b6) were used. Fractures were induced in mice tibiae at the age of 3 months and left to heal from this point onwards. The mice were sacrificed 3, 7, 14 and 21 days following crack and tibiae were removed for the study. Unfractured tibiae served as control.

Bone regeneration was assessed employing Raman spectroscopy. Spectra were recorded through a microscope from several points of the wounded area. It was shown that remineralization took place unevenly. Some spots set out to heal later than others. A low Raman mineralization index signaled onset of bone regeneration on the specific spot. A few micrometers further, regeneration delayed appearing the index value of the "old" bone. Finally, 21 days after fracture, the healing process seemed to be completed.

Fractured mice tibiae were also studied with X-ray microtomography. Bone mineral density was calculated in all cases. Surprisingly, no significant differentiation was noted which was attributed to the bulk character of the analysis.
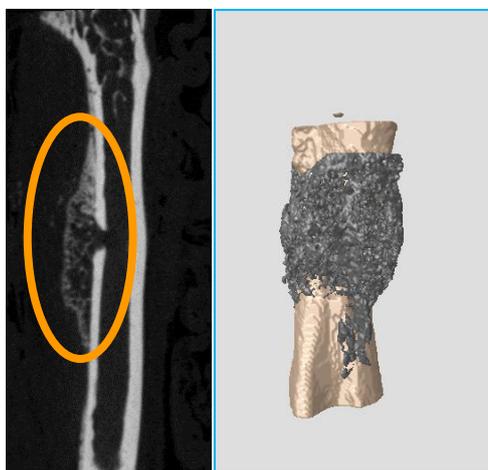
Picture. X-ray microtomographic illustrations of a fractured tibia. The crack and the callus around it are shown. 2D (left) and 3D (right) picture.

X-ray fluorescence mapping of $Gd_{0.6}Eu_{0.4}VO_4$ nanoparticles in tissues

E. Proiou[1*], F. Pinakidou[1], E. C. Paloura[1], M. Abdesselem[2], N. Pétri[2,3], T. Gacoin[3], , C. Laplace-Builhé[4], A. Alexandrou[2], M. Katsikini[1]

[1]School of Physics, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[2]Laboratoire d'Optique et Biosciences, Ecole Polytechnique, Institut Polytechnique de Paris, CNRS, INSERM, 91128 Palaiseau Cedex, France
[3]Laboratoire de Physique de la Matière Condensée, Ecole Polytechnique, Institut Polytechnique de Paris, CNRS, 91128 Palaiseau Cedex, France
[4]Photon Imaging and Flow Cytometry, CNRS, INSERM, Gustave Roussy Cancer Campus, 114, rue Edouard Vaillant, 94805 Villejuif Cedex, France

*E-mail: eproiou@physics.auth.gr

Lanthanide-ion based nanoparticles attract a lot of interest for theranostic applications due to their high luminescence efficiency, their magnetic properties, and their low cytotoxicity [1,2]. The purpose of the current work is to investigate the distribution, structure, and stability of 30 nm $Gd_{0.6}Eu_{0.4}VO_4$ nanoparticles (NPs) prior to and after the injection of their colloidal solution into biological tissue. $GdVO_4:Eu^{3+}$ NPs can be used as luminescent probes, oxidant sensors, contrast agents for Magnetic Resonance Imaging and for extracting information on the local temperature with submicrometric spatial resolution [3,4]. To visualize the spatial distribution and the mechanisms of nanoparticle diffusion in mouse ear tissues, we resorted to X-Ray Fluorescence (XRF) maps of their constituents (V, Eu, Gd) which were recorded at the BESSY-II storage ring at the Helmholtz Zentrum Berlin. Comparison of the XRF maps recorded using the emission lines of the three elements reveals that their distributions are the same indicating the integrity of the nanoparticles within the tissues. It is also observed that upon injection the NPs diffuse in the tissue following specific pathways, as shown in a representative XRF map of the Gd distribution in Fig. 1. Branching of the colloidal solution upon injection is evident. To identify variations in the nanoparticle structure, X-ray Absorption Near Edge Structure (XANES) spectra were recorded at the V K- and Eu, Gd L_3- edges prior to and after the injection in the ear tissues. The XANES results reveal that the nanoparticles do not decompose after the injection. These results pave the way for the investigation of the long-term *in vivo* fate of such nanoparticles.
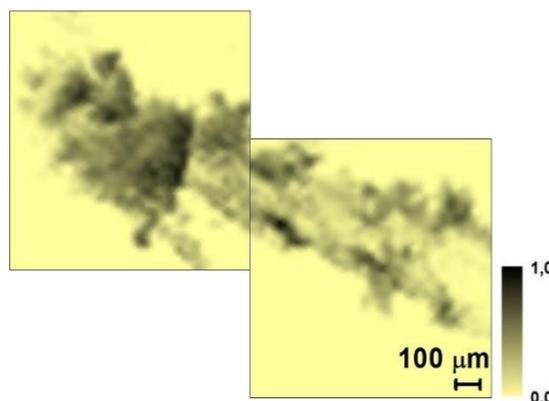

Figure 2: Gd XRF map of mouse ear tissue after injection of 10 μL of $GdVO_4$:Eu NP colloidal solution (concentration of 5 mM in vanadate ions, nanoparticle concentration of approx. 50 nM).

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Estimation of the degree of crystallinity of PLA/PHSu block copolymers with XRD and DSC

I. Chrysafi[1], D.N. Bikiaris[2]

[1]Laboratory of Advanced Materials and Devices, Department of Physics, Faculty of Sciences, Aristotle University of Thessaloniki, Thessaloniki, GR-54124, Macedonia, Greece

[2]Laboratory of Polymers Chemistry and Technology, Department of Chemistry, Faculty of Sciences, Aristotle University of Thessaloniki, GR-54124, Macedonia, Greece

E-mail: iochrysa@physics.auth.gr

Crystallinity refers to the degree of structural order of a solid. The properties of a polymer can be significantly influenced by their degree of crystallinity Xc. Moreover, the chemical structure and thermal history can alter Xc. The degree of crystallinity can be determined by several experimental techniques such as X-Ray diffraction (XRD) [1], Differential Scanning Calorimetry (DSC) [2] and Fourier Transform Infrared Spectroscopy (FTIR). In the present work, the degree of crystallinity of Poly(lactic acid) (PLA), Poly(hexylene succinate) (PHSu) and their copolymers was studied by the comparison of two techniques, XRD and DSC[3]. Polarizing light microscope (PLM) was also used to observe the crystallization process. The results come in qualitative agreement with each other. Differences of approximately 5 % arise most probably from the in principle different techniques, in addition to the two completely different methods of analysis and evaluation. Both techniques in conjunction with PLM revealed a semicrystalline morphology.
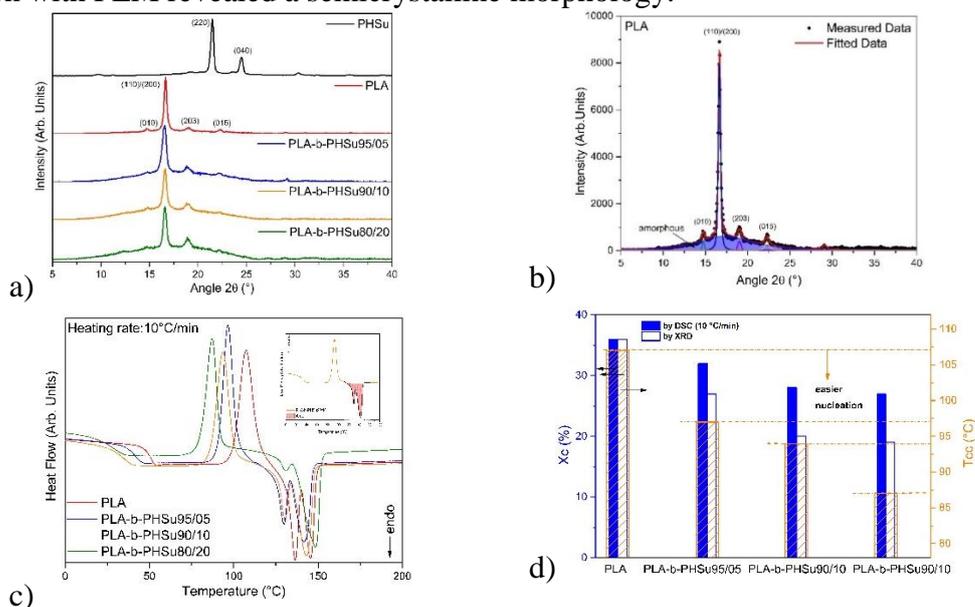

Figure 3. X-Ray Diffractograms of PHSu, PLA and their copolymers (a), Fitting profile of PLA (b), DSC heating thermograms of PLA and its copolymers(c), Xc by DSC and XRD against Tcc in the form of columns (d).

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Determination of stabilizers in nitrocellulose-based propellants before and after ageing

A. Hatziantoniou [1], M. Achilleos [1], C. Colocassidou [1], E. Ioannou- Papayianni [2]

[1] National Guard Laboratory, General Staff of National Guard
[2] State General Laboratory, P.O.Box 28648, 2081, Nicosia, Cyprus

Email: geefdyp5@cytanet.com.cy

Smokeless powder has been developed in the 1800s in order to replace black powder and is the primary propellant in civilian and military ammunition. These types of propellants are nitrocellulose-based, divided into three different categories (single, double, triple based) each category containing key additives such as stabilizers, other energetic materials, plasticizers etc. The prediction of the lifespan of propellants is of high significance not only for economical and performance but most importantly for safety reasons. High temperatures (>30°C) or high moisture content (>65%) can lead to the degradation of stabilizers which can cause chemical instability and therefore self ignition.

The National Guard Laboratory (NGL) was established in 2013 and its main purpose is to determine the stability of the propellants for the safety of civilians and military personnel. NGL uses two different techniques, Heat Flow Calorimetry (HFC) and High Performance Liquid Chromatography (HPLC) which are both validated [1]. HFC is a measure of the decomposition rate (calculated from the recorded heat flow curve) and yields information regarding the stability of propellants as well as the prediction of the lifespan [2]. Using HPLC, qualitative and quantitative determination of five initial and two daughter stabilizers present in the propellant before and after artificial ageing (the ageing of propellants is carried out artificially by HFC) is evaluated. From the results obtained separately from the abovementioned techniques is possible to predict whether the propellant is suitable for safe storage.

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Optimization of mixed-mode HPLC method intended for analysis of adrenergic drugs

B. Svrkota, J. Krmar, A. Protić, M. Zečević, B. Otašević

University of Belgrade – Faculty of pharmacy, Vojvode Stepe 450, 11221 Belgrade, Serbia

E-mail: svrkotabojana@gmail.com

One of widely used drug group refers to drugs that act through adrenergic system. In following research, the intent was to develop High Performance Liquid Chromatography (HPLC) method for separation of adrenergic drugs. Therefore, selected mixture of analytes contained bisoprolol (BP), fenoterole (FT), doxazosin (DOX), tetrahydrazoline (TH) and lofexidine (LOF). One of the advantages of this method reflects in expanding knowledge about the analytical behaviour of lofexidine, which is limited. Since all of selected analytes are basic in nature, mixed-mode column, which includes reverse phase (RP) and weak cation exchange (WCX) interactions, was considered as adequate for this HPLC analysis. The pronunciation one of two separation mechanisms depends on mobile phase content and pH, and as a consequence has the potential of selectivity modulation [1].

Analyses were performed on HPLC Thermo Acclaim Mixed Mode WCX-1 (3 μm, 2.1 x 150 mm) column. During screening phase, factors of significant influence on cationic analytes behaviour were determinated. For that reason, column temperature (30-38°C) and mobile phase composition parameters (acetonitrile content (30-50% (v/v)), pH (3.8-5.6) and ionic strength (20-40 mM) of aqueous part of mobile phase) were selected for optimization. Design of Experiments (DoE) was used for simultaneous optimization of selected factors. Experimental plan was in line with face-centered Central Composite Design. Optimization goals were set in order to adequately separate all critical peak pairs, and execute the analysis in reasonable time. Since selectivity in mixed-mode HPLC can be governed through mobile phase content, selectivity factor (α) values of critical peak pairs were set as optimization goals ($\alpha_{BP/FT}$, $\alpha_{TH/BP}$, $\alpha_{LOF/DOX} > 1.2$). The retention factor of last eluting analyte (k_{DOX}) was desired not to be greater than 10, in order to assure the rational analysis time. Experimental plan and mathematical models were obtained with Design-Expert 7.0.0 software. Obtained models were statistically evaluated (R^2, pred. R^2 and adj. $R^2 > 0.99$). The most pronounced effect on followed responses had the organic solvent content, whose increase lead to $\alpha_{TH/BP}$ enhancing, and had the opposite effect on $\alpha_{BP/FT}$, $\alpha_{LOF/DOX}$ and k_{DOX}. Higher ionic strength corresponded to better separation of BP from both FT and TH. This can be assigned to competitive behaviour between analytes and ions present in mobile phase [2]. Time analysis shortening resulted as a consequence of higher values of organic solvent, ionic strength and temperature. However, mobile phase pH had the opposite effect, which can be related to more expressed cationic interactions and greater retention of basic analytes [2].

Taking into account effects of all factors, and according to generated Derringer's desirability function for multiobjective decision making, the values of the factors that give the optimal responses are selected to be 44% of acetonitrile, ionic strength 36 mM, temperature of 38°C and mobile phase pH of 5.1.

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Development and validation of a LC-DAD-ESI-MS method for the determination of phenolic compounds of the solid residues from the essential oil industry

M. Irakli[1], P. Chatzopoulou[1], C. Biliaderis[2]

[1]Hellenic Agricultural Organization - Demeter Institute of Plant Breeding and Genetic Resources, Thermi, 57001 Thessaloniki, Greece

[2]Department of Food Science and Technology, School of Agriculture, Aristotle University of Thessaloniki, P.O. Box 235, 54124 Thessaloniki, Greece

E-mail: irakli@ipgrb.gr

Medicinal and aromatic plants play a vital role in health of people and recently they have been exploited as promising ingredients to develop novel products in sectors like pharmaceutical, cosmetic, food and pesticide industries. A feasible use can be the production of essential oils, by steam distillation processes, to obtain quality products in a traceable supply chain. However, a considerable amount of solid residues remained after their essential oil distillation, is currently unexploited, as they represent a major disposal problem for the industry concerned. Nonetheless, they are also promising sources for the recovery of health-promoting phenolic compounds which may be a profit in a sustainable way from low-cost raw material. These compounds may be used as natural antioxidants and functional food ingredients. Herein, the potential use of solid residues produced by the steam-distillation of medicinal aromatic plants as promising and unexploited bioactive matrices, is limited. Thus, the aim of the present study was to optimise and validate a simple, reproducible and sensitive LC-DAD-ESI-MS method for the simultaneous determination of 44 phenolic compounds in solid residues from the essential oil industry. These compounds were distributed in five major categories, namely, hydroxycinnamic acid derivatives (10 compounds), hydroxybenzoic acid derivatives (6 compounds), flavonoids (25 compounds), phenolic terpenes (carnasol and carnosic acid) and chalcones (arbutin).

The phenolic compounds were separated on a Poroshell 120 EC-C$_{18}$ (4.6 x 150 mm, 4 μm), thermostated at 35 °C, using a linear gradient elution system consisting of 0.1% (v/v) formic acid in water and acetonitrile, with a flow rate of 0.5 mL/min. Identification of the phenolic compounds was carried out by comparing retention times, UV and mass spectra with those of authentic standards. An analytical method validation procedure was conducted comprising linearity, inter-day and intra-day precision (repeatability), detection and quantification limits (LOD/LOQ). The validation data showed good linearity (R^2 > 0.992), precision (RSD < 15%), LOD (0.018 – 1.483 μg/mL), LOQ (0.054 – 4.494 μg/mL) and short chromatographic run (< 35 min). Finally, the method was applied to oregano (*Origanum vulgare subsp. hirtum*), rosemary (*Rosmarinus officinalis* L.) and sage (*Salvia fruticosa Mill*), before and after essential oil distillation, and more than 20 phenolic compounds were quantified in all the samples analyzed. The major phenolic compound found in oregano, rosemary and sage extracts were rosmarinic acid, while carnasol and carnosic acid were quantified in both rosemary and sage extracts in a considerable amount. As a result, it was confirmed that the proposed method was highly reliable for the determination of the phenolic species in aromatic plants extracts.

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The potential of biopartitioning micellar chromatography to predict the soil sorption of pesticide compounds

C. Stergiopoulos, F. Tsopelas, M. Ochsenkühn-Petropoulou

Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering, National Technical University of Athens, Greece

E-mail: chrisxp3@hotmail.com

Modern agricultural practices use a huge amount of pesticides to control unwanted pests. Many of such agrochemicals persist in the soil and plant systems for long time and pose a risk of migration into the drinking water sources and food chain [1]. The mobility, fate, and transformation of agrochemicals depend largely on the soil organic matter content and the hydrophobicity of compounds [2]. Chromatographyposesa powerful technique for the measurement of physicochemical parameters of organic compounds [3]. Chromatographic retention has proven to be a good surrogate to hydrophobicity measurements due to the higher accuracy and easier experimental performance [4]. The potential of Biopartitioning Micellar Chromatography (BMC) to predict the soil sorption of pesticides was investigated in this work. For this purpose, the retention of 39 structurally-diverse pesticides was determined on a Discovery RP-18 column using micellar mobile phases consisted of polyoxyethylene(23) lauryl ether (Brij35), sodium dodecyl sulfate (SDS) and cetrimonium bromide (CTAB) surfactants above critical micelle concentrations. Measured retention factors served to establish linear BMC models with soil sorption coefficient K_d and soil organic carbon sorption coefficient K_{OC} values, compiled from literature sources. Most models were improved upon inclusion of additional physicochemical parameters. The predictive performance of BMC was compared with the one of octanol-water partitioning (log P), the traditionally used physicochemical property for ecotoxicological endpoint estimation. BMC models using Brij35 and SDS surfactants were generally followed by better statistics than those derived with CTAB, while Brij35 BMC models showed the best statistics of all three surfactants, suggesting that its use mimics better partitioning processes of pesticides into soil systems. Predictions based on BMC chromatography were comparable or even superior to those based on log P.

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Monitoring of spent coffee grounds detoxification efficiency by high performance liquid chromatography

Z. Juglová, P. Diviš, H. Maňáková

Brno University of Technology, Faculty of Chemistry, Purkyňova 118, Brno 61200, Czech Republic

E-mail: xcjuglova@vutbr.cz

Coffee represents important global commodity and constitutes a significant fraction of the export economy of many countries. Coffee as brewed drink prepared from roasted coffee beans is one of the most widely consumed beverages in the world. Consumption of this beverage generates huge amounts of solid residues. An estimated percentage of around 90 % of the brewed coffee ends up though in the form of so-called Spent Coffee Grounds (SCG). With an estimated average of 11 grams of fresh ground coffee going into each cup, around 381,000 tonnes of ground coffee are brewed every year, resulting in an estimated quarter of a million tonnes of wet SCG [1]. This material is valorised according to the latest trends and a number of valuable products can be obtained from it. Fresh SCG or SCG after valorisation can also be used as a fertilizer due to the content of some elements and the large water retention capacity [2, 3]. However, the use of SCG as a fertilizer is hindered by the content of caffeine, tannins and phenolics in SCG which acts as phytotoxins.

In the presented work, we studied detoxification of SCG using oxidizing agents, so that SCG could be used for fertilizer preparation. Detoxification efficiency was tested using HPLC by determining the individual phenolic compounds content in the SCG water extract (1:10 solid-water ratio). An Agilent Infinity 1260 HPLC with DAD detector was used during the study. Stationary phase was formed by a Kinetex EVO C18 column and mobile phase consisted from the mixture of acetonitrile and water with 2 % formic acid. The results of this study proved successful detoxification of SCG using the proposed method. The concentration of individual phenolic compounds in the detoxified SCG fell from tens of mg/kg below the detection limit of HPLC (0.3 mg/kg chlorogenic acid, 0.2 mg/kg caffeic acid, 0.1 mg/kg ferulic acid and 0.6 mg/kg gallic acid). Content of caffeine in detoxified SCG fell by more than half in comparison to fresh SCG. In addition, the possibility of repeated use of an oxidizing agent for SCG detoxification was studied in this work. The results of this study proved that detoxified SCG has significant potential for use in the agricultural industry as a fertilizer.

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Determination of the aroma profile of commercial Authentic Greek strained yoghurts and yoghurt desserts using SPME coupled with GC-MS

G. Kartsiolas[*], M. Zoumpourlis, T. Massouras

Laboratory of Dairy Research, Agricultural University of Athens, Iera Odos 75, Votanikos, 11855 Athens, Greece

*E-mail: gkartsiolas@gmail.com

The aroma profile of commercial Authentic Greek strained yoghurts and yoghurt desserts (desserts with animal fat and desserts with vegetable oils), produced from cow milk, was studied. Eighty-four samples taken from the greek market, with fat contents less than 3%, between 3% and 6% and more than 7%, were used. All samples were analyzed using Solid Phase Microextraction and Gas Chromatography–Mass Spectrometry analysis the day after purchase. Analysis of Variance and Principal Component Analysis were performed to determine the significant differences in volatiles levels. A total of 53 volatile compounds were identified, among them all those considered as major aroma components of yoghurt (i.e. acetaldehyde, diacetyl, acetoin, acetone, 2-butanone, 2,3-pentanedione, acetic acid). Among carbonyl compounds acetaldehyde was the most abundant in all samples followed by diacetyl. The results showed that concentrations of acetaldehyde, diacetyl and total ketones were significantly higher ($p<0.05$) in strained yoghurts compared to yoghurt desserts. From methyl ketones 2-heptanone, 2-nonanone, 2-undecanone, and 2-dodecanone were not detected in yoghurts with vegetable oils. These ketones are related to fat content of yoghurt. Among short chain free fatty acids, acetic acid was the most abundant in all yoghurts, followed by octanoic, hexanoic and decanoic acids. Sorbic acid was found only in yoghurt desserts with vegetable oils in concentrations ranging from 8ppm to 208ppm. Ethanol was detected only in two yoghurt desserts with animal fat. The increased fat content did not seem to have a negative effect on the release of volatile substances. The observed differences in the aroma profile would be applicable to predict flavor, quality control, or shelf-life of the commercial Authentic Greek strained yoghurts and yoghurt desserts.

Development and validation of an UHPLC-qTOF-MS method for the determination of PET and PBT cyclic oligomers in food simulants

D. Diamantidou[1,2], O. Mastrogianni[3], E. Tsochatzis[4], J. A. Lopes[5], G. Theodoridis[1,2], N. Raikos[2,6], H. Gika[2,6], S. Kalogiannis[2,7]

[1]Laboratory of Analytical Chemistry, Department of Chemistry, Aristotle University of Thessaloniki, 541 24, Thessaloniki, Greece

[2]Biomic_AUTh, Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center, B1.4, Thessaloniki, 10th km Thessaloniki-Thermi Rd, P.O. Box 8318, GR 57001

[3]Laboratory of Toxicology, Forensic Service of Thessaloniki, Thessaloniki, 56334, Greece

[4]Department of Food Science, iFOOD, Centre for Innovative Food Research, Aarhus University, Agro Food Park 48, Aarhus N, 8200, Denmark

[5]European Commission, Joint Research Centre, Geel, Belgium

[6]School of Medicine, Aristotle University of Thessaloniki, Thessaloniki, 54124, Greece

[7]Department of Nutritional Sciences and Dietetics, International Hellenic University, Thessaloniki, GR-574 00, Greece

E-mail: ddiamanti@chem.auth.gr

Polyethylene terephthalate (PET) and polybutylene terephthalate (PBT) are among the most frequently used Food Contact Materials (FCMs) in beverage packaging and kitchenware, respectively. These items may end-up containing different substances with potential to migrate into food, such as, which may contain PET and PBT cyclic oligomers, formed during polymerization. The European Union (EU) legislation has not defined specific migration limits for PET cyclic oligomers[1]. But that is not the case for the PBT cyclic oligomers (dimer to pentamer), which are regulated and can be used as additives in FCMs following specific limitations[1]. However, their risk for human health still remains unclear due to the lack of toxicological data [2]. Moreover, according to the European Food Safety Authority's (EFSA) *in silico* calculation of the Threshold of Toxicological Concern (TTC), both PET and PBT cyclic oligomers are considered as Cramer III toxicity substances, indicating a potential for significant health effects [3,4,5].

The aim of our study was to develop, optimize and validate an analytical method for the quantification of cyclic PET (dimer to pentamer) and PBT (dimer to tetramer) oligomers, by UHPLC-qTOF-MS analysis and positive electrospray ionization. The separation was performed on a Waters BEH C18 (150 x 2.1 mm, 1.7µm) column under gradient elution conditions. The proposed method was validated in terms of linearity ranges, limits of detection (LODs) and quantification (LOQs), precision and accuracy. LODs ranged from 1.7 to 6.7 µg L^{-1}, while method's accuracy, expressed as percentage relative error, ranged within the acceptable values of ±0.1 and ±20.5% for the low concentrations. Finally, the method was successfully applied to the analysis of food simulants following the EFSA recommended migration testing conditions for plastic FCMs (10 days at 60°C) [1].

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Exploring zwitterionic HILIC chromatography using the newly introduced BEH Z-HILIC column for targeted and untargeted LC-MS metabolic profiling.

A. Lioupi[1], C. Virgiliou[1], H. Gika[2], G. Theodoridis[1], I. D Wilson[3], K. Smith[4]

[1]Laboratory of Analytical Chemistry, School of Chemistry, Aristotle University of Thessaloniki, 54124, Thessaloniki, Greece
[2]Laboratory of Forensic Medicine and Toxicology, School of Medicine, Aristotle University of Thessaloniki, 54124, Thessaloniki, Greece
[3]Division of Systems Medicine, Department of Metabolism, Digestion and Reproduction, Imperial College London SW7 2AZ, UK
[4]Waters Corporation, Milford, Massachusetts, USA

E-mail: liouarte@chem.auth.gr

Column selection plays a key role in LC-MS-based metabolomics studies and great attention has been given to the development of novel and improved separations in the quest to obtain the most comprehensive metabolic profiles of biological samples in a single analysis. Mixed-mode phases can offer advantages in that direction as they can provide alternative selectivities to the typical RP or HILIC separations. Here, the performance of a newly introduced zwitterionic HILIC column was explored for the chromatography of a wide range of small hydrophilic metabolites with diverse chemical properties under different mobile phase compositions and pH values. The column is designated to retain polar and charged analytes by exploiting the combination of HILIC mechanisms with anion as well as cation exchange interactions. Typical chromatography parameters such as peak shape, retention time, signal-to-noise ratio, peak asymmetry and peak width were evaluated based on triplicate injections of standard mixtures. These mixtures were formed from a total of 89 hydrophilic metabolites comprising important metabolic intermediates such as aminoacids and their derivatives, amines, organic acids, sugars, purines, pyrimidines and derivatives.

The optimum chromatographic parameters were then applied to a targeted (UHPLC-MS/MS) as well as an untargeted (UHPLC- QTOF MS) system for the analysis of liver tissue extracts from two groups of mice (control *vs* treated). The potential of the chromatographic systems to obtain a comprehensive metabolic profile and capture meaningful metabolic information was shown; the performance of the BEH Z-HILIC column regarding sensitivity of detection and metabolite coverage was assessed. It is demonstrated that the BEH Z-HILIC column can detect a wide variety of metabolites from different chemical classes and polarities utilising one-dimension separation and it was proven to have high potential for LC-MS-based metabolomics applications.

QSRR Modelling for Analytes Retention Prediction in LC/MS by Applying Different Machine Learning Algorithms

T. Liapikos[1,2], D. Kodra[1,2], K. Kademoglou[1,2], D. Diamantidou[1,2], O. Begou[1,2], C. Zisi[1,2], A. Pappa-Louisi[1], G. Theodoridis[1,2]

[1] Department of Chemistry, Aristotle University of Thessaloniki, 541 24, Thessaloniki, Greece
[2] Biomic_AUTh, Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center, B1.4, Thessaloniki, 10th km Thessaloniki-Thermi Rd, P.O. Box 8318, GR 57001

E-mail: tliapikos@chem.auth.gr

Liquid Chromatography coupled to Mass Spectrometry (LC-MS) offers high throughput analysis of the metabolome and lipidome. Efficient separation of analytes depends on their Retention Time (t_R), determined by physicochemical interactions with the stationary and mobile phase. Analyte t_R values are useful in untargeted LC-MS pipelines, as they aid in reducing false positive identifications for unknown molecules. However, such information is rarely leveraged, due to t_R inherent variability, given that t_R values largely depend on the specific chromatographic and instrumental properties applied. As a result, retention prediction methods have been developed based on the structural and physicochemical characteristics of analytes. Such methods employ regression models, harnessing machine learning algorithms mapping experimentally derived t_R analytes with various structural and physicochemical descriptors, known as quantitative structure retention relationships (QSRR) models. In this study, 12 Machine Learning (ML) Regression algorithms, namely Multiple Linear, Lasso, Ridge, Bayesian Ridge, XGradientBoost, AdaBoost, Random Forest, Partial Least Squares, Support Vector Machine (linear/non-linear), Passive Aggressive and ElasticNet, were tested for their ability to perform t_R predictions. Each algorithm was evaluated on publicly available RP LC-MS retention data sets[1,2] using various experimental setups and on experimental chromatographic data (>900 metabolites), including Merck/MSMLS library, analysed in different chromatographic columns. For each analyte data set, a total of 200 Molecular Descriptors (MDs, features) were calculated using the RDKit library (open-source ChemInformatics software). Finally, through a filtering process (e.g., removing features showing low variance among molecules or a correlation greater than a certain limit), the final set of features for each data set was determined. A number of feature subsets were defined and evaluated, based on both the importance of the features in the ML algorithm used, and on the application of a variety of Genetic Algorithm techniques. Additional QSRR models were developed by using dimensionality reduction techniques (such as Principal Component Analysis and Partial Least Squares Regression), on the original set of features. All analyses were performed using Python programming language.

Our study shows that each algorithm performance strongly depends on the dataset used, making clear the necessity for an initial performance inspection of the algorithms when dealing with new sets of data, in order to attain the best prediction results possible.

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Fabrication of a novel polydopamine-modified luffa sponge for the sorption-based extraction in sample pretreatment

P. Anagnostou, C. Stalikas, T. Chatzimitakos

Laboratory of Analytical Chemistry, Department of Chemistry, University of Ioannina, 45110 Ioannina Greece

E-mail: cstalika@uoi.gr

Luffa sponge is a natural and biodegradable material, commonly referred to as vegetable sponge. It is derived from matured dried fruit of *Luffa cylindrical* and mainly consists of cellulose, lignin and a small amount of hemicellulose. Luffa displays remarkable mechanical properties, revealing a physically strong texture and forming a natural 3D network. It is a low density, high-porosity, eco-friendly and low-cost sponge, with large specific surface area, that shows great potential to be used as a sorbent [1].

A limited number of reports have been published, trying to modify Luffa sponge and improve its sorptive capacity. In this work, the 3D porous sponge was coated with polydopamine (PDA) layers through a novel approach. As a proof-of concept, we examined the sorption performance of Luffa@PDA, for three groups of substances with environmental impact, namely, phenols, benzophenones and parabens, so that a new analytical method can be established.

Our results demonstrate that the bare luffa sponge achieves a yield of extraction to some degree, for all analytes (i.e., methylparaben, ethylparaben, propylparaben, butylparaben, 2-nitrophenol, 3-nitrophenol, 4-nitrophenol, 3-methyl-4-nitrophenol, benzophenone-2, benzophenone-3, benzophenone-8, *4 OH-benzophenone)*. However, PDA coating increases by 150% the extraction yield. It is noteworthy, that the duration of polymerization was found to be a critical parameter. Luffa sponges coated with PDA for a few hours (5-7 h) exhibited better performance compared to Luffa@PDA sponges prepared by conducting the polymerization for more hours (e.g., 16 h). Therefore, it can be inferred that not only PDA but also the non-coated functional groups of Luffa sponge are responsible for the extraction of the analytes from the aqueous sample, since more time of polymerization results in a higher coating of PDA on the surface of Luffa. The reproducibility of the extraction from different batches of Luffa@PDA (polymerized for 7 h) expressed as the relative standard deviation of five analyses was <6%.

All the above showcase that natural luffa sponge is a promising sorbent material, whose sorption properties can easily and cost-effectively be enhanced through the polymerization of dopamine.

Figure 1: Different kinds of Luffa sponge A) Bare Luffa B) Luffa@PDA

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Development of chromatographic conditions for separation of fifteen phenolic acids with the use of silica-based monolithic columns

M. Staniak, M. Wójciak, S. Dresler, J. Sawicki

Department of Analytical Chemistry, Medical University of Lublin, Chodźki 4a, 20-093 Lublin, Poland

E-mail: michalstaniak@umlub.pl

High performance liquid chromatography (HPLC) is one of the most common analytical techniques used for the analysis of plant secondary metabolites. Huge diversity of packing materials for chromatographic columns enables the selection of the appropriate column for a specific analytical purpose. Monolithic filling is characterized by the presence of macro and mesopores in the monolithic structure of silica. First generation of chromatographic columns based on monolithic silica appeared on the commercial market in 2000. Improvement of the production process allowed to introduce the second generation of monolithic columns in 2011. The main advantages of this type of filling are the possibility of using high flow rates of the mobile phase and a lower risk of column clogging, which is very important in the analysis of samples with rich matrix including plant extracts.

The aim of this study was to develop analytical method enabling the separation of a group of fifteen phenolic acids with the use of a commercially available monolithic column. The elaborated method was successfully applied for the analysis of extracts from leaves of *Hypericum perforatum* L.

Identification of malodorous compounds in a non-food product by using different chromatographic techniques

S. Zauner, E. Rosenberg

Vienna University of Technology, Institute of Chemical Technologies and Analytics, Getreidemarkt 9/164, A-1060 Vienna, Austria

E-mail: sabine.zauner@tuwien.ac.at

The acceptance of products on the market is not only strongly dependent on their quality, appearance and physico-chemical and sensoric properties but also often on their smell. The study presented here is concerned with a non-food product which is produced by polymerization that involves a thiol-ene click reaction. During the production process, an unpleasant odour is formed which leads to a low acceptance of the product on the market.

The sensitivity of the human nose often exceeds that of any detector in analytical chemistry. In order to achieve a qualification of the odour-active components in the samples concerned, headspace analysis was performed and Solid Phase Micro Extraction (SPME) was used to improve sensitivity. To increase the analytes' gas phase concentration, the samples were thermally equilibrated at 70 °C and then extracted for 10 minutes using the SPME fibre. Preliminary experiments showed that Carboxene/PDMS coating was the most suitable one for this analysis. After the enrichment the analytes are chromatographically separated, and different combinations and variations of detection methods were applied. Since odorous substances are often compounds containing heteroatoms, gas chromatography with an atomic emission detector (AED) was used in addition to qualitative GC/MS analysis. Due to the complexity of the sample with more than 140 components in the gas phase, additional GC/FID measurements with an olfactometric detection port were performed to qualify the odours that were detected.

The application of individual chromatographic methods and their combinations with various modes of detection, allowed to trace back the possible odour-causing compounds to seven substances. Five of them could be identified as sulphur-containing.

Development of a negative-thermal gradient GC for fast gas chromatography and its application for the study of volatile products formed in lithium-ion batteries

B. Klampfl, R. D. Müller, E. Rosenberg

Vienna University of Technology, Institute of Chemical Technologies and Analytics, Getreidemarkt 9/164
A-1060 Vienna, Austria

E-mail: bernhard.klampfl@tuwien.ac.at

Improvements to the longevity, energy-density, cost-efficiency and charging times of lithium-ion batteries are crucial aspects for the viability of this form of energy storage. Work is going on in many laboratories all over the world to improve LIBs by optimizing their electrolyte and establishing a deeper understanding of their degradation and stabilization mechanisms.

To investigate the degradation and aging mechanisms of the electrolyte *in-situ*, a real-time-(*operando)* method is required. Identifying and quantifying with high time resolution gas species that are formed during the use of LIBs is crucial to understand the dynamics of the process. To achieve this, a GC/MS system equipped with a fast-probing system to enable short injection intervals will be employed. Conventional temperature programmed gas chromatography has too long cycle times for this purpose, which prompted us to look for a different approach. Short columns with direct heating systems allowing for rapid heating rates appear to be a suitable solution. Once that direct heating of the GC capillary is adapted, this opens further new possibilities to improve the separation performance of this system, e.g. by using a thermal gradient in space. The potential of such a system is currently under investigation and will be discussed.

We will report here a concept and experimental set-up that has been developed and evaluated for its ability to produce a spatially resolved temperature gradient along the GC column. Heating rates of several hundred °C/min could be achieved while maintaining a gradient of at least 40 °C along the column.

Use of a negative thermal gradient should improve the resolution for broad injected solute zones [1] by focusing peaks as they move along the column. Additionally, changes in the selectivity compared to classical GC are possible, due to the higher interaction of analytes with the stationary phase at lower temperatures. The shaping of this spatially resolved gradient should also allow to account for varying volatility ranges of complex analytes. These aspects should make it a viable alternative to existing heating methods in gas chromatography. As a result, a negative-thermal gradient GC should be a feasible alternative for any *in-situ* or *on-line* analysis in processes that require a high time resolution.

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Analyte-modulated optical properties of organic and inorganic nanoparticle suspensions for biothiol sensing using consumer electronic devices as detectors

T. G. Choleva, A. Sfakianaki, A.G. Vlessidis, D.L. Giokas*

Department of Chemistry, University of Ioannina, 45110, Ioannina, Greece

E-mail: tacholeva@gmail.com

In this work we report on the development of new sensing methods that are based on the optical changes induced by biothiols in organic and inorganic gold nanoparticle assemblies. The first sensing strategy relies on the ability of thiols to inhibit the catalytic enlargement of gold nanoparticle (AuNP) seeds in the presence of ACl_4^- ions and trigger their aggregation while the other on the ability of thiols to quench the color of $AuBr_4^-$ coated cationic micelle aggregates. Both assays were performed by monitoring the color changes in the RGB color system, using a flatbed scanner operating in transmittance mode as an inexpensive microtiter plate photometer. The analytical merits of both methods were satisfactory in terms of detection limits (1-15 μM), recoveries (80-115%), and reproducibility (RSD=1-10%).
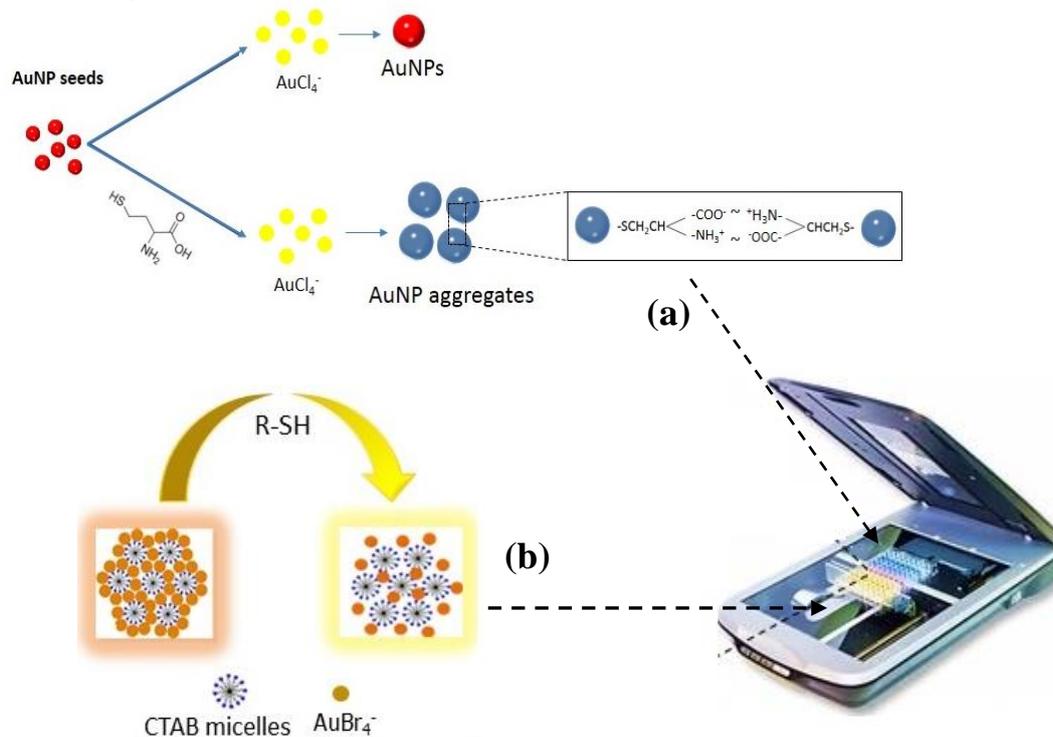
Graphical representation of the potential mechanisms of biothiol sensing based on (a) the effect of biothiols on the growth and aggregation of gold nanoparticles and (b) gold-coated CTAB micellar assemblies.

Keywords: biothiols, nanoparticles, optical detection

Breath alkanes study of the effectiveness of spirulina supplement in recovering from exercise-induced muscle damage in humans

K. Mikedi[1], P. Zeimpekis[1], S. Karma[1], P. Athanasakis[1], I. Katsoulakis[1], A. Pappa[1], A. Krokidas[2], T. Nomikos[2], M. Krokida[1]

[1] Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering, National Technical University of Athens
[2] Department of Nutrition and Dietetics, School of Health Science and Education, Harokopio University

E-mail: mikedi@chemeng.ntua.gr

Human expired air contains hundreds of Volatile Organic Compounds (VOCs) that could provide with important insights into environmental exposure, dietary habits, lifestyle, diseases and medication. Determining VOCs in breath for studying various health issues has been significantly increased since 1990. Breath analysis has certain advantages over analysis of other body liquids (blood, urine, sweat); it is non-invasive and pain-free, sample is always available, long sampling and preconcentration makes the analysis sensitive to low levels of VOCs. In the current study, breath analysis is used for studying the effectiveness of spirulina supplement in recovering from exercise induced-muscle damage (EIMD). EIMD is a pathophysiological phenomenon occurred after intense or accustomed exercise and it is characterized by the acute activation of oxidative stress and inflammation. Intake of supplements, such as, spirulina can potentially attenuate EIMD and accelerate recovery of athletes. Spirulina was selected since it is considered a well-known microalgae genus with significant content of proteins, vitamins, pigments, fatty acids, sterols, having proved health benefits and hence, athletes prefer to consume. For this work, a group of 8 healthy volunteers of mild physical activity participated in a randomized, cross-over, double-blind intervention study. The intervention started with a supplementation period of 15 days where the volunteers consumed daily 42mg/Kg body weight Spirulina Nigrita or maltodextrin as placebo. By the end of the supplementation period, the volunteers followed an eccentric exercise protocol that is able to induce muscle damage. The volunteers consumed the supplement three more days post-exercise. Samples of exhaled breath have been collected in Tedlar bags prior to supplementation (baseline), prior to exercise, and 1h, 24h, 48h and 72h post-exercise. Breath samples were analysed by Thermal Desorption-Gas Chromatography/Mass Spectrometry (TDU-GC/MS). Ergo-physiological methods and blood analysis were also used as reference methods. Analysis of the chromatographic data targeted to straight-chain alkanes and monomethylalkanes (C5-C9), which are among the oxidative stress biomarkers originated from lipid peroxidation of polyunsaturated fatty acids [1]. Preliminary results have shown a significant interindividual variability with no obvious correlation with classical markers of exercise-induced muscle damage while the kinetic are similar after the placebo and spirulina intervention. A different set of molecules are currently investigated for the ability to assess post-exercise inflammation and oxidative stress.

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Fabric phase sorptive extraction coupled to porous graphitized carbon liquid chromatography for the quantitation of parabens in human breast milk

A. Parla[1], I. Balatsouka[1], G. Mavrogeorgos[2], A. Kabir[3], V. Samanidou[4], K.G. Furton[3], D. Gennimata[5], I. Vinkovic Vrcek[6], I. Panderi[1]

[1]Laboratory of Pharmaceutical Analysis, Division of Pharmaceutical Chemistry, Faculty of Pharmacy, National and Kapodistrian University of Athens, Panepistimiopolis, Zografou, 157 71 Athens, Greece
[2]P.&A. Kyriakou Children's Hospital, Athens, Greece
[3]International Forensic Research Institute, Department of Chemistry and Biochemistry, FL, USA
[4]Laboratory of Analytical Chemistry, Department of Chemistry, Aristotle University of Thessaloniki, 54124
[5]General Hospital "Korgialenio-Benakio National Red Cross", Erithrou Stavrou 1, 115 26 Athens, Greece
[6]Institute for Medical Research and Occupational Health, Croatia.

E-mail: irenepanderi@gmail.com and anthiparla@hotmail.com

Human breast milk is the optimum source of nutrition during the first six months of life with several health and social benefits. Drug treatment during breast feeding raises several issues including the risk of exposure to drugs and chemicals passing through the breast milk to the infant. Milk is slightly more acidic than human plasma and contains significantly more lipids and fewer proteins. Thus, drugs and chemicals with low molecular weight, low plasma protein binding and high lipophilicity tend to concentrate more into the breast milk [1,2]. Parabens have been widely employed as preservatives since the 1920s for extending the shelf life of foodstuffs, medicines, and daily care products. Given the fact that there are some legitimate concerns related to their potential multiple endocrine disrupting properties, the development of novel bioanalytical methods for their biomonitoring is crucial. In recent years, the development of novel sample preparation procedures following the philosophy of Green Analytical Chemistry is a matter of growing interest among the analytical and bioanalytical scientists. To this purpose, we thought that it would be of interest to develop a method for the quantitation of parabens in human breast milk samples using a novel, eco-friendly, and efficient fabric phase sorptive extraction (FPSE) technique. In this work a fabric phase sorptive extraction-high performance liquid chromatography method (FPSE-HPLC) has been developed and validated for the quantification of parabens in human breast milk. This technique uses a natural or synthetic fabric membrane as a substrate that is chemically coated in the form of a very thin but spongy coating of sol–gel organic–inorganic hybrid sorbent. In this study, a sol–gel Carbowax® 20M coated FPSE membrane was used for sample preparation of the human breast milk samples [3]. Chromatographic separation was performed on a porous graphitized carbon analytical column which allowed the elution of the analytes within 10 min. The assay was linear over a concentration range of 20 to 500 ng mL^{-1} for all the analytes and it was applied to the analysis of parabens in real samples.

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Simultaneous quantitation of three ultraviolet filters in suncare products by high-performance liquid chromatography with photodiode-array (PDA) detection

P. Feidias[1], I. Balatsouka[1], E. Kalogria[1], S. Papageorgiou[2], A. Varvaresou[2], I. Panderi[1]

[1]Laboratory of Pharmaceutical Analysis, Division of Pharmaceutical Chemistry, Faculty of Pharmacy, National and Kapodistrian University of Athens, Panepistimiopolis, Zografou, 157 71 Athens, Greece.
[2] Laboratory of Chemistry-Biochemistry-Cosmetic Science, Department of Biomedical Sciences, University of West Attica, 122 10 Athens, Greece.

E-mail: irenepanderi@gmail.com

Sun protection is not a modern occurrence. For centuries, people have used various means to protect themselves from solar ultraviolet (UV) radiation exposure. Early on, such protection was largely achieved by clothing and wide-brim hats. Generally speaking, the optical region of the solar spectrum consists of ultraviolet (UV), visible, and infrared (IR) radiation. The UV region has the highest energy and is responsible for the damage caused to skin and hair [1]. The UV region is divided into UVA (320-400nm), UVB (290-320nm), and UVC (200-290nm). UVC, with the highest energy, is the most dangerous but is absorbed before it reaches the earth. The development of modern sunscreens began in earnest during the 1930s. Through the course of the 20th century, numerous UV filters having unique characteristics were introduced with the common property of reducing solar UV, principally short wavelengths (ie, UVB [290–320 nm]). More recently, much effort has been devoted to the development of UV filters that absorb at longer wavelengths of UV or UVA (ie, 320–400 nm). Chemical UV filters are organic molecules formed in general by one or more benzene rings and/or carbonyl groups, allowing a high electron delocalization that provides them with a high molar absorptivity in both UVA and UVB ranges [2,3].

4-Methylbenzylidene camphor (4-MBC), octyl methoxycinnamate (OMC) and avobenzone (AVO) are three chemical UV filters which are widely used in sunscreen formulations, and, to this purpose, we thought that it would be of interest to develop a method for the quantitation of these UV filters in a suncare product. Regarding the commercial sample preparation, we used methanol as diluent and the ultrasonic bath for the extraction of the UV filters. Chromatographic separation was performed on a reversed-phase phenyl analytical column with photodiode-array detector (PDA), which allowed the elution of the analytes within 12 min. The assay was linear over a concentration range of 0,4 to 6 μg mL^{-1} for 4-MBC, 1,2 to 18 μg mL^{-1} for OMC and 0,32 to 4,8 for AVO and it was applied to the analysis of the UV filters in the suncare products.

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Determination of 19 psychoactive substances in biological samples by ultra-high-performance liquid chromatography–tandem mass spectrometry

S. Karampela[1, 2] J. Smith[2], I. Panderi [1,*]

[1]Laboratory of Pharmaceutical Analysis, Division of Pharmaceutical Chemistry, Faculty of Pharmacy, National and Kapodistrian University of Athens, Panepistimiopolis, Zografou, 157 71 Athens, Greece.
[2] Division of Forensic Science, Department of Safety and Homeland Security, State of Delaware, USA.

E-mail: irenepanderi@gmail.com

Over the past decade, the trend towards substance abuse has changed, and many new psychoactive substances have emerged as "legal high" alternatives to traditional illicit drugs. This phenomenon was accompanied by increased incidences of people in intensive care units of hospitals after substance use with unknown pharmacological and/or toxicological action. The acute and chronic effects of new psychoactive sub-stances (NPSs) are not always known, and safety data on their toxicity are usually not available. By the end of 2018, the European Monitoring Centre for Drugs and Drug Addiction (EMCDDA) recorded more than 730 NPSs, 55 of which were detected for the first time on the drug market in Europe [1]. The forensic analysis of phenylethylamine derivatives is important because they can be involved in drug-related deaths. Fatal intoxication due to the recreational use of 3,4-methylenedioxymethamphetamine (MDMA) and 3,4-methylenedioxy-N-ethylamphetamine (MDEA) has been reported in Italy in 1996. MDMA overdosage can cause hyperthermia, cardiac arrhythmias, and renal failure, leading to death. More recently, 77 deaths where MDMA was detected in the body were reviewed and of these cases, 59 deaths had MDMA present in whole blood.

An ever-increasing need exists within the forensic laboratories to develop analytical processes for the qualitative and quantitative determination of a broad spectrum of new psychoactive substances [2]. Phenylethylamine derivatives are among the major classes of psychoactive substances available on the global market and include both amphetamine analogues and synthetic cathinones. In this work, an ultra-high-performance liquid chromatography-positive ion electrospray ionization tandem mass spectrometric method (UHPLC-ESI-MS/MS) has been developed and fully validated for the determination of 19 psychoactive substances, including nine amphetamine-type stimulants and 10 synthetic cathinone derivatives, in biological samples. The separation was achieved on a Poroshell 120 EC-C18 analytical column with a gradient mobile phase of 0.1% formic acid in acetonitrile and 0.1% formic acid in water in 9 min. The dynamic multiple reaction monitoring used in this work allowed for limit of detection (LOD) and lower limit of quantitation (LOQ) values of 0.5 and 2 ng mL^{-1}, respectively, for all analytes both in premortem and postmortem whole blood samples. A quadratic calibration model was used for the 12 quantitative analytes over the concentration range of 20–2000 ng mL^{-1}, and the method was shown to be precise and accurate. The method was applied to the analysis of real cases and proved to be a valuable tool in forensic and clinical toxi-cology.

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Label-free chemiluminescence detection of cysteine in biological fluids based on its catalytic effect on the reduction of gold ions to gold nanoparticles by luminol

T. Choleva[1], E.A. Akrivi[2,3], N. Kourkoumelis[2], D.L. Giokas[1] A.G. Vlessidis[1], G.Z. Tsogas[4]

[1] Laboratory of Analytical Chemistry, Department of Chemistry, University of Ioannina, 45110, Ioannina, Greece
[2] Department of Medical Physics, School of Health Sciences, University of Ioannina, Greece
[3] Neurology Clinic, University Hospital of Ioannina, Greece
[4] Laboratory of Analytical Chemistry, School of Chemistry, Faculty of Sciences, Aristotle University of Thessaloniki, GR-54124, Thessaloniki, Greece

E-mail: tacholeva@gmail.com

Biothiols are naturally occurring biomolecules that play a significant role in a variety of biological procedures [1]. In the human organism, cysteine (Cys), homocysteine (Hcy) and glutathione (GSH) are the most abundant biothiols and their concentrations are related to several clinical disorders and diseases [2]. The determination of biothiols is customarily based on the reactivity of their sulfhydryl group which enables the determination of the total concentration of biothiols but not their discrimination. In this work we demonstrate a selective assay for the determination of L-cysteine, based on the enhancement of the chemiluminescence signal generated when luminol is oxidized by gold ions towards the formation of gold nanoparticles. By appropriately regulating the experimental conditions and the sequence of reactions it is possible to discriminate Cys from GSH and Hcy, enabling its selective determination in protein-free blood plasma samples. The analytical features of the method in terms of sensitivity, linear range, recoveries, and reproducibility were satisfactory suggesting that the method holds promise to the speciation of biothiols in biological fluids.

Keywords: Chemiluminescence, gold nanoparticles, biothiols

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Voltammetric Determination of Rivaroxaban

M. Kartashov[1,2], J. Barek[1,2], E.I. Korotkova[2]

[1] Charles University, Faculty of Science, Department of Analytical Chemistry, UNESCO Laboratory of Environmental Electrochemistry, Albertov 6, CZ 128 43 Prague 2, Czech Republic
[2] National Research Tomsk Polytechnic University, Tomsk, Russia

E-mail: maks99661@gmail.com

Rivaroxaban is an oral anticoagulant from the group of direct factor Xa inhibitors. It is used to treat deep vein thrombosis and pulmonary embolism, as well as to prevent thrombi formation during atrial fibrillation and after hip or knee surgery [1]. Rivaroxaban could be considered for thromboprophylaxis in high VTE risk COVID-19 outpatients [2].
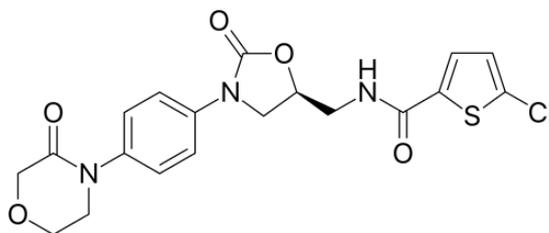
Figure 1. Chemical structure of rivaroxaban

At the moment, there is only one voltammetric technique intended for the determination of rivaroxaban [3], which shows the need to develop new voltammetric techniques and to study the electrochemical behavior of rivaroxaban in more details. The aim of this work is to study the voltammetric characteristic of the electrochemical redox reaction rivaroxaban and to develop a new "green" voltammetric method for the determination of rivaroxaban in dosage forms using various types of carbon electrodes to minimize the processes of their passivation. The influence of pH on cathodic voltammetric signal of rivaroxaban was investigated using graphite working electrode. Ag|AgCl was used both as an auxiliary and a reference electrode. LSV technique was used at the scan rate 100 mV/s. The maximum cathodic peak was observed in an alkaline medium (pH 10) at a potential of -1.68 V, in an acidic medium (pH 4), a maximum anodic peak was found at a potential of 1.38 V. Calibration curve was constructed under optimal conditions.

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Adsorptive stripping voltammetry as a convenient tool for the determination of Ti(IV) in plant samples on the example of nettle and horsetail

M. Adamczyk, M. Grabarczyk

[1]Department of Analytical Chemistry, Institute of Chemical Sciences, Faculty of Chemistry, Maria Curie-Sklodowska University, 20-031 Lublin, Poland

E-mail: marzena.adamczyk@poczta.umcs.lublin.pl

Both nettle and horsetail are very popular weeds found almost all over the globe. Among the wide range of applications, the most frequently mentioned is their use in cosmetics, improving the health condition of hair and nails. In addition to various biologically active ingredients with pro-health properties, these weeds are also characterized by a high content of titanium, which plays a key role in plant cultivation.

This work describes the determination of titanium content in the dried leaves of nettle as well as horsetail commercially available by three different producers (A, B, C) using the adsorptive stripping voltammetry method. In the voltammetric procedure, mercury film silver based electrode (Hg(Ag)FE) and chloranilic acid were used as a working electrode and a complexing agent, accordingly. The voltammetric analysis was preceded by the preparation of extracts from all tested plant materials by high pressure microwave-assisted mineralization. The standard adsorptive voltammetric measurements were run in a solution containing 0.1 mol L^{-1} acetate buffer pH $= 4.2$ and 1×10^{-3} mol L^{-1} chloranilic acid. The measurements were performed by differential pulse adsorptive stripping voltammetry from deaerated solutions. The accumulation of the complexes Ti(IV)-chloranilic acid onto the Hg(Ag)FE electrode were conducted at the potential of -0.3 V for 30 s while stirring the solution. At the end of accumulation time, the stirrer was switched off, and, after the equilibration time of 5 s, a differential pulse voltammogram was recorded, while the potential was scanned from -0.4 V to -0.9 V. The scan rate and pulse height were 20 mV s^{-1} and -50 mV, respectively. The titanium peak appeared at -0.6 V [1]. The developed procedure turned out to be a simple and fast tool for the determination of titanium in the prepared extracts. The sample matrix did not interfere with the determination of titanium. All determinations were carried out by the standard addition method.

The obtained results confirmed that both nettle and horsetail are rich in titanium. In the preparations tested in this study, made from commercially available dry materials of stinging nettle leaves, the titanium content was 56.0 ± 2.2 (C), 65.0 ± 3.5 (B) and 69.0 ± 3.2 μg/g (A) in dry matter of nettle [1]. However, the titanium content in the tested horsetail is slightly lower than in nettle and amounts to: $37,3 \pm 2,5$ (A), $41,8 \pm 2,2$ (B) and 56.3 ± 3.0 μg/g (C). It was proved that this simple and time-saving voltammetric procedure is suitable for high accuracy analysis of plant extracts.

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Selenium biotransformation by lactic acid bacteria, bifidobacteria and yeasts

V. Kantorová[1], A. Kaňa[1], G. Krausová[2], I. Hyršlová[2], O. Mestek[1]

[1]University of Chemistry and Technology Prague, Czech Republic
[2]Dairy research institute, Ltd., Prague, Czech Republic

E-mail: kantorov@vscht.cz

Selenium is an important trace element for human health. In its organic forms, such as selenoamino acids occurs as a component of selenoproteins and selenoenzymes, e.g. glutathione peroxidase, which is known for its antioxidative properties. The daily intake of selenium lower than 0.1 mg per kg of human body weight leads to selenium deficiency. This may cause several disorders of cardiac muscle. On the other hand, the intake over 1 mg per kg of body weight is considered toxic. [1]
Selenium is consumed from the food, and its natural sources are liver, sea food, and brazil nuts. [2] It´s clear that those sources are not the usual example of daily meals of an average Czech person. The lack of selenium then might be compensated by food supplements, especially selenium enriched yeasts, which could accumulate up to 3 g·kg^{-1} of selenium. [3] However, the daily consumption of tablets is not the best "user friendly" way. Functional foods like dairy products, using selenium enriched lactic acid bacteria (LAB) and yeasts, is a promising approach. In our study, we´ve obtain an overview, how LAB and yeasts (15 strains, 8 genuses) incorporate and transform selenite after being exposed to selenium enriched growth media. The total content of selenium was determined by using inductively coupled plasma mass spectrometry (ICP-MS) after microwave assisted digestion of the samples. The highest content of selenium, 45 mg·g^{-1} (d.w.) was found in *Streptococcus thermophilus*, but most of the samples has average content of Se lower than 1 mg·g^{-1} (d.w.). Speciation analysis was performed by coupling ion pair reversed phase liquid chromatography with ICP-MS, after extraction with proteinase XXIII. Major species (abundance up to 60%) detected were selenocystine, selenomethionine and one unknown species, while being major in *Bifidobacterium* strains. Other detected species as methylselenocysteine, selenite and selenate were minor, with content in units of percent. Most of the samples also showed the ability of producing selenium nanoparticles (SeNP). The transmission electron microscopy (TEM) provided an overview about the sizes and position of SeNP. It was shown, that most of the strains produce Se NPs extracellularly, two strains can produce both intra- and extracellularly, and three strains do not produce any particles at all. Se NPs average diameter vary from 20 to 350 nm. The total content of selenium in nano form was determined using single-particle ICP-MS. This multiple-step analysis allowed us to determine the mass balance of various selenium forms in enriched LAB and yeasts.

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Chromium fractionation in solid samples: the tannery waste scenario

E. Kokkinos, A. Zouboulis

Laboratory of Chemical & Environmental Technology, Department of Chemistry, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece

E-mail: evgenios@chem.auth.gr

Aiming in a sustainable future, numerous studies were published in recent years, regarding the further valorization of different wastes in the general framework of circular economy. A relevant work revealed the chromium and energy recovery from waste (sludge) produced during the treatment of tannery wastewaters [1]. In order to achieve higher chromium extraction efficiency, this waste was thermally treated. As a result, the initial insoluble trivalent chromium was oxidized to its hexavalent form, presenting much higher solubility. In the present study, the fractionation of chromium's different "species"/compounds in the aforementioned samples, i.e. the initial waste, the corresponding ash after energy recovery (400° C) and the thermally treated ash after chromium recovery (700° C), are investigated.

The chromium bound in the samples was determined by the application of sequential extractions. The different conditions/extractants are shown in Table 1, where the corresponding chromium fraction was obtained by using a different solvent. The liquid to solid ratio was equal to 45 and, if required, the step was repeated until zero chromium concentration in the produced leachate was observed.

According to the results, Cr oxides/hydroxides are the dominant "species" in the initial waste, mainly attributing to the presence of Cr(III). On the other hand, the thermally treated waste presents significantly lower distribution, regarding Cr(III) percentage, because its oxidation towards Cr(VI) is favored; therefore, the water soluble fraction, which is attributed to Cr(VI), increased proportionally.

Table 1. Chromium fractionation in the tannery waste and in the corresponding ashes [2].

Step	Fraction	Extractant	Cr (%)		
			Waste	**400 ºC**	**700 ºC**
1.	Water soluble	$0.015M\ KH_2PO_4$	0	48.6	90.4
2.	Exchangeable	$1M\ NH_4Cl$	0.1	0	0
3.	Organic bound	$0.1M\ Na_4P_2O_7$	4.8	1.2	1.2
4.	Carbonate bound	$1M\ NaCH_3COO$, pH 5	0.1	0	0
5.	Oxides and hydroxides	$0.04M\ NH_2OH.HCl$ in 25% v/v acetic acid, pH 2	95	50.1	8.3
6.	Chromium sulfide	$1M\ HNO_3$	0	0.1	0.1
7.	Residual	$HNO_3 + HClO_4 + HF$	0	0	0

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Chemical investigation of insect attractants and repellents using portable membrane inlet mass spectrometry.

S. Giannoukos,[1] A. Michaelakis[2], A. Pappa[1]

[1] National Technical University of Athens, School of Chemical Engineering, 157 73, Athens, Greece
[2] Department of Entomology and Agricultural Zoology, Benaki Phytopathological Institute, 8, S. Delta str., 14561 Kifissia, Athens – Greece

E-mail: stamatisgiannoukos@gmail.com

This work focuses on the development, testing and characterization of an olfactometer for the chemical investigation of volatile organic compounds (VOCs) with insect attractant and repellent activity. The olfactometer has a Y-shape design and comprises of four chambers: a) for hosting the under-investigation insect population, b) for hosting the chemical analytes of interest, c) a control chamber and d) a decision chamber. A portable membrane inlet (MI) triple-filter quadrupole mass spectrometer (QMS) was combined with the developed olfactometer and employed for the first time to detect and monitor continually, both qualitatively and quantitatively, VOCs with mosquito repellent activity. On-line monitoring of such compounds in field conditions (i.e. out of the laboratory) is of great importance and allows the chemical comprehension of the olfactory behaviour of insects with potential negative impact on public health. Portable membrane inlet mass spectrometry is a powerful, simple analytical technique ideal for *in-situ* applications that offers high sensitivity (low limits of detection), selectivity, fast (within seconds) and accurate analysis, with no sample preparation requirements. Representative compounds tested include: (-)-a-pinene, γ-terpinene, (+/-)-linalool, (-)-b-pinene, p-cymene, R-(+)-limonene, b-myrcene, (-)-linalool, butyl hexanoate, ethyl butyrate, o-xylene. Gas phase experiments were performed at concentration levels from low ppb to low ppm using two techniques for gaseous standards generation: a) static dilution bottles and b) a built-in-house vapour generator. Results obtained showed very good linearity within the examined concentration range, low ppb limits of detection and fast response times (within sec).

Distinguishing 1- and 3-methylhistidine isomers based on combined chromatographic and ion abundance characteristics

I. Sampsonidis[1], M. Marinaki[2], A. Pesiridou[2], H. Gkika[3], G. Theodoridis[2], N. Siachos[4], G. Arsenos[4], S. Kalogiannis[1]

[1]Department of Nutritional Sciences and Dietetics, International Hellenic University, Sindos Campus, Thessaloniki, GR57400, Greece

[2]School of Chemistry, Faculty of Sciences, Aristotle University of Thessaloniki, University Campus, Thessaloniki, GR54124, Greece

[3]School of Medicine, Aristotle University of Thessaloniki, University Campus, Thessaloniki, GR54124, Greece

[4]Faculty of Veterinary Medicine, Aristotle University of Thessaloniki, University Campus, Thessaloniki, GR54124, Greece

E-mail: isampson@nutr.teithe.gr, kalogian@ihu.gr

Since first isolated from human urine hydrolysates, 3-methylhistidine (3-MH) has been employed as a muscle protein turnover marker in a wide range of human and animal studies and is widely accepted as a marker for determining periods of increased muscle catabolism. Like all endogenous compounds, quantitative determination of 3-MH requires arduous method development, especially since its isomer, 1-methylhistidine (1-MH), which is a product of anserine breakdown, is widely present in most biological matrices of interest [1].

Traditionally, the individual determination of 1- and 3-MH requires either the use of ion chromatography or the utilisation of a derivatisation technique in combination with high performance liquid chromatography, to avoid co-elutions and determine the two isomers separately, and with gas chromatography, to enhance their volatility and facilitate the analysis. This is mainly due to the fact that the two compounds present almost identical retention behaviour in reverse phase chromatography and poor gas chromatographic behaviour as non-derivatised entities [2,3,4].

Here, we examine the possibility of facilitating unique spectral and ion abundance differences during collision induced dissociation between the two isomers along with different retention behaviour in certain hydrophilic interaction liquid chromatography conditions in order to develop a liquid chromatography coupled to tandem mass spectrometry method for the simultaneous separation and analysis of the two isomers without the need for derivatisation. Such a method will enable the analysis of large numbers of samples, that are usually associated with muscle catabolism studies, in an efficient and economic friendly manner.

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Combination of library-free and library-based metabolomics protocols formthe characterization of anti-cancer drug Carfilzomib administrated to mice

I. Barla[1], P. Efentakis[2], S. Lamprou[2], P-E. Nikolaou[2], E. Kastritis[3], MA. Dimopoulos[3], E. Terpos[3], I. Andreadou [2], N. Thomaidis [1], E. Gikas[1]

[1]National and Kapodistrian University of Athens, Department of Chemistry, Athens, Greece
[2]National and Kapodistrian University of Athens, Department of Pharmacy, Athens, Greece
[3]National and Kapodistrian University of Athens, Department of Medicine, Athens, Greece

*E-mail: vgikas@chem.uoa.gr

Introduction: Metabolomic profiling studies are powerful weapons for the comprehension of biological systems. Metabolites include all compounds of low-molecular-weight and have pivotal role, as metabolome is affected by every external or internal perturbation. Due to this sensitive response, the investigation of metabolites is frequently employed to provide "clues" in several health issues eg the definition of pathophysiological processes and the discovery of disease biomarkers including those of drug efficacy and drug toxicity.

Purpose: The main purpose of this study was to develop a metabolomics protocol for the characterization of renal homeostasis of mice treated with Carfilzomib (Cfz), an antineoplastic agent indicated for multiple myeloma. Despite Cfz's curative action, this agent is also associated with cardiotoxicity in humans, causing hypertension as the most common cardiovascular side effect. Considering that renal function plays a decisive role in blood pressure regulation, we sought to investigate the renal contribution in Cfz-induced hypertension. In order to research this observation, an experiment was performed to resemble the clinical findings in vivo : Twelve male mice were divided into two groups: i) Control Group: Treated with 0.9% Normal Saline ii) Cfz-Group: Treated with 8mg/kg of Carfilzomib for seven alternate days. The metabolomic study was applied on plasma, urine, and kidney samples obtained from the experimental groups at the end of the treatment.

Experiment: Three experimental treatment protocols, according to each sample type, were developed. The general steps of sample treatment involved protein precipitation and metabolites pre-concentration. An additional homogenization step was applied in the case of the kidneys. Data were acquired on a *Bruker Maxis Impact* QTOF/MS coupled to a *Thermo Dionex Ultimate 3000* LC. HILIC and Reversed Phased chromatography were employed respectively for small chain polar and non-polar metabolites. Additionally, both positive and negative ionization modes were implemented to collect as much information as possible.

Data treatment: Target screening based on an in-house metabolites data base and non-target screening for detection of unknown metabolites were performed. Referring to non-target analysis, several software tools were combined for data transformation, peak picking, signal correction, univariate and multivariate statistical analysis, and annotation.

Results: Target screening revealed differences between treated and control samples. Especially the kidney tissues analysis revealed increase in urea cycle metabolites (L-Alanine, L-Glutamine, glutamate, aspartate) and taurine content. Non-target multivariate analysis was employed for samples' classification providing several characteristic features, additional to those of target screening. Those features were annotated to specific biomarkers using online libraries.

Targeted metabolomics analysis for quantification of organic acids in urine by GC-MS/MS

T. Mouskeftara, C. Virgiliou, G. Theodoridis, H. Gika

[1]Department of Medicine, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[2]Biomic AUTh, Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center B1.4, 10th km Thessaloniki-Thermi Rd, P.O. Box 8318, GR 57001 Thessaloniki, Greece
[3]Department of Chemistry, Aristotle University of Thessaloniki, 54124, Greece

E-mail: gkikae@auth.gr

Organic acids (OAs) consist of mono-, di- and tricarboxylic acids or phenolic acids normally found as intermediates in a variety of intracellular metabolic pathways. The present study was to develop and validate a sensitive, accurate and effective method for the quantification of 48 OAs in human urine in the context of metabolomics applications in clinical or nutritional studies. Organic acids to be included in the study were selected based on their role as metabolic markers for Krebs cycle abnormalities, neuro-transmittion, nutritional deficiencies, antioxidant deficiencies, yeast and clostridia overgrowth, fatty acid metabolism. The determination of OAs was performed after simple removal of macromolecules and dual derivatization (methoxamine, MSTFA 1% TMCS), followed by GC-MS/MS (EVOQ 456 Bruker) analysis. To select the most appropriate sample preparation protocol for all 48 analytes, different solvents that have been applied in the literature for the extraction of low molecular weight metabolites or organic acids were considered: extraction was tested with MTBE, ethyl acetate, after sample acidification, whereas a more general protocol was applied by the addition of methanol. Thereafter different derivatization protocols with derivatization reagents as methoxamine, MSTFA, BSTFA were applied. The method was extensively validated, including the study of stability, and ensuring data accuracy, according to current guidelines for bioanalytical methods and applied to the analysis of human urine samples. To the best of our knowledge, no other fully validated method has been reported to cover such a wide range of OAs in urine using small volumes of urine and GC-EI-MS/MS technique.

An Alternative Technique to Calculate Mineral Carbon and Total Organic Carbon

K. Selekou[1], E. Chamilaki [2], V. Koulos [3], N. Pasadakis [1, 2]

[1] Institute of Petroleum Research, Foundation for Research and Technology (FORTH/IPR)
[2] School of Mineral Resources Engineering Department, Technical University of Crete
[3] BD Inventions P.C, Thessaloniki

E-mail: kselekou@ipr.forth.gr, ehamilak@mred.tuc.gr , vasilios.koulos@bd-inventions.com, pasadaki@ipr.forth.gr

A new promising technique, using FOGL Digital Soil Calcimeter (FOGL), by BD Inventions P.C, is employed to determine elemental carbon. Samples are treated with hydrochloric acid (HCl) to produce carbon dioxide that is measured and automatically converted into the elemental carbon percentage contained in the inorganic matter. Elemental Analysis (CHNS) is a well-established technique used to determine total elemental carbon. However, CHNS requires samples first to undergo acid-treatment, a difficult and time-consuming process, in order for the carbonates to be removed, and then the total organic carbon (TOC) is determined.

In this work, it is proposed that FOGL and CHNS (excluding the acid treatment) are combined, to determine organic carbon content. Both techniques are applied on two sets of samples, the first being mixtures of $MgCO_3$ and $CaCO_3$ in known concentrations, and the second sediments containing carbonate salts, from different regions of Greece. The inorganic carbon is obtained and compared using FOGL, Rock-Eval, and CHNS analysis in both sample sets, and the techniques are evaluated. It is demonstrated that FOGL provides consistent analytical results with the other two techniques. Furthermore, the TOC of the sediment samples was determined with Rock-Eval analysis and then compared to the calculated one; subtraction of the FOGL inorganic carbon from the CHNS total carbon. Results are comparable and satisfactory. To this end, we propose that CHNS combined with FOGL analysis, can replace the difficult and time consuming pretreatment step needed in the CHNS TOC method, and the Rock-Eval analysis when not accessible.

Passive sampling of pharmaceuticals and pesticides as a tool to assess their occurrence in marine aquacultures

E. Makou, V.I. Boti, I. K. Konstantinou, T.A. Albanis, D.G. Hela

Laboratory of Industrial Chemistry, Department of Chemistry, University of Ioannina, 45110 Ioannina, Greece

E-mail: dchela@uoi.gr

Increasingly widespread reports suggest the urgent demand of monitoring emerging contaminants, including pharmaceuticals and pesticides, in the marine environment and specifically in aquaculture waters. Passive sampling has been shown as a promising alternative to conventional water sampling for the monitoring of pharmaceuticals and pesticides[1]. The use of polar organic chemical integrative sampling (POCIS) passive samplers as a semi-quantitative and qualitative tool for screening of the selected target compounds (7 antibiotics, 13 pesticides and 25 pharmaceuticals /psychiatrics) in seawater has been studied. Data on the presence of these analytes are presented, that were collected twice a year using passive sampling techniques (with POCIS in pharmaceutical and pesticide configuration) in a fish farm placed on the Mediterranean Sea. Detection and quantification of the selected target compounds was carried out with liquid chromatography- high- resolution linear ion trap/ Orbitrap mass spectrometry (LC-HR-LTQ/Orbitrap-MS). Full scan in positive ionization (PI) mode was acquired, for identification and quantification purposes, at mass resolving power of 60,000FWHM, over a mass range of 120–500 Da. For confirmatory purposes, a data-dependent acquisition (full MS/dd-MS2) based on Collision Induced Dissociation (CID), was performed. Data-dependent MS2 analysis resolution was set at 15,000 FWHM and the ions were isolated in the LTQ ion trap and fragmented by using normalized collision energy (NCE) at 35%. The mass tolerance window was set to 5 ppm. During a two-season monitoring study, among the selected compounds, two antibiotics (trimethoprim in both seasons-April and September, sulfadiazine in September) and one antifouling compound (irgarol 1051 in september) were detected at the low ppt levels indicating the need for an organised monitoring alarm system of the marine aquaculture surrounding environment. Trimethoprim is a diaminopyrimidine antimicrobial agent and in veterinary medicine it is commonly used in combination with a sulphonamide (such as sulfadiazine) in a concentration ratio of 1:5 [2]. On the other hand, Irgarol-1051 is booster biocide that has been used to prevent biofouling on submerged surfaces such as boats, navigational buoys, underwater equipment and ships in marine environment. All the detected compounds were indentified on the basis of their retention time, formation of the protonated molecular ion $[M + H]^+$ and their main fragment ion (trimethoprim 291.1452/230.1162, sulfadiazine 251.0579/156.0114 and irgarol 254.1434/198.0811). Despite the compounds found, the detected levels do not pose any serious threat to the marine organisms.

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Investigation of drug-protein interactions using immobilized protein chromatography and its potential to model aquatic toxicity

C. Stergiopoulos[1], P. Danias[1], E. Notari[1], L.A. Tsakanika[1], A. Tsantili- Kakoulidou[2], F. Tsopelas[1]

[1] Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering, National Technical University of Athens, Iroon Polytechniou 9, 15780 Athens.
[2] Laboratory of Pharmaceutical Chemistry, School of Pharmacy, University of Athens, Panepistimiopolis, Zografou, 157 71 Athens, Greece.

E-mail: ftsop@central.ntua.gr

The evaluation of ecotoxicity is a necessity according to the REACH regulation "no data no market" [1]. Experimental determination of median toxicity in aquatic media, i.e. lethal concentration (LC_{50}) or effective concentration (EC_{50}) and Bioconcentration Factor (BCF) is based on standardized procedures based on biological organisms [2, 3]. However, this classic approach is lengthy, expensive and undergo ethical limitations, while in some cases its efficiency is questioned. Therefore, new laboratory-based tools with medium to high throughput need to be developed. In this aspect, immobilized plasma protein chromatography, using Human Serum Albumin (HSA) and Alpha-1 acid glycoprotein (AGP) stationary phases have been successfully used in drug discovery and development to model biological processes in the human body, such as the percentage of plasma protein binding and volume of distribution of drugs or the penetration of drugs through blood-brain barrier [4]. As the role of drug- protein interactions to aquatic toxicity and bioconcentration may be significant especially in the case of ionic species [5], immobilized plasma protein chromatography can prove as a valuable tool for ecotoxicological purposes.

In the present work, the potential of Immobilized protein chromatography to model aquatic toxicity of structurally-diverse drugs was studied. For this purpose, retention factors on HSA and AGP stationary phases were measured and compared to n-octanol- water partitioning in an effort to unravel elution mechanism. Retention factors were further used to model BCF and LC_{50} values of a series of aquatic organisms, compiled from literature sources. Linear models were established, which can be improved upon introduction of additional physicochemical parameters. For reasons of comparison, corresponding models were derived by replacing HSA/AGP retention factors with octanol-water partition coefficients (logP).

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EXSA: The European X-ray Spectrometry Association

D. Eichert[1,2] on behalf of EXSA Executive Committee[2]

[1]ELETTRA – Sincrotrone Trieste, S.S. 14 Km 163.5 in Area Science Park, 34149 Basovizza, Trieste, Italy
[2]Konkoly-Thege M. út 29-33. 26.ép.fsz.11., H-1121 Budapest, Hungary

diane.eichert@elettra.eu, excom@exsa.hu

X-ray spectrometry field of research is composed by many X-ray based techniques, which allows to shed some light into the properties of matter, in a complementary way. From micro-analytical to spatially nano-resolved technique, and time-resolved to energy-resolved technique, a plethora of investigations can be undertaken for various types of materials, and in many fields of research. For instance, X-ray fluorescence techniques are used qualitatively as well as quantitatively for sensitive and simultaneous elemental analysis. XRS is especially well-fitted for contributing in addressing some global challenges such as in Energy, Nanoelectronics, Health, and Environment as well as for the preservation of Cultural Heritage.

EXSA is a non-profit organization founded in 2004 for promoting cooperation and scientific exchanges between X-ray spectroscopists and analysts within Europe. It brings together users of X-ray spectrometry in various fields of research as well as manufacturers of X-ray instrumentation and developers of X-ray methodologies. EXSA aims to stimulate interaction and communication between young and experienced scientists, between academia and industry, thus fostering scientific progress and innovation.

EXSA supports the participation of talented young scientists in scientific events and co-organizes summer schools as well as dedicated workshops on novel and emerging X-ray spectrometry topics. This year, EXSA has put together its 1st Virtual Conference on X-ray Spectrometry, with four hot topics: XRS in environmental applications, XRS for novel materials including batteries, nanostructures, etc.; Instrumental development for XRS and its applications, and Fundamental parameters (https://www.exsa.hu/conf2021/).

EXSA awards scientists for their exceptional contributions to the field of X-ray spectrometry. Two awards for young X-ray spectrometrists and one award for outstanding scientific achievements are bestowed usually during the EXRS conferences.

EXSA is supporting the Fundamental Parameters Initiative and its workshops (https://www.exsa.hu/fpi.php), aiming for better knowledge of FPs as an essential requirement for accurate quantitative X-ray Analysis.

EXSA gathers more than 125 members from academia and industry and is sponsored by more than 12 companies. Since its foundation, EXSA has supported more than 50 workshops and conferences, among them the IMA Conference 2019 and 2021.

For more information, please visit www.exsa.hu.

Wednesday 22 September 2021

POSTER SESSION 2

Food Analysis | Quality control | Environmental Analysis | Spectroscopy

Chair: D. Giokas, R. Tzimou-Tsitouridou

Application of Raman spectroscopy for the detection of olive oil adulteration with seed oils

G. Bagkakis, P.K. Revelou, M. Xagoraris, E. Xera, C. Pappas, P. Tarantilis[*]

Chemical Laboratories, Agricultural University of Athens, Athens, Greece

[*]E-mail: ptara@aua.gr

Olive oil is a product of high nutritional value and economic importance. Due to its growing popularity, it is frequently a target of fraudulent practices such as adulteration with low-cost seed oils (blends). Taking into consideration the Commission Implementing Regulation 29/2012, which allows the mixing of seed oils with olive oil and the presence of a graphic element of olive oil is allowed provided that there is at least 50% content, this issue becomes crucial. Therefore, a need arises for the development of reliable, rapid, and low-cost methods to detect olive oil adulteration. In the present study, two Greek olive oil cultivars (Koroneiki and Megaritiki) were mixed with two different seed oils (sunflower oil and soybean oil). The percentage of seed oil in olive oil was varied from 0 to 60% (v/v) in 10% increments. Then a methodology was developed using Raman spectroscopy and chemometrics to determine the percentage of sunflower oil and soybean oil, in olive oil blends. Raman spectra were acquired in triplicates, and processed with the OMNIC software (Version 9.0 Thermo Nicolet). TQ Analyst software version 9.0 (Thermo electron Corporation, Madison, WI) was used for chemometric analysis. The Partial Least Squares (PLS) method was implemented using the spectral region at 1242-1283 cm^{-1}, which corresponds to the =C–H bending (scissoring) vibration of the *cis* RHC=CHR group. The statistical accuracy of the PLS chemometric model was described by the correlation coefficient (r), along with the root-mean-square error of prediction (RMSEP) and the root-mean-square error of calibration (RMSEC). The optimum number of factors was determined based on the minimum value of root-mean-square error of cross validation (RMSECV) and the first principal component was proposed for both PLS models. Very satisfactory results were obtained for the blends of olive oil with sunflower oil (r = 0.9788, RMSEC = 3.91, RMSEP = 3.93, RMSECV = 5.77) and soyabean oil (r = 0.9425, RMSEC = 7.14, RMSEP = 6.47, RMSECV = 8.25). The percentage of seed oil in adulterated olive oil was successfully determined by applying the PLS method to the Raman spectra. Raman spectroscopy combined with chemometrics proved to be a powerful tool for the detection of olive oil adulteration significant for quality control applications.

High resolution mass spectrometry study of olive oil phenolic constituents in Megaritiki cultivar

E. Xera, P.K. Revelou, M. Xagoraris, C. Kanakis, C. Pappas, P. Tarantilis[*]

Chemical Laboratories, Agricultural University of Athens, Athens, Greece

[*]E-mail: ptara@aua.gr

Olive oil is derived from the fruits of the olive tree (*Olea europaea* L.) through physical or mechanical methods. The Mediterranean diet encompass the regular consumption of olive oil which contributes to a lower risk of developing cancer and coronary heart disease. The health benefits acquired from olive oil consumption can be mainly attributed to the presence of antioxidant phenolic constituents. Consequently, the study of the phenolic profile of olive oil is of great importance as it is associated with its nutritional and commercial value. Megaritiki cultivar (*Olea europaea* var. *argentata*) is cultivated mainly in Attica, Peloponnese, and Central Greece regions. Although this cultivar is one of the most commercial in Greece, research on individual phenolic compounds is still scarce. In this study, 20 constituents belonging to phenolic acids, phenolic alcohols, flavones, and secoiridoids, were studied by high resolution mass spectrometry in Megaritiki olive oil samples from Attica, Peloponnese, and Central Greece. The targeted analysis experiments were performed in negative electrospray ionization mode with an Agilent 6530 Quadrupole Time of Flight LC-MS system (QTOF LC-MS), coupled with an Agilent 1290 Infinity UHPLC system and an autosampler (Agilent Technologies, Santa Clara, USA). Nucleoshell Bluebird RP 18, particle size: 2.7 μm, length: 100 mm, ID: 4.6 mm was used as a column and the mobile phase consisted of a gradient (A: acetonitrile/0.1% acetic acid; B: H_2O/0.1% acetic acid). Phenolic acids were mainly identified in olive oil samples from Peloponnese and Central Greece, such as syringic acid, *p*-coumaric acid, caffeic acid, protocatechuic acid, *p*-hydroxybenzoic acid, cinnamic acid, and hydroxytyrosol acetate. In Megaritiki olive oil from Attica, the phenolic alcohol hydroxytyrosol, the flavones apigenin and luteolin, the secoiridoids oleacein, oleocanthal, and oleuropein aglycon, and various phenolic acids were identified. The identification was performed using commercially available standards and the publicly available databases ReSpect for phytochemicals and human metabolome database (HMDB). This study serves as a valuable contribution for a deeper investigation of the understudied Greek olive oil Megaritiki cultivar and will contribute to the enhancement of its commercial value.

Volatile compound characterization of olive oil from Koroneiki cultivar by SPME-GC/MS and differentiation according to geographical origin

P.K. Revelou, M. Xagoraris, E. Xera, C. Kanakis, C. Pappas, P. Tarantilis*

Chemical Laboratories, Agricultural University of Athens, Athens, Greece

*E-mail: ptara@aua.gr

Solid phase microextraction-gas chromatography-mass spectrometry (SPME-GC/MS) is a well-established technique for the study of volatile compounds of olive oil. The authentication of geographical origin of olive oil is crucial since it affects its commercial value. In this study 41 olive oil samples from three Greek regions (12 Heraklion, 10 Lakonia, 19 Messinia) were analyzed using a divinylbenzene/carboxen/polydimethylsiloxane (DVB/CAR/PDMS) fiber (Supelco, Bellefonte, PA, USA) and a Thermo GC-TRACE ultra, interfaced to a Thermo Mass Spectrometer DSQ II (Thermo Scientific Inc., Waltham, MA, USA). Statistical analysis was performed with XLSTAT ver. 2020.3.1.0 software (Addinsoft Deutschland, Andernach, Germany). Forty-seven volatile compounds were identified and semi-quantified using b-ionone as an internal standard. Subsequently, the olive oil samples were divided into three groups (Heraklion, Lakonia, Messinia) and the forward stepwise algorithm was applied for the identification of the most discriminating variables (volatiles) for the geographical differentiation of samples. Two chemometric models were developed for the classification of olive oil samples using the linear discriminant analysis (LDA) and quadratic discriminant analysis (QDA). The chemometric models were validated utilizing the cross-validation method.

Five volatiles, namely, tetradecane, 1-penten-3-ol, 2-hexenal, 1-pentanol and 2-hexanone were selected by the stepwise algorithm for the discrimination of olive oil samples. Using these volatiles as explanatory variables, LDA was performed where correct classification and cross validation were 90.2 % and 85.4 %, respectively. Applying the QDA algorithm 90.2 % of the samples were correctly classified while after the cross-validation method this percentage was lower, at 70.7 %. The above results indicate the high potential of the SPME-GC/MS technique in combination with the forward stepwise algorithm as well as the LDA and QDA algorithms for the differentiation of olive oil according to geographical origin.

The use of FTIR-ATR and SPME-GC-MS for botanical origin differentiation of unifloral honeydew and blend honeydew Greek common honey

M. Xagoraris[1], P.K. Revelou[1], F.P. Vardaka[2], E. Alissandrakis[2,3], P.A. Tarantilis[1], C.S. Pappas[1]

[1]Laboratory of Chemistry, Department of Food Science and Human Nutrition, Agricultural University of Athens, 75 Iera Odos, 11855 Athens, Greece
[2]Laboratory of Quality and Safety of Agricultural Products, Landscape and Environment, Department of Agriculture, Hellenic Mediterranean University, Stavromenos PC 71410, Heraklion, Crete, Greece
[3]Institute of Agri-Food and Life Sciences Agro-Health, Hellenic Mediterranean University Research Center, Stavromenos PC 71410, Heraklion, Crete, Greece

E-mail: ealiss@hmu.gr

In recent years, there has been an increasing demand for honeydew honey, with subsequent high export activity in global markets. Honeydew honey is crucial for Greek beekeeping, with production exceeding 70% of total annual production. The most popular honeydew honeys in Greece are pine and fir honey, making their authenticity of high interest to producers, consumers, and traders. In the present work, a total of 38 honey samples (14 fir, 12 pine, 12 honeydew blends) were provided directly from beekeepers. All honey samples were confirmed of their origin with physicochemical and pollen analyses. Using solid phase micro extraction (SPME), a fraction of volatile compounds was studied using gas chromatography coupled with mass spectrometry (GC-MS). Also, Fourier transform infrared (FTIR) spectra of honey samples were obtained using attenuated total reflection (ATR) spectroscopic technique. Botanical origin discrimination was achieved performing stepwise linear discriminant analysis (SW-LDA) and cross-validation method using SPSS v.25 software. Eight volatile compounds, octane; octanal; undecane; nonanal; nonanol; decanal; methyl nonanoate and nonanoic acid were selected from stepwise algorithm as the most significant for the differentiation. Also, the spectral regions 1390–945 and 847–803 cm^{-1} presented statistical significance and were selected with higher absolute coefficients for the discrimination, correspond mainly to the sugars (fructose and glucose). SPME-GC-MS indicate a classification rate of 86.8% while FTIR 92.1%. Moreover, cross-validation resulted to a 60.5% and 81.66% correct classification, respectively. The Wilks' λ values for each chemometric model were indicative of the power of the models ($p < 0.05$). SPME-GC-MS is a widely used method for honey authentication, yet in this case, spectroscopic data allowed better discrimination results with promising future potential uses. In addition, FTIR spectroscopy is a simple, rapid, economical and environmentally friendly (not solvents used) method for honey authentication.

GC-MS and LC-QTOF-HRMS for volatile and phenolic analysis of strawberry tree honey from Greece

M. Xagoraris[1], P.K. Revelou[1], E. Savvidaki[2], E. Lazarou[1], E. Alissandrakis[2,3], P.A. Tarantilis[1], C.S. Pappas[1]

[1]Laboratory of Chemistry, Department of Food Science and Human Nutrition, Agricultural University of Athens, 75 Iera Odos, 11855 Athens, Greece
[2]Laboratory of Quality and Safety of Agricultural Products, Landscape and Environment, Department of Agriculture, Hellenic Mediterranean University, Stavromenos PC 71410, Heraklion, Crete, Greece
[3]Institute of Agri-Food and Life Sciences Agro-Health, Hellenic Mediterranean University Research Center, Stavromenos PC 71410, Heraklion, Crete, Greece

E-mail: ealiss@hmu.gr

Strawberry tree (*Arbutus unedo* L.) honey is one typical Mediterranean honey with bitter taste unique aroma profile and biological activity. In Greece, strawberry tree honey usually is not harvested but left for the bees without much commercial interest. However, in recent years this type of honey presents a notable interest from a specific group of consumers due to its distinct and rare organoleptic characteristics. Also, a remarkable economic importance at local markets is reported, with 4 to 8 times higher prices than other honey types[1]. Despite its economic interest, research on the chemical properties of strawberry tree honey are scarce. In the present work the volatile and phenolic compounds of four samples were studied. Two volatile fractions were received, using isopropanol and hexane and analyzed by gas chromatography coupled with mass spectrometer (GC-MS). The phenolic compounds were extracted with acetonitrile and analyzed by liquid chromatography combined with time-of-flight high-resolution mass spectrometry (LC/Q-TOF/HRMS). The main volatile compounds, mainly norisoprenoids, which were identified in isopropanol fraction were 1,3,3-trimethyl-7-oxabicyclo[4.1.0]heptane-2,5-dione (40.84%), 5-(hydroxymethyl)furan-2-carbaldehyde (9.49%), 6-butyl-3-methoxycyclohex-2-en-1-one (3.12%), and 3,5-dihydroxy-6-methyl-2,3-dihydro-4H-pyran-4-one (1.84%). Respectively, the main volatiles of hexane extract were 3-hydroperoxyhexane (3.60%), 2-hydroperoxyhexane (3.52%), 3,4,5-trimethylphenol (2.56%), 3,5,5-trimethylcyclohex-2-en-1-one (2.32%), and (*E*)-3,5,5-trimethyl-4-(3-oxobut-1-en-1-yl)-cyclohex-2-en-1-one (2.17%). In addition, 30 phenolic compounds were identified with homogentisic acid, abscisic acid and unedone being the most characteristics. In conclusion, strawberry tree honey has a special chemical composition and organoleptic characteristics, while it presents biological activity. Greek legislation does not clearly define physicochemical and mellisopalynological characteristics for this honey variety, so further study is needed towards the definition of its monofloral characteristics.

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SPME/GC-MS study of volatile compounds from strawberry tree and autumn heather honeys

M. Xagoraris[1], E. Lazarou[1], P.K. Revelou[1], E. Alissandrakis[2,3], P.A. Tarantilis[1], C.S. Pappas[1]

[1]Laboratory of Chemistry, Department of Food Science and Human Nutrition, Agricultural University of Athens, 75 Iera Odos, 11855 Athens, Greece
[2]Laboratory of Quality and Safety of Agricultural Products, Landscape and Environment, Department of Agriculture, Hellenic Mediterranean University, Stavromenos PC 71410, Heraklion, Crete, Greece
[3]Institute of Agri-Food and Life Sciences Agro-Health, Hellenic Mediterranean University Research Center, Stavromenos PC 71410, Heraklion, Crete, Greece

E-mail: ealiss@hmu.gr

Strawberry tree (*Arbutus unedo* L.) and autumn heather (*Erica manipuliflora* Salisb.) are important beekeeping plants of Greece. Six monofloral honeys (four strawberry tree, two autumn heather) were analyzed by means of Solid Phase Micro-Extraction, followed by Gas Chromatography coupled to Mass Spectrometry (GC-MS). A DVB/CAR/PDMS fiber was employed, and benzophenone saw used as internal standard. The volatile compounds with higher concentration (μg/ g of honey expressed as benzophenone) from strawberry tree honey samples, were α-isophorone (2.50-8.12), 3,4,5-trimethyl-phenol (0.20-4.62), 2-hydroxy-isophorone (0.06-0.53), 4-oxoisophorone (0.38-0.46), and β-isophorone (0.02-0.43). Regarding heather honey samples, the most abundant compounds were 1-methoxy-4-propyl-benzene (1.22-1.40), p-anisaldehyde (0.97-1.28), p-anisic acid (0.35-0.58), 2-furaldehyde (0.52-0.57), and benzaldehyde (0.41-0.56). Norisoprenoids are potent floral markers for strawberry-tree honey; β-isophorone is found exclusively in the volatile fraction of this type of honey, while also α-isophorone, 4-oxoisophorone and 2-hydroxy-isophorone could be considered as additional marker compounds. The analysis of autumn heather honey revealed that phenolic compounds are the most abundant and p-anisaldehyde, 1-methoxy-4-propyl-benzene, and p-anisic acid could serve as potent marker compounds. In conclusion, marker compounds for the determination of the botanical origin for these honeys could be identified as several norisoprenoids and phenolic components were found exclusively or in higher concentrations compared to common Greek honey varieties.

Application of optical spectroscopic techniques and multivariate statistical analysis as a method of determining the percentage and type of adulteration of extra virgin olive oil

G. Stavrakakis[1,2], A. Philippidis[1], M. Velegrakis[1]

[1]Institute of Electronic Structure and Laser, Foundation for Research and Technology-Hellas (IESL-FORTH), 700 13 Heraklion, Crete, Greece
[2]Department of Materials Science and Technology, University of Crete, 700 13 Heraklion, Crete, Greece

E-mail: vele@iesl.forth.gr

The adulteration of extra virgin olive oil (EVOO) with low quality edible oils is a major problem in the industry leading to economic frauds as well as customers' health issues. A number of techniques have been used in the recent years for the characterization of extra virgin olive oil adulteration [1,2]. In the current work, the spectroscopic techniques of absorption and fluorescence in combination with statistical analysis have been exploited to classify and quantify the adulteration of EVOO with different seed oils and pomace oil. The specific spectroscopic methods are rapid and with low cost. The results of the present work indicate that both techniques are suitable for olive oil quality control, with absorption at visible region (400-800 nm) being superior for the specific number and types of oil samples. Specifically, visible absorption spectroscopy presents adulteration detection limits below 10%, a percentage within the range of commercial interest. The model accuracy of classification for the mixtures of adulterated oils is above 90%. We present a facile method for obtaining simultaneously the amount of adulteration and the type of the adulterant in EVOO.
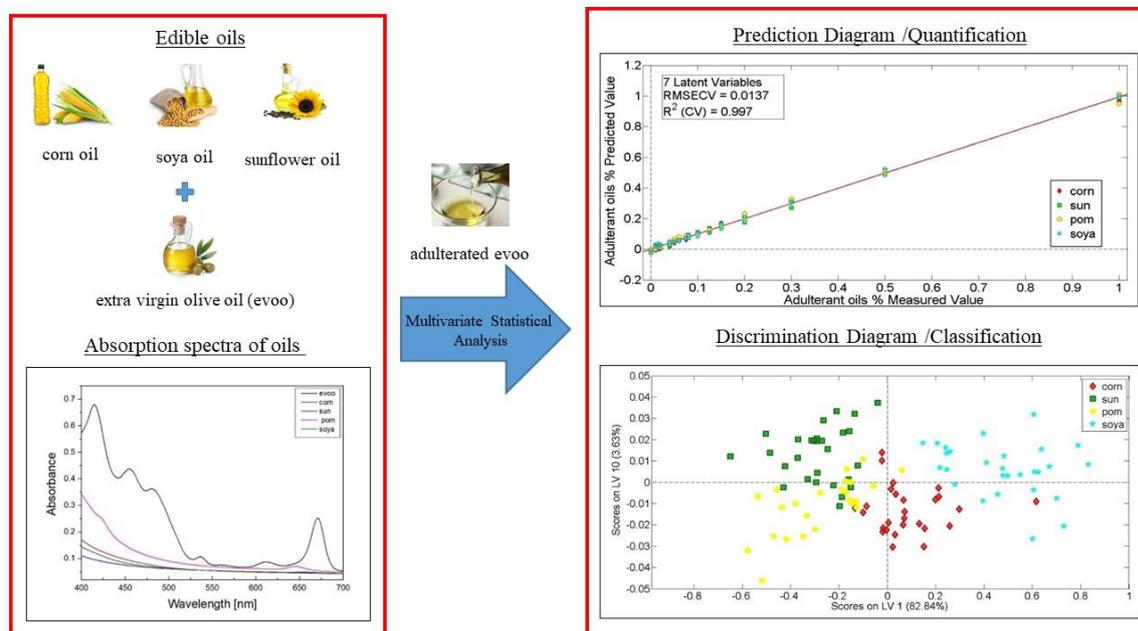
Figure 1. Absorption spectra of different types of oils and the exported multivariate statistical analysis diagrams

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Optimized methods for the extraction of C-phycocyanin and β-carotene from *Arthrospira* spp. (Spirulina) and their application to microalgae isolated from Greece

A. Paraskevopoulou[1], T. Kaloudis[1], S. Gkelis[2], U. Lortou[2], M. Panou[2], A. Hiskia[1], T.M. Triantis[1*]

[1] Institute of Nanoscience and Nanotechnology, NCSR Demokritos, Athens, Greece
[2] Department of Botany, School of Biology, Aristotle University of Thessaloniki,
GR-541 24 Thessaloniki, Greece

* E-mail: t.triantis@inn.demokritos.gr

Cyanobacteria belonging to *Arthrospira* spp.*,* commercially known as *'Spirulina',* are gaining more and more interest, especially in the food supplements sector, due to their high content in proteins, nutrients, and compounds with beneficial properties. Among these chemicals, phycocyanin (C-PC) and β-carotene are two of the most popular natural pigments. They are commercially available in several forms and used in the food industry, mainly as color additives in various food products [1]. Pigment-rich extracts are described as antioxidant, anti-inflammatory, and anti-tumoral agents with great interest for applications in food and cosmetical industries due to their bioactive potential and their green product label [2,3]. The quantity and purity of these metabolites in the crude extracts can be affected by the employed extraction techniques and conditions, so there is a need for optimized methods that can bring value and market interest to the final products.

This work focused on the optimization of C-PC and β-carotene extraction techniques from Spirulina biomass and their application to lab-cultured microalgae strains isolated from diverse habitats of Greece. Selected Spirulina samples were further fractionated in order to evaluate the antioxidant activity of different fractions. Several techniques of cell disruption were tested for C-PC extraction (sonication, magnetic stirring, etc.) and optimized experimental parameters included an initial pre-treatment of the sample for 2 h, at 25 $^\circ$C, using a phosphate buffer (C=0.01 M) and a solvent to biomass ratio equal to 10:1 followed by a 5 min sonication. In case of β-carotene, optimum extraction conditions included intense stirring for 1.5 h using a mixture of acetone and ethanol. Both methods were applied under optimum conditions for C-PC and β-carotene extraction. The results showed that C-PC content ranged from 1.1 to 61.4 mg C-PC/g d.w., while C-PC purity ratio (A_{620}/A_{280}) ranged from 0.15 to 1.16. Although *Arthrospira* spp. have shown low C-PC content values (2.4 – 3.6 mg C-PC/g d.w.), their crude extracts exhibited elevated C-PC purity (0.71-0.85). β-Carotene content ranged from 0.1 to 180.7 μg/g d.w. with *Arthrospira* spp. being among the strains with the highest content (135.3 μg/g d.w.). Extraction and fractionation of selected Spirulina samples were performed using solvents with different polarities (water/methanol, methanol, chloroform, acetone/ethanol, ethyl acetate, and n-butanol) and the antioxidant activity of all fractions was evaluated using the DPPH and ABTS assays. Results revealed that methanolic extracts exhibited the highest antioxidant activity, followed by acetone/ethanol extracts. The present study clearly demonstrates that significant amounts of C-PC and β-carotene can be extracted under optimal conditions from local microalgae strain cultures, while a fractionation protocol applied to Spirulina biomass samples produced fractions enriched with metabolites that exhibit high antioxidant activity.

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Optical spectroscopy as an alternative tool for honey analysis

E. Boumpoulaki[1,2], E. Orfanakis[1,3], A. Zoumi[1], A. Philippidis[1], M. Velegrakis[1]

[1]Institute of Electronic Structure and Laser, Foundation for Research and Technology-Hellas (IESL-FORTH), Heraklion, Greece
[2]Department of Chemistry, University of Crete, Heraklion, Greece
[3]Department of Materials Science and Technology, University of Crete, Heraklion, Greece

E-mail: vele@iesl.forth.gr

The quality and the botanical origin of honey affects its commercial value and is an important issue for consumers. Therefore, it attracts both commercial and research interest. The aim of the present study is the application of optical spectroscopic techniques such as FT-IR, Raman, Absorption and Fluorescence spectroscopy coupled with machine learning-based modeling for honey analysis. These techniques offer a quick monitoring of the characteristic substances contained in honey. The findings of this work demonstrate that spectroscopic techniques provide a rapid, useful tool that requires no sample pre-treatment for the evaluation of honey according to its botanical origin and physicochemical parameters.

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The isotopic approach to the authenticity of traditional products from Cyprus and Aegean islands for strengthening their identity

E. Ioannou Papayianni, E. Tzioni, C. Savvidou, C. Louka, R. Kokkinofta*

State General Laboratory, P.O.Box 28648, Nicosia, Cyprus

E-mail: sglsnif@cytanet.com.cy

Local products are an integral part of an area's local heritage as they contribute to the adoption of a region's "identity", creating bonds between its products, their unique characteristics, the geoclimatic conditions existing in the region and the culture. Recently the consumers have diverted their interest towards local traditional products, which possess unique quality characteristcs. The increased demand of those products is an economic opportunity for areas that are able to diversify and properly promote their local traditional products.

In the framework of "AGROFOOD" project funded by the INTERREG V-A Greece- Cyprus 2014-2021, the specific characteristics of indigenous Cypriot products (wines, alcoholic beverages, juices, honey) were studied and compared to similar products from the Greek islands of the northern Aegean, in order to differentiate and certify their origin. The results were processed using multivariate chemometric techniques.

The combined information from isotopic ratios of $^{13}C/^{12}C$ and $^{18}O/^{16}O$ by IRMS and D/H by SNIF-NMR spectroscopy, has been observed to create a unique isotopic fingerprinting of the products studied. It is believed that the differentiation of local products is related to the unique geological and climatic conditions existing in the islands.
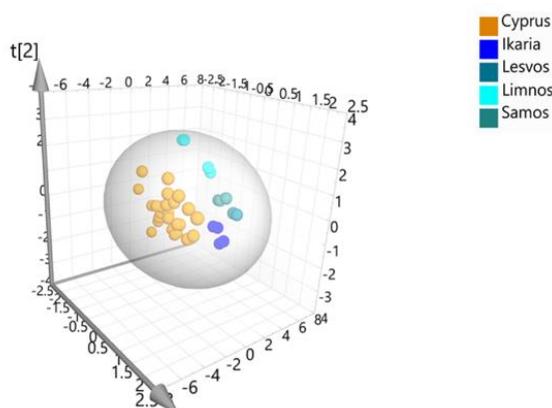
Figure 1: Scatter plot of wines from different origin, according to their stable isotopic fingerprint.

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Simultaneous speciation analysis of arsenic and selenium species in seafood and onion samples by HPLC-ICP-MS

K. Karaś[1], D. Jakkielska, A. Zioła-Frankowska[2], M. Frankowski[1]

[1]Department of Analytical and Environmental Chemistry, Faculty of Chemistry, Adam Mickiewicz University, Poznan, Poland
[2]Department of Analytical Chemistry, Faculty of Chemistry, Adam Mickiewicz University, Poznan, Poland

E-mail: katarzyna.karas@amu.edu.pl

Human health is one of the most important area discussed at the world's top congresses. The number of elements that have negative effect on human health is constantly growing and should be carefully monitoring. The toxicity of elements mainly depends on the form, in which these chemical species occur. This specific form of element may have different impact to living organism.

In order to assess potential toxicity of each element form, it is needed to use speciation analysis, which allow to separate the interesting components of the material and give the qualitative and quantitative data about the composition of tested mixture.

The main purpose of this work was to develop a new method for simultaneous speciation analysis of arsenic (As(III) - arsenite, As(V) - arsenate, DMA - dimethylarsinic acid, MMA - monomethylarsonic acid, As-B - arsenobetaine) and selenium (Se(IV) - selenite, Se(VI) - selenate, Se-methionine, Se-cystine) forms, which was applied to analysis of seafood and onion samples available on Polish market. Determination was made using hyphenated technique - high-performance liquid chromatography inductively coupled plasma mass spectrometry (HPLC-ICP-MS). This technique allowed to create conditions for the chromatographic separation and determination of target forms with high sensitivity and with impressive lower detection limit.

Taking into account results obtained from speciation analysis, the highest level of arsenic in seafood samples were found in red argentine shrimps and the lowest one in white shrimp. However, in the case of arsenic forms in the onion samples, no significant differences in the concentration levels between the types of onions were found. The highest concentration of selenium in seafood samples was determined in red shrimp, but the lowest one in argentine red shrimp. In the case of onion samples, the highest level of selenium was observed in green sweet onion sample, but the lowest one in green bio-onion. Obtained results was compared with results from ICP-MS analysis.

It should be underlined that the analysis was performed in one analytical cycle and the proposed method presented good selectivity, linearity and sensitivity for both arsenic and selenium species. It may be also good tool to provide necessary data of arsenic and selenium species in samples with similar matrix and nature.

Analysis of polyester cyclic oligomers of polyethylene terephthalate in pasta with modified QuEChERS clean-up and UHPLC-qTOF-MS

E.D. Tsochatzis [1], J. Alberto Lopes[2], H. Gika[3,4,5], G. Theodoridis[3,4,5]

[1]Department of Food Science, Aarhus University, Agro Food Park 48. 8200 Aarhus N. Denmark
[2] European Commission, Joint Research Centre, Geel, Belgium
[3]FoodOmicsGR Research Infrastructure, AUTh node, , GR 57001 Thessaloniki, Greece.
[4]Biomic AUTh, Center for Interdisciplinary Research and Innovation, GR 57001, Thessaloniki, Greece.
[5] Aristotle University of Thessaloniki, Thessaloniki, Greece.

E-mail : Emmanouil.Tsochatzis@food.au.dk

An Ultra High-Performance Liquid chromatography method quadruple time-of-flight mass spectrometry has been developed for the analysis of 11 cyclic polyesters oligomers, extracted with 1,1,1,3,3,3-hexafluoro-2-propanol (HFIP)/trifluoroethanol (TFE)/ethanol (2:2:4 v/v/v) following a modified QuEChERS clean-up with alumina/primary secondary amine, in pasta. Target analytes were polyethylene terephthalate (PET) 1st series cyclic dimer to heptamer, polybutylene terephthalate (PBT) dimer to pentamer and a polyurethane oligomer (Figure 1). Standard addition method was applied for the calibration, and the limits of quantification ranged from 3.2 to 17.2 ng g^{-1}. Recoveries ranged from 86.4 to 109.8%, RSDs were lower than 12% for all analytes, and matrix effect never exceeded ± 2.5%
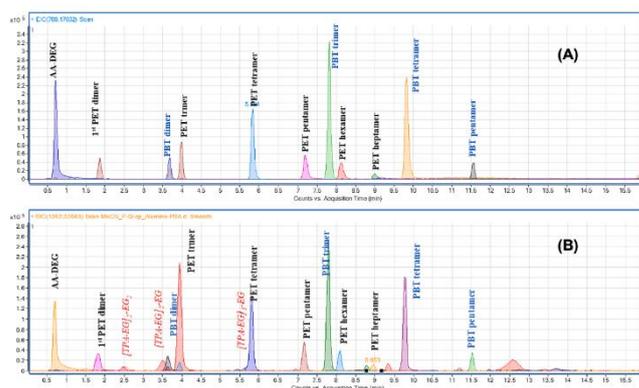
Figure 1. UHPLC-QTOF-MS extracted Ion Chromatogram of (A) analytical standards mixture at a level of 0.1 µg kg^{-1} 1 for each polyester cyclic oligomer; (B) spiked pasta sample (0.10 µg kg^{-1}).

The method was successfully applied to real commercial pasta samples, where the PET 1st series cyclic trimer was the most abundant oligomer, being found in all tested samples. The 1st series PET cyclic dimer and tetramer, as well as 1,4,7-trioxacyclotridecane-8,13-dione, were found in considerable amounts. Traces of the 2nd and 3rd series PET cyclic dimers were also found (Table 1).

Table 1. Analysis of target analytes in pasta samples.

Target analytes	(ng g^{-1})*			
	ML1 (Paper/ PET)	ML2 (Paper/ PET)	ML3 (Plastic incl. PET)	ML4 (Plastic incl. PET)
AA-DEG	10.2 ± 0.7	12.7 ± 1.2	14.2 ± 1.4	10.5 ± 0.7
[TPA-EG]$_2$	<LOD	<LOD	19.2 ± 1.6	17.4 ± 1.7
[TPA-BG]$_2$	<LOD	<LOD	<LOD	<LOD
[TPA-EG]$_3$	60.5 ± 4.4	45.2 ± 4.2	140.2 ± 12.9	164.1 ± 13.5
[TPA-EG]$_4$	16.2 ± 1.6	17.2 ± 0.9	22.3 ± 2.2	24.9 ± 2.5
[TPA-EG]$_5$	<LOD	<LOD	<LOD	<LOD
[TPA-BG]$_3$	<LOD	<LOD	<LOD	<LOD
[TPA-EG]$_6$	<LOD	<LOD	<LOD	<LOD
[TPA-EG]$_7$	<LOD	<LOD	<LOD	<LOD
[TPA-BG]$_4$	<LOD	<LOD	<LOD	<LOD
[TPA-BG]$_5$	<LOD	<LOD	<LOD	<LOD
Total content; sum (ng g^{-1})	70.7	75.1	195.9	216.9

*All samples were analysed in triplicates (n = 3).

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Wide scope screening of >900 pesticides in olive oil using LC-ESI/GC-APCI-QToF-MS

S.K. Drakopoulou[1], S.E. Kokolakis[1], M.E. Dasenaki[1,2], A.S. Kritikou[1], N.S. Thomaidis[1]

[1]National and Kapodistrian University of Athens, Department of Chemistry,
Laboratory of Analytical Chemistry, Panepistimiopolis Zografou, 15771 Athens, Greece
[2]National and Kapodistrian University of Athens, Department of Chemistry,
Laboratory of Food Chemistry, Panepistimiopolis Zografou, 15771 Athens, Greece

E-mail: ntho@chem.uoa.gr

Olive oil has been acknowledged as food of high nutritional value with several health benefits derived from its consumption. Being declared as an integral part of the Mediterranean diet, it is highly ranked in the global market, with its commercial price and availability varying according to olive oil production per year. To increase agricultural production and eliminate the risk of crop loss, producers often resort to pesticides' administration. However, the extensive use of pesticides poses serious risks to consumer health, as part of them could pass into crude olive oil at alarming concentrations, with reference to Maximum Residues Limits (MRL) (*Regulation (EC) No 396/2005*).

In this framework, an efficient QuEChERS-based method was developed, appropriate for the extraction of pesticides of different chemical properties. Taking full advantage of two different High Resolution Mass Spectrometry (HRMS) approaches, Liquid Chromatography and Gas Chromatography coupled to hybrid mass spectrometry analyzer of type quadrupole time of flight (LC/GC-QTOF-MS), detection of more than 900 pesticides was successfully achieved. Olive oil extracts were analysed using reversed phase Liquid Chromatography (RPLC) in positive and negative ionization mode with an electrospray source (ESI), while gas chromatography system was operated in positive ionization with a source of Atmospheric Pressure Chemical Ionization (APCI). The sample preparation methodology was common for both techniques, differentiated only on reconstitution step, which makes the method highly applicable in routine analysis. The method was fully validated according to the *European Commission Directive SANTE /11813/2017*. Thus, its performance characteristics were evaluated in terms of linearity, accuracy and precision, while the limits of detection and quantification of the method were determined and matrix effect was estimated. Finally, twenty olive oil samples from Lesvos Island were analysed with the proposed methodologies and were found to be in compliance with European legislation, detecting pesticides' residues below the established MRLs.

Holistic untargeted LC-HRMS-based metabolomics workflow for food authenticity studies: Exploiting both RPLC- & HILIC-HRMS platforms capability for citrus juices authenticity investigation

A.S. Kritikou[1], S.K. Drakopoulou[1], M.E. Dasenaki[1,2], E.Gikas[1], N.S. Thomaidis*,[1]

[1]National and Kapodistrian University of Athens, Department of Chemistry,
Laboratory of Analytical Chemistry, Panepistimiopolis Zografou, 15771 Athens, Greece
[2]National and Kapodistrian University of Athens, Department of Chemistry,
Laboratory of Food Chemistry, Panepistimiopolis Zografou, 15771 Athens, Greece

E-mail: ntho@chem.uoa.gr

Fruit juices authenticity is a challenging task, as the high cost of the fruit and the possibility of poor harvest, conflicting with high consumer demand, makes juice industry susceptible to adulteration. One of the most common frauds is to add a co-fruit to the authentic juice and, specifically, one of the most common practices is the adulteration of orange juice with tangerine, lemon and/or grapefruit. Therefore, the development of analytical methods to detect the adulteration of orange juices is of great interest to guarantee the food authenticity demanded by food producers, consumers and regulatory bodies. So far, approaches based mainly on conventional analytical techniques have been used to detect severe adulteration practices through the measurement of selected physicochemical indicators. However, they are often unable to detect small differences that could be indicative of low-level adulteration. On the other hand, methodologies based on chemical analysis combined with statistics have proved to be rather reliable, developing models for the determination of geographical origin or quality brand of foodstuffs and fraud detection. One of these approaches includes Liquid Chromatography-Mass Spectrometry (LC-MS) applied using both Reverse-Phase (RP) and Hydrophilic Interaction Liquid Chromatography (HILIC). HILIC is globally acknowledged as an emerging separation mode, being an attractive complementary tool to the widely used RP chromatographic separations in metabolomics.

In the present study, the potential of RPLC- and HILIC coupled to High Resolution Mass Spectrometry (HRMS) platforms is fully exploited to study the crucial authenticity issue of citrus fruit juices, applying a metabolomics approach. Integrated untargeted screening strategies were developed and applied to discover potential biomarkers for the reliable detection of orange juice adulteration with tangerine, grapefruit and lemon juice. Robust classification and prediction models were built with the use of both unsupervised and supervised techniques (principal component analysis (PCA) and partial least squares discriminant analysis (PLS-DA)), which were able to distinguish between citrus juices, as well as the orange juice adulteration down to 1% adulteration level. Characteristic m/z markers for different citrus juices were detected and several marker compounds were identified. The results obtained from this study clearly demonstrate that Mass Spectrometry (MS)-based metabolomics have the potential to be used as a reliable screening tool for the rapid determination of fruit juice adulteration. In conclusion, HILIC's unique capabilities in the determination of polar compounds undoubtedly enhance the comprehensiveness of metabolite detection, offering an additional depth of coverage in food authenticity metabolomic investigations.

Volatile-compound fingerprinting by both Headspace-SPME(HS-SPME) and static Headspace as a tool for the assessment of honey's authenticity

P.-L.P. Gialouris[1], G.A. Koulis[1,2], M.E. Dasenaki [2,1], N.S. Thomaidis[1]

[1] Laboratory of Analytical Chemistry, Chemistry Department, National and Kapodistrian University of Athens, Panepistimiopolis Zographou, 15771 Athens, Greece
[2] Laboratory of Food Chemistry, Chemistry Department, National and Kapodistrian University of Athens, Panepistimiopolis Zographou, 15771 Athens, Greece

Email: ntho@chem.uoa.gr

Abstract: Volatile compounds are a key aspect of honey as they provide its unique aroma[1]. Thus, they substantially influence honey's quality, along with consumers' preferences, in both national and worldwide markets. This study focuses on determining honey's volatile compounds for the characterization and botanical classification of Greek honeys based on their volatile profile. For this purpose, two methods, a static HS-GC-MS and a HS-SPME-GC-MS, were developed, optimized, validated and compared, in order to evaluate their suitability for analyzing the volatile composition of honey. 154 Greek honey samples from 12 different botanical origins (pine, fir, oak, thyme, heather, orange, chestnut, cotton, arbutus, carob, sage and vanilla) were collected from various geographical regions of Greece, during the harvesting period of 2019 and they were all analyzed with both methods.Compared to static HS, HS-SPME proved to be more efficient, recovering a wider variety of analytes, with a higher sensitivity. Target and suspect screening workflows, coupled with chemometric approaches, were then applied for the detection and quantification of volatile compounds.As a result, several biomarkers were identified and were evaluated as indicative of honey's botanical origin, such as Isophorone, which can be characterized as a potential marker of Arbutus honey.

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Determination of Vitamin D_2 in UV-B treated Mushrooms

D.Z. Lantzouraki[1,2], G. Sotiroudis[1], T. Tsiaka[1,2], G. Koutrotsios[3], N. Kalogeropoulos[4], G.I. Zervakis[3], P. Zoumpoulakis[1,2]

[1]Institute of Chemical Biology, National Hellenic Research Foundation 48, Vas. Constantinou Ave., 11635 Athens, Greece
[2]Department of Food Science and Technology, University of West Attica, Ag. Spyridonos, 12243 Egaleo, Greece
[3]Laboratory of General and Agricultural Microbiology, Department of Crop Science, Agricultural University of Athens, 11855 Athens, Greece
[4]Department of Nutrition and Dietetics, School of Health Science and Education, Harokopio University, 70 El. Venizelou Str., 17661 Athens, Greece

E-mail: dlantzouraki@eie.gr

Vitamin D deficiency is nowadays one of the most common nutritional deficiencies worldwide associated with a number of health risks in certain populations, such as hyperparathyroidism, rickets and osteomalacia. Among the few natural sources containing sufficient levels of vitamin D, commonly consumed mushroom species can provide substantial amounts of ergocalciferol (vitamin D_2) after natural or artificial UV-light treatment, and thus they are recognised as the only non-animal dietary source of vitamin D [1]. Within this framework, the present study aims to investigate the UV treatment as a means to rapidly increase vitamin D content in king oyster mushrooms (*Pleurotus eryngii*) in comparison to non-treated cultivated mushrooms. UV-B irradiation was used for the post-harvest irradiation under the controlled environment of a thermostabilized chamber. For the efficient recovery of vitamin D from the UV-B exposed mushrooms, an analytical protocol was developed, including a hot alkaline hydrolysis work-up with short extraction time and clean-up steps of the extract. Moreover, a selective and sensitive targeted Liquid Chromatography-Tandem Mass Spectrometric methodology was developed for the quantitative determination of the vitamin D forms (D_2 and D_3) in the studied samples. The LC-MS analysis revealed high ergocalciferol concentrations in the UV-treated mushrooms ranging from 50- to 120- fold higher vitamin D_2 compared to the control group.

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Elemental metabolomics nutritional value of Polish and Greek honeys - Mn, Cu and Mg

S.A. Drivelos[1], G. Danezis[1], M. Halagarda[2], S. Popek[2], C.A. Georgiou[1]

[1]Chemistry Laboratory, Department of Food Science and Human Nutrition, Agricultural University of Athens, 75 Iera Odos, 118 55 Athens, Greece

[2]Department of Food Product Quality, Cracow University of Economics, ul. Sienkiewicza 5, 30-033 Krakow, Poland

E-mail: cag@aua.gr

Honey is a supersaturated solution of sugars containing small amounts of proteins, enzymes, amino acids, trace elements, vitamins, aroma compounds and polyphenols. Very small amounts of trace elements such as Mn and Cu are essential for human health, while other elements such as Mg are required in relatively large amounts in a diet [1]. There is also a group of elements called ultra-trace elements, which are investigated for possible biological function. In order to prevent nutrient deficiencies, but also to reduce the risk of chronic diseases such as osteoporosis, cancer and cardiovascular diseases, scientific food committees around the world have set specific limits for the intake of each element at values adjusted to different population groups (children, adolescents, pregnant or elderly). The European Commission has set Nutritive Reference Values (NRV) for adults [2]. The World Health Organization (WHO)/ Food and Agriculture Organization (FAO) and the United States Department of Agriculture (USDA) have also established Recommended Dietary Allowances (RDAs) indicating the amount of an individual nutrient that people need for good health depending on their age and gender. According to FAO, 4.24 grams of honey are consumed daily per capita in Greece [3]. This value is one of the highest in the world. Based on the median of samples from Greece and Poland the contribution of honey on Mn, Cu and Mg NRV, adequate intake (AI) and RDAs of various population groups is small, ranging from 1.4% (AI Lactation 31–50 years of age for honeys from Greece) to 5.6% (NRV and AI Pregnancy 31–50 years of age for honeys from Poland) for Mn, from 1.0% (RDA Lactation 31–50 years of age for honeys from Poland and Greece) to 1.5% (RDA Males 31–50 years of age for honeys from Poland and Greece) for Cu and from 0.09% (RDA Males 31–50 years of age for honeys from Poland) to 0.31% (RDA Lactation 31–50 years of age for honeys from Greece) for Mg. Beyond nutritive elements Pb, as an example of toxic elements, has been also studied. The FAO / WHO Joint Expert Committee on Food Additives (JECFA) has established a Provisional Tolerable Weekly Intake (PTWI) for several toxic elements and especially heavy metals. Results for the typical 75 kg adult show 0.03% Pb intake from honeys of both countries.

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Utilising Head-Space Solid Phase Microextraction (HS-SPME) coupled to Gas Chromatography-Mass Spectrometry (GC-MS) for the classification of Greek wines based on their volatile composition

M. Marinaki[1,2,3], I. Sampsonidis[3,4], A. Lioupi[1,2,3], K. Zinoviadou[5], N. Thomaidis[3,6], G. Theodoridis[1,2,3]

[1] Laboratory of Analytical Chemistry, School of Chemistry, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[2] Biomic AUTh, Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center B1.4, 10th km Thessaloniki-Thermi Rd, P.O. Box 8318, GR 57001, Thessaloniki, Greece
[3] FoodOmicsGR Research Infrastructure, AUTh Node, Center for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center B1.4, 10th km Thessaloniki-Thermi Rd, P.O. Box 8318, GR 57001, Thessaloniki, Greece
[4] Department of Nutritional Sciences and Dietetics, International Hellenic University - P.O. 141, Sindos, 57400, Thessaloniki, Greece
[5] Perrotis College, American Farm School, Marinou Antipa 54, P.O. Box 60097, 570 01, Thermi, Thessaloniki, Greece
[6] Laboratory of Analytical Chemistry, Department of Chemistry, National and Kapodistrian University of Athens, Zografou 157 84, Athens, Greece

E-mail: marinakimaria2605@gmail.com

Wine is a complex mixture of hundreds of chemical species that are present at various concentration levels. Several factors, including geographical origin, grape variety and general vinification processes, contribute significantly to the production of this widely consumed beverage. Aroma is one of the most important quality attributes of wine. It consists mainly of volatile organic compounds (VOCs - also referred as the volatilome) that can be arranged in several chemical classes such as terpenes, esters, alcohols, acids, alkanes, aldehydes etc. The study of the wine volatilome is part of wineomics, which is the study of wine metabolites (also part of Foodomics, which is the application of -omics technologies in the Food and Nutrition domains). Several cutting-edge analytical techniques can be used to study the wine volatilome including Gas Chromatography - Mass Spectrometry (GC-MS) in combination with Headspace-solid phase microextraction (HS-SPME) [1,2].

In the present work, arduous method development was applied in order to reach optimal HS-SPME conditions to assay the volatilome of a collection of Greek wine samples. Design of Experiments (DOE)assisted optimization of the analysis. Nine different Greek grape varieties, seven white and two red, and six different regions were analysed. Raw GC-MS data was processed using specific tools and multivariate statistical analysis techniques, such as principal component analysis (PCA) and Partial Least-Squares Discriminant Analysis (PLS-DA) were applied to the resulting datasets in order to evaluate sample classifications.

Preliminary results indicate that the classification of samples according to grape colour and variety is achievable. The possibility of determining the geographical origin of wines of the same variety is also highlighted.

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Simplifying edible oil analysis by automating the transesterification process through the development of a 3D-printed continuous flow system

A. du Preez, B. Khuzwayo, R. Meijboom, C. Arderne, E. Smit

Research Centre for Synthesis and Catalysis, Department of Chemical Sciences, University of Johannesburg, PO Box 524, Auckland Park, Johannesburg, South Africa

E-mail: esmit@uj.ac.za

Automation introduces a myriad of advantages to sample preparation and processing steps. However, the implementation of automated procedures is often hindered by high costs. Herein, a continuous flow system was developed to automate the base-catalyzed transesterification of triacylglycerols in edible oils. The costs involved were reduced drastically by manufacturing the main flow system components (e.g., syringe pumps, connectors, and reactors) using 3D printing technology (see Figure 1).

The automated flow method was compared to manual batch protocols. Internal standards were used to ensure that transesterification was complete and that saponification did not occur. Notably, lower catalyst concentrations (1 – 2 wt. %) were required to achieve complete transesterification in the continuous flow method. The 3D-printed flow system was successfully used to determine the fatty acid content of nine commercial edible oils and four oil blends. Results obtained, expressed as a percentage of total fatty acid methyl ester (FAME) content, were comparable with certified and literature values. Furthermore, statistical analysis indicated that the peak area ratios of observed FAMEs could be used to screen for the authenticity of avocado oil, macadamia oil, and olive oil.
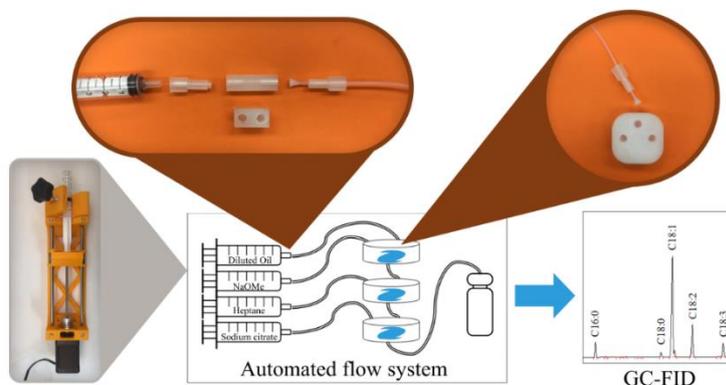
Figure 1. Automated 3D printed flow system for the transesterification of triacylglycerols in edible oils.

Lipidomics investigation of traditional Greek yoghurt skin (petsa) utilising RP-UHPLC-TOFMS following exhaustive liquid-liquid extraction

A. Pesiridou[1,2,3], I. Sampsonidis[1,4], S. Kalogiannis[1,4], T. Moschakis[5], A. Pechlivanis[1,2,3], G. Theodoridis[1,2,3]

[1]FoodOmicsGR_Research Infrastructure, Auth Node, CIRI-AUTH, Thessaloniki, Greece
[2]Department of Chemistry, Aristotle University of Thessaloniki, Thessaloniki, Greece
[3]Biomic_Auth, Bioanalysis and Omics Laboratory, CIRI-AUTH, Thessaloniki, Greece
[4]Department of Nutritional Sciences & Dietetics, International Hellenic University, Thessaloniki, Greece
[5]Department of Food Science and Technology, School of Agriculture, Aristotle University of Thessaloniki, Thessaloniki, Greece

E-mail: anapesiridou@gmail.com

Nowadays, food products of high nutritional value are becoming more popular and so is yoghurt, which is one of the most widely consumed dairy products. Yoghurt is a food rich in proteins, probiotics, vitamins, minerals and potentially beneficial fats, such as conjugated linoleic acid (CLA). Lately, yoghurt has gained a lot of popularity due to its various proposed health benefits and therapeutic properties [1].

Traditional Greek yoghurt is prepared following a traditional recipe, directly inside the final container, without applying any standardisation or homogenisation of milk, resulting in the formation of a thin layer at the surface of the yoghurt. This top layer, also called skin or in Greek known as "petsa", is rich in fat and provides the traditional Greek yoghurt with special organoleptic characteristics that make it significantly different from any other yoghurt types [1].

Since traditional Greek yoghurt's skin is not homogenous, a modified lipid extraction technique, based on liquid-liquid extraction (based on methyl tert- butyl ether) was developed, in order to achieve reliable lipid extraction. After solvent evaporation, the extracts were resuspended in a generic lipidomics solvent system and analysed using RP-UHPLC-TOFMS, in both positive and negative ionisation modes, employing a generic and well-established lipidomics protocol[2]. Chemometric analysis and lipid identification were performed using MS-Dial[3] and related lipids investigation software. To the best of the authors' knowledge, a thorough lipidomic profiling of the characteristic yoghurt's skin has not yet been performed. Results indicate a highly diverse lipid profile, worthy of further investigation, especially since it is derived from non-homogenized milk.

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The determination of acrylamide in baby foods as an exemplary case study for the role of metrology in food safety

C. Alexopoulos[1], A. Georgopoulou[2], P. Giannikopoulou[1], E. Kakoulides[1], E. Skotidaki[1], E. Stathoudaki[1], V. Schoina[1]

[1]EXHM/GCSL-EIM, Chemical Metrology Service, General Chemical State Laboratory, 16 A. Tsocha Street, 115 21 Athens, Greece
[2]Division of Food and Alcohol, General Chemical State Laboratory, 16 A. Tsocha Street, 115 21 Athens, Greece

E-mail: metrology.gcsl@aade.gr

Metrology institutes constitute the pinnacle of the traceability chain for measurements. Their role is to provide reference procedures and materials that are characterised by the highest accuracy coupled to the lowest possible uncertainty, providing a link to the SI. In the field of food safety, such methods and materials are of vital importance, considering the requirements for monitoring processing contaminants in food at lower concentration levels. For the past two decades acrylamide has drawn considerable attention due to its toxicity and its presence in food undergoing heat processing [1]. Stringent performance requirements for its determination in matrices such as baby foods have been recently introduced in the European legislation [2]. Using this as an exemplary case study and working within the context of the EU research infrastructure METROFOOD-RI [3], EXHM/GCSL-EIM developed, validated and accredited a new method for the determination of acrylamide in baby food and infant formula. The method employs liquid-liquid extraction, dSPE clean-up and quantification by LC-IDMS. Using the new method, a five-fold reduction of the limit of detection was made possible, bringing the limit of quantification to levels lower than 5 μg/kg. The procedure was proved to be equivalent to 'traditional' approaches employing solid phase extraction, since it showed comparable performance in a relevant interlaboratory trial and was introduced in the 2021 EURL-PC meeting. The new method has the potential to characterize materials accurately with uncertainties that are fit for the certification of reference materials at acrylamide concentrations lower than the current benchmark levels.

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Optical Spectroscopic Techniques in Detecting Fuel Adulteration

N. Fragkoulis[1,2*], M. Velegrakis[1], P. Samartzis[1]

[1] Institute of Electronic Structure and Laser (IESL) – Foundation for Research and Technology-Hellas, (FORTH), N. Plastira 100, Vassilika Vouton, 70013 Heraklion Crete, Greece

[2] Department of Chemistry – University of Crete, Panepistimioupoli Vouton, 70013 Heraklion Crete, Greece

*E-mail: nfragkoulis@iesl.forth.gr

Adulteration of fuels is a problem of high importance to the economy and to the environment. Spectroscopic techniques, such as FTIR, UV-NIR absorption, Raman and others, have been used to quantify adulteration. In this contribution we present our results from the comparison of four spectroscopic techniques (FTIR, UV absorption, NIR absorption and fluorescence) in detecting four common types of fuel adulteration: unleaded in super unleaded gasoline, heating diesel in car diesel, marine diesel in car diesel and lubricants in car diesel. The spectroscopic data were analyzed using multivariate statistical methods [Partial Least Squares (PLS) Regression]. Fluorescence, UV and NIR absorption exhibit R^2 values above 93% for car diesel adulteration with marine or heating diesel, while adulteration of super with unleaded gasoline is best quantified with fluorescence.

This study demonstrates the potential of optical spectroscopy as a fast and low-cost probe of fuel adulteration.

Fatty acids composition of traditional, strained and set type Greek yogurt, manufactured from cow's, sheep's and goat's milk.

M. F. Kontodimou *[1], E. Zoidou [1], G. Papadopoulos [2], T. Massouras [1]

[1]Laboratory ofDairy Research, Agricultural University of Athens, IeraOdos 75, Votanikos, 11855 Athens, Greece
[2]Laboratory ofPlant Breeding and Biometry, Agricultural University of Athens, IeraOdos 75, Votanikos, 11855 Athens, Greece

*E-mail: marla_kon@hotmail.com

Greek yogurt is one of the most popular fermented milk products in the world. The profile of fatty acids is known to influence the nutritional and health prospects of the respective product. The objective of this study was to investigate the fatty acid profile of traditional, strained, and set type Greek yogurt, manufactured fromcow's, sheep's and goat'smilk. One hundred and eighteen (118) Greek yogurt samples were collected from the Greek market and analyzed for fatty acid composition by Gas chromatography.One-way analysis of variance (ANOVA) and multiple range tests were applied, to determine significant differences among yogurt samples, regarding yogurt type, kind of milk, and fat content. A total of 25 fatty acids were identified. Palmitic acid wasthe most abundant fatty in all samples, followed by oleic, stearic, and myristic acid.Significant differences for saturated (SFA); C6: 0, C8: 0, C10: 0, C11: 0, C12: 0, C13: 0, C16: 0, C18: 0, C20: 0 and the unsaturated (UFA); C14: 1 , C16: 1, C17: 1, C18: 2n6t, C18: 3n3, CLA, C20: 3n6 fatty acidswere observed between traditional, strained, and set type yogurt. The kind of milk showed a significant effect(P-Value <0,001)in the profile of all the examined saturated fatty acids as well as most of the unsaturated. Significant differences(P-Value <0,001) were also observed infat content for the (SFA) C4: 0, C6: 0, C8: 0, C10: 0, C12: 0, C13:0, C15: 0, C16: 0 , C20: 0 and the (UFA) C14: 1, C16: 1, C17: 1, C18: 1n9t, C18: 3n3, CLA, C20: 3n6 fatty acids. Additionally, a comparison of specific nutritional indices between yogurt from cow, sheep, and goat milk in terms was attempted.The milk type did not significantly affect the sum of SFA. In contrast, a higher percentage of MUFA and PUFA fatty acids was found in yogurt made from sheep milk. Finally, the n6: n3 ratio of fatty acids was found to be closer to the desired values in sheep and goat milk yogurt rather than in cow milk yogurt.

Metabolomic profiling of women with thyroid disorder

Ch. Fotakis[1], G. Moros[1], A. Kontogeorgou[2], N. Iacovidou[2], Th. Boutsikou[2], P. Zoumpoulakis[1,3]

[1] Institute of Chemical Biology, National Hellenic Research Foundation, 48 Vas. Constantinou Ave., 11635, Athens, Greece

[2] Department of Neonatology, Medical School, National and Kapodistrian University of Athens, Greece

[3] Department of Food Science and Technology, Univeristy of West Attica, Ag. Spyridonos, Egaleo, 12243, Athens, Greece

E-mail: pzoump@eie.gr

The importance of thyroid hormones (THs) is well established in preserving energy homeostasis, regulating significant metabolic processes, fetal development as well as programming and functioning of the mature thyroid gland. Evidently, it is essential to frequently monitor thyroid function during pregnancy, investigate how it affects the metabolic profile and apply therapeutic interventions on indication. Serum blood provides a rational compartment to interpret metabolite variations. In this context, we implemented an NMR-metabolomics approach on ninety maternal blood serum samples with both controlled and uncontrolled (overactive and underactive) thyroid. We extracted multivariate statistical analysis models (PCA, OPLS-DA) to pinpoint metabolite differences and detect biomarkers. Resonance peak identification was accomplished by facilitating an in-house fully automated metabolite identification platform, Metaboneer[1] and a series of 2D experiments (gCOSY, zTOCSY, gHMBCad, gHSQCad).

We ascribed the role of biomolecules such as unsaturated lipids, cholesterol-VLDL, LDL2/VLDL2, n-acetylated glycoproteins as well as metabolites such 1-methylhistidine and methionine to the hypothyroidism specimens, thus verifying the relation of abnormal lipid panels to patients with hypothyroidism. NMR metabolomics proves an invaluable tool for precision therapy, as it enables the monitoring of therapy response, allowing clinicians to act when the thyroid despite the medication remains unregulated. Our results highlight the impact of hypothyroidism on metabolites' composition during pregnancy.

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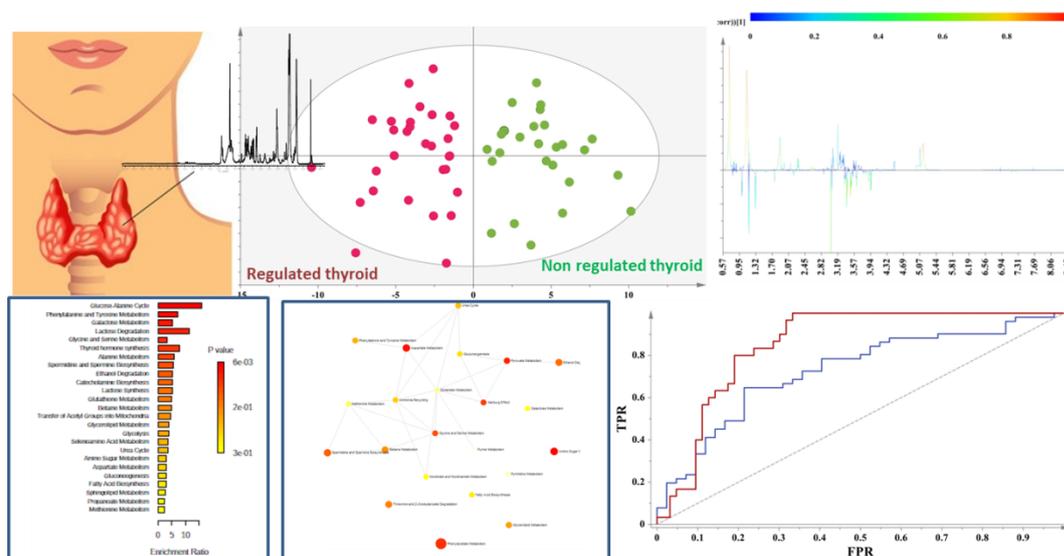
Multiresidue analysis of pharmaceuticals and pesticides in fish feed by QuEChERS method coupled to liquid chromatography-high-resolution Orbitrap mass spectrometry

P.P. Martinaiou, V.I. Boti, T.A. Albanis

Laboratory of Industrial Chemistry, Department of Chemistry, University of Ioannina, 45110 Ioannina, Greece

E-mail: vboti@uoi.gr

The extensive and ubiquitous use of emerging contaminants especially in aquaculture sector is a matter of major importance due to the consequences to human health, ecological systems and food safety [1]. Thus, it is crucial to comprehensively investigate the occurrence of pharmaceuticals and pesticides in fish feeds so as to assess the potential impacts on aquatic ecosystems. In the present study, an accurate and quick quantitation QuEChERS (Quick, Easy, Cheap, Effective, Rugged, and Safe) method for the analysis of the targeted categories of emerging contaminants in fish feed matrices, had been developed and validated. It was followed by ultra high-performance liquid chromatography- high- resolution linear ion trap/ Orbitrap mass spectrometry (UHPLC-LC-HR-IT/Orbitrap-MS). After the selection of a modified AOAC 2007.01 QuEChERS [2] method for pharmaceuticals and antibiotics analysis and the EN15662 [3] one for pesticides, crucial parameters were optimized such as the extraction solvent and the clean-up sorbents identity and amounts. The methods were then validated in terms of recoveries, limits of quantitation (LOQ) and detection (LOD), methods linearity, and precision at three spiking levels (40, 80 and 160 ng g^{-1}). The matrix effect was also estimated. Most of the analytes achieved good recoveries (49.2 to 127.5%). Limits of quantification were below 25 ng g^{-1} for the target pesticides and below 50 ng g^{-1} for the selected antibiotics and pharmaceuticals group. Linearity, expressed as a correlation coefficient, was greater than 0.9954 in all cases. Over of 95% analytes achieved satisfactory repeatability and reproducibility results with relative standard deviations (RSDs) <12.4% and 19.3% for the pesticides and antibiotic/pharmaceuticals group, respectively. The matrix effect study revealed signal suppression for most of analytes. The validated methodologies were then applied to the determination of the target compounds residues in fish feed samples of different composition.

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Analysis of Emerging Contaminants in seawater and aquaculture fish, using liquid chromatography-high-resolution Orbitrap mass spectrometry

P. P. Martinaiou, V. I. Boti, T. A. Albanis

Laboratory of Industrial Chemistry, Department of Chemistry, University of Ioannina, 45110 Ioannina, Greece

E-mail: vboti@uoi.gr

One of the main risks associated with the marine environment and specifically with aquaculture waters, is the occurrence of emerging contaminants such as pharmaceutical and pesticide residues [1][2]. To evaluate this occurrence, this study points at the development and validation of accurate and quick quantitation methods for the analysis of these categories of emerging contaminants, with a focus on seawaters and fish tissues. In the present survey, the SPE and QuEChERS (Quick, Easy, Cheap, Effective, Rugged, and Safe) methods had been optimized, validated, and applied in seawater and fish respectively, followed by ultrahigh-performance liquid chromatography- high- resolution linear ion trap/ Orbitrap mass spectrometry (UHPLC-LC-HR-IT/Orbitrap-MS). After the selection of the most suitable methods not only for pharmaceuticals and antibiotics analysis but also for pesticides, some crucial parameters were optimized and then, the methods were validated in terms of recoveries, limits of quantitation (LOQ) and detection (LOD), methods linearity, and precision at three spiking levels. The matrix effect was also estimated. Most of the analytes achieved good recoveries. Linearity, expressed as a correlation coefficient, was greater than 0.99 in all cases. Most of the analytes achieved satisfactory repeatability and reproducibility results. Thus, the methods were found to comply with current European legislation [3]. The proposed methodologies were successfully applied to the determination of the target compounds residues in seawater and aquaculture fish samples after a one-year monitoring study.

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Investigation of Furaltadone Hydrolytic and Photolytic degradation in aqueous environmental systems

C. V. Efthymiou, V. I. Boti, T. A. Albanis

Laboratory of Industrial Chemistry, Department of Chemistry, University of Ioannina, 45110 Ioannina, Greece

E-mail: vboti@uoi.gr

Nitrofurans are a series of pharmaceutical substances used as therapeutic agents to promote animal health, but they may provoke significant effects to natural ecosystems, and in many cases, potential harmful issues on human health. Furaltadone (FTD) is an antibiotic drug that belongs to the nitrofuran group. It is used as an antibacterial agent to prevent bacterial and protozoan infections. The parent compounds, like FTD, are metabolized resulting in stable metabolites that exhibit adverse effects as mutagenesis, teratogenesis and carcinogenesis in humans. For this reason, FTD, along with other nitrofurans, were banned from use in the EU in 1993 [1]. Nevertheless, due to the cost-effectiveness of nitrofurans, they have been produced and supplied in many developing countries. For example, imported aquaculture, poultry and pork products consist of nitrofuran residues and metabolites from FTD [2]. Despite the interest, a few studies focus on the fate of furaltadone residues in the environment. This study aims to investigate the behaviour of FTD in aquatic systems like lakes, rivers and sea, by observing its hydrolysis and photolytic degradation. For the hydrolysis determination, samples of FTD were observed in various pH values (4-10) for 30 days. In order to simulate degradation processes that take place in the aquatic environment, photodegradation experiments were carried out under natural artificial solar irradiation conditions. Photodegradation under simulated irradiation conditions was performed using a Suntest CPS + apparatus from Heraeus (Hanau, Germany) equipped with a xenon arc lamp (1500 W) at an irradiance of 750 W/m^2, as well as special glass filters which blocked the transmission of wavelengths below 290 nm. FTD concentrations were determined by an LC-UV/Vis system from Shimadzu (Kyoto, Japan) using a gradient program. Concerning hydrolysis, results showed that this occurred for FTD after 30 days and only for pH 4 and 10. On the other hand, direct photolysis of FTD in distilled water and buffered solution occurred in both cases in less than one hour indicating the liability of this compound to photodegradation as the main degradation route in the aquatic environment.

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Determination of pharmaceuticals and metabolites in sewage sludge using a modified QuEChERS extraction and UHPLC LTQ/Orbitrap MS

K. Miserli[1], C. Kosma[1], I. Konstantinou[1,2]

[1]Department of Chemistry, University of Ioannina, 45110, Greece
[2]University Research Center of Ioannina (URCI), Institute of Environment and
Sustainable Development, Ioannina 45110, Greece

E-mail: iokonst@uoi.gr

In the last years, emerging contaminants have become a popular research venue due to their devastating effects on human health as well as on the environment. Pharmaceuticals (PhACs) are an important group of emerging contaminants that released continuously in the environment from wastewater treatments plants (WWTPs) and they can have a biological response even though at very low concentrations [1]. Sewage sludge is produced as a solid matter in primary and secondary treatments. However, conventional WWTPs are not able to remove or degrade completely emerging pollutants resulting in the presence of PhACs in sewage sludge after wastewater treatment. PhACs are found in sludge at low ppb-ppt levels and their analysis and detection is a difficult task due to the complexity of sewage sludge matrices [2]. A modified quick, easy, cheap, effective, rugged and safe (QuEChERS) based methodology with a dispersive solid-phase extraction (d-SPE) clean-up (Figure 1) followed by ultra-high-performance liquid chromatography–high-resolution linear ion trap Orbitrap mass spectrometry (UHPLC-LTQ/Orbitrap MS), operated in positive ionization mode, was adopted to investigate 33 multiclass pharmaceuticals in sewage sludge. The analytical method was validated in terms of linearity, recovery, intra and inter-day precisions, uncertainty (%U)/Horrat ratio at three spiking levels, matrix-effects (ME), process efficiency (PE) and limits of detection and quantification (Table 1). The developed methodology was applied to sludge samples from the WWTP of Ioannina city and 26 from the 33 PhACs were detected at concentrations up to 19.1 ng g^{-1} dw. Amisulpride, clozapine and amitriptyline were detected at higher concentration levels up to 506, 135 and 239 ng g^{-1} dw, respectively.

Table 1: Method performance parameters for spiking concentration level 100 ng g^{-1}

	%Recovery (n=5)	LOD (ng g^{-1})	%RSD (n=5)	R^2	%U	%ME	%PE
33 PhACs	62-111	0.5-9.5	2-14	>0.9912	5.3-23.3	<±20	55-96

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Figure 1: QuEChERS extraction.

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Extraction step — Clean up step
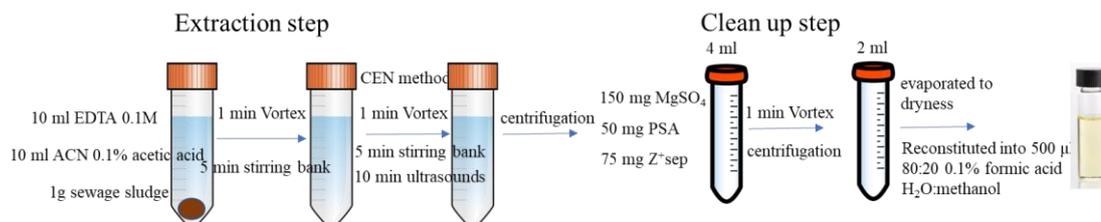
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Photocatalytic degradation of antibiotic pharmaceuticals using nitrogen-modified TiO$_2$ nanostructures

P. P. Falara[1,2], M. Arfanis[2], K. Kordatos[1]

[1] School of Chemical Engineering, National Technical University of Athens, 9 Iroon Polytechniou St., 15780 Zografou, Athens, Greece.

[2] Institute of Nanoscience and Nanotechnology, National Center for Scientific Research "Demokritos", 15341 Agia Paraskevi, Athens, Greece.

E-mail: pin.falara@gmail.com, kordatos@mail.ntua.gr

Pharmaceuticals' increasing detection in natural and wastewater bodies is becoming a worldwide environmental issue and threat to human health owing to their solubility, persistence and high toxicity. Among the existing water treatment methods, heterogeneous TiO$_2$ photocatalysis is one of the most promising and viable technologies to tackle aqueous pollution. Remarkably, the combination of adsorption and photocatalysis is a promising strategy to diminish the concentration of hazardous pharmaceuticals in wastewater without causing secondary pollution. Doping TiO$_2$ with various elements (e.g. nitrogen (N), carbon (C), sulfur (S) and halides) can radically broaden and shift the photoresponse and thus significantly enhance the photocatalytic activity. Herein, nitrogen-modified TiO$_2$ nanostructures were synthesized by the gel combustion method based on the calcination of an acidified alkoxide solution, using urea as nitrogen source. The materials were thoroughly characterized and successfully tested for the adsorption and photocatalytic degradation of three model antibiotic pharmaceuticals of emerging concern, ciprofloxacin, tetracycline and oxytetracycline dehydrate. The three pharmaceuticals follow first order degradation kinetics and their initial reaction constants were determined as 0.047, 0.028 and 0.045 min^{-1}, respectively.

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Micropollutants' degradation through heterogeneous catalytic ozonation - application of continuous flow pre-industrial unit

E. Kaprara[1], S. Psaltou[2], Ch. Nannou[2,3], M. Salapasidou[4], P.-A. Palasantza[4], P. Diamantopoulos, D. Lambropoulou[2,3], A. Zouboulis[2], M. Mitrakas[1]

[1] Department of Chemical Engineering, Aristotle University of Thessaloniki, GR 54124, Thessaloniki, Greece
[2] Department of Chemistry, Aristotle University of Thessaloniki, GR 54124, Thessaloniki, Greece
[3] Centre for Interdisciplinary Research and Innovation (CIRI-AUTH), Balkan Center, Thessaloniki, 10th km Thessaloniki-Thermi Rd, GR 57001
[4] AKTOR S.A., Wastewater Treatment Plant of Touristic Area of Thessaloniki "AINEIA", N. Michaniona, Thessaloniki, Greece

E-mail: manasis@eng.auth.gr

The aim of this study was to evaluate the potential of heterogeneous catalytic ozonation for micropollutants removal from wastewater through a continuous flow system. For this purpose, a pre-industrial level unit was designed, constructed and operated at AINEIA's municipal wastewater treatment plant (WWTP) established in N. Michaniona-Thessaloniki, Greece. The unit comprised of a column packed with sand for the preliminary post-filtration and removal of suspended solids, a custom-made porous PTFE hollow fiber membrane (provided by Markel Corporation, PA, USA) for the contact/dilution of ozone gas to water, a column packed with an appropriate solid material (zeolite, PET), applied as potential catalyst for the ozonation treatment of micropollutants and a sand/activated carbon column for the final biological stabilisation of treated wastewater. The secondary (biologically treated) effluent of WWTP was introduced into the pilot unit at a flow rate of 200 L/h, while ozone gas supply was provided by an existing generator, used to disinfect the treated wastewaters before disposal to the sea, which can produce 190 g O_3/Nm^3. Micropollutants concentrations were determined at the influent of WWTP, the secondary effluent before and after conventional disinfection (by ozonation) and at different treatment stages of the pilot unit, by using Liquid Chromatography–High Resolution Mass Spectrometry. In total, pharmaceuticals and personal care products (PPCPs), belonging to 38 different therapeutic classes were studied. Out of them, compounds belonging to 27 different classes were detected. The highest mean effluent concentrations were observed for antihypertensives (366.4 ng/L), insect repellents (247.85 ng/L), antifungals (96.50 ng/L), antihistamines (42.38 ng/L), caffeine (37.98 ng/L) and adrenergic receptors (23.60 ng/L). Mean concentrations for the rest of compounds ranged below 20 ng/L. Degradation rates obtained by the application of conventional disinfection (through ozonation) ranged between 32% and complete removal, depending on the specific contaminant, with an average value of 88%. Experimental results from the treatment of secondary effluent in the pilot unit revealed that part of micropollutants was removed through filtration of residual suspended solids (i.e. an average of 42%), while degradation carried out at catalytic ozonation column led to complete removal of them, when applying zeolite as catalyst, and an average of 80% removal by using PET. In the latter case, the biological stabilisation of the effluent after catalytic ozonation led to complete degradation for the majority of examined contaminants and an average removal rate of about 90%. Comparison of micropollutants' residual concentrations after the application of conventional (single) and catalytic ozonation indicated the enhancement of contaminants removal through ozonation by the introduction of a proper solid material, acting as catalyst. Specifically, application of zeolite led to smaller or equal residual concentrations for the 95% of examined contaminants, while for the case of PET use the respective percentage was 82%.

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Enhanced techniques for sensitive immunochromatographic detection of antibiotics and their application for control of honey

D.V. Sotnikov, L.V. Barshevskaya, A.V. Zherdev, B. B. Dzantiev

A.N. Bach Institute of Biochemistry, Research Center of Biotechnology of the Russian Academy of Sciences, 119071 Moscow, Russia

E-mail: boris.dzantiev@mail.ru

Immunochromatographic (lateral flow) test systems have become widespread and are successfully used in medical diagnostics, food quality control, environmental monitoring, etc. Immunochromatographic analysis (ICA) allows the quick and easy screening of a large number of samples without the use of complex equipment or the involvement of trained personnel. However, the need for high sensitivity in the detection of many significant toxicants and biomarkers limits the use of ICA in its traditional form. Therefore, the development of approaches for increasing the sensitivity of ICA is in high demand. This work presents two such approaches implemented for the determination of antibiotics in honey as an example of contaminated food.

A competitive form of ICA with the indirect introduction of the marker into the registered immune complexes was implemented for the determination of sulfanilamide (SA). The indirect introduction technique allows one to overcome the limitations of traditional competitive ICA caused by the impossibility of independently varying the concentrations of the marker and specific antibodies immobilized on nanoparticles. When switching to indirect labeling, instead of one reagent (i.e., a marker with specific antibodies), two independent reagents (i.e., free specific antibodies and conjugates of nanoparticles with immunoglobulin-binding protein G) are used. The developed ICA of sulfanilamide was tested in the control of honey samples. With digital registration, the SA detection limit was 0.37 ng/mL. The visual limit of detection, corresponding with the disappearance of staining in the analytical zone of the test strip, was 100 ng/mL. The achieved parameters correspond to regulatory demands for the maximum allowable content of sulfonamides in foods (0.1 mg/kg). Note the versatility of the used conjugate of gold nanoparticles with protein G, i.e., its applicability in the detection of various compounds.

The second proposed method of ICA provides for two stages of competitive binding of an analyte to antibodies. The corresponding test system was developed for the determination of chloramphenicol and tested in the control of contaminated honey samples. The visual detection limit for chloramphenicol was 1 ng/mL, which is 10 times lower compared to the limit for traditional ICA based on the same immunoreagents. With digital registration, the detection limit of chloramphenicol was 0.01 ng/mL.

The proposed ICA techniques, in combination with the use of portable video digital detectors for the quantitative assessment of the amount of formed immune complexes, can be used for the highly sensitive, simple, and economical determination of various low-molecular-weight compounds.

Insights into the impact of 2,6-dichloro-1,4-benzoquinone and 2,6-dichloro-3-hydroxy-1,4-benzoquinone on human keratinocytes metabolism through an NMR metabolomic study

D. G. Meintani[1], T. G. Chatzimitakos[1], A. I. Kasouni[2], C. D. Stalikas[1]

[1]Laboratory of Analytical Chemistry, Department of Chemistry, University of Ioannina, 45110 Ioannina, Greece, E-mail: cstalika@uoi.gr
[2]Laboratory of Biophysical Chemistry, Department of Biological Applications and Technologies, University of Ioannina, 45110 Ioannina, Greece

E-mail: cstalika@uoi.gr

The 2,6-dichloro-1,4-benzoquinone (DCBQ) and its derivative 2,6-dichloro-3-hydroxy-1,4-benzoquinone (DCBQ-OH) are disinfection by-products and emerging pollutants in the environment. In this study, HaCaT cells were exposed to concentrations that induced their death at an average of 20% (IC_{20}) and 50% (IC_{50}) for the two substances, and a semiquantitative metabolomic study was performed using ^{1}H-NMR. For DCBQ-OH, an exposure of cells was carried out in DMEM at pH=7, for 24 h. To avoid transformation to its hydroxy-derivative the more cytotoxic DCBQ (it has lower IC_{50}) was studied in DMEM at pH=5, at the limited exposure time of 30 min. Differences in the metabolic pathways were recorded. In the control samples of DCBQ-OH, 21 metabolic pathways were recorded, whereas 26 and 20 pathways were detected when exposed to IC_{20} and IC_{50}, respectively. Some metabolic pathways were deactivated while others were activated. Changes like the activation of alpha-linolenic acid, biotin and glutathione metabolism, occurred. Alpha-linolenic acid is a cell membrane component, biotin takes part in gluconeogenesis [1] and glutathione is a scavenger of reactive oxygen species [2]. Moreover, the deactivation of glycolysis/gluconeogenesis in IC_{50} was recorded, among others. A tentative quantification of metabolites revealed increased concentration of the metabolites when cells are exposed to IC_{20} and a decreased one when exposed to IC_{50}, suggesting their inability to function properly. In the control samples of DCBQ, 22 metabolic pathways were recorded, whereas 31 and 30 pathways were recorded when the cells were exposed to IC_{20} and IC_{50}, respectively. Some of the changes include the activation of inositol phosphate, lipoic acid and riboflavin metabolism, among others. Inositol phosphate is involved in transferring messages from the receptors of membrane to the interior [3], lipoic acid is a free radical scavenger and riboflavin is an energy generator[4]. In contrast, biotin metabolism was deactivated. The metabolites found in all samples exposed to DCBQ, indicate a dose dependent decrease of saccharides which provide cells with energy. Steroids increase when cells are exposed to IC_{20} and decrease when exposed to IC_{50}. It can be concluded from the above that DCBQ-OH and DCBQ induce oxidative stress and membrane damage. Although both chemical factors stress cells, exposure to DCBQ leads cells to activate the transport of messages for better adaption and demands of energy are higher, which leads to the activation of riboflavin and the consumption of saccharides. The aforementioned pathways as well as others of equal or less importance were noticed in cells exposed to DCBQ. Our findings underline the importance to further study their impact on human.

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Unbiased and class-independent optimization of Headspace Solid Phase Micro-Extraction conditions for the discovery of volatile classification markers in grape marc spirits using the Response Surface Methodology

I. Sampsonidis[1,2], M. Marinaki[2,3,4], A. Nakas[5,6], A.N. Assimopoulou[5,6], S. Kalogiannis[1,2] G. Theodoridis[2,3,4]

[1]Department of Nutritional Sciences & Dietetics, International Hellenic University, Thessaloniki, Greece;
[2]FoodOmicsGR_Research Infrastructure, Auth Node, CIRI-AUTH, Thessaloniki, Greece
[3]Department of Chemistry, Aristotle University of Thessaloniki, Thessaloniki, Greece
[4]Biomic_Auth, Bioanalysis and Omics Laboratory, CIRI-AUTH, Thessaloniki, Greece
[5]Laboratory of Organic Chemistry, School of Chemical Engineering, Aristotle University of Thessaloniki, Thessaloniki, Greece
[6]Natural Products Research Centre of Excellence (NatPro-AUTh), CIRI-AUTh, Thessaloniki, Greece

E-mail: isampson@nutr.teithe.gr

Optimization of extraction conditions in analytical chemistry is traditionally performed following the one-factor-at-a-time (OFAT) approach while consulting the literature or previously acquired in-house knowledge. The OFAT approach, however, apart from being time and labour intensive, does not provide the practitioner with knowledge on factor interactions and may even lead to false optima. On the contrary, the design of experiments approach (DOE) allows for more accurate optima detection, alongside the acquisition of process specific knowledge while performing a minimum number of experimental runs [1,2].

Optimization of Headspace Solid Phase Micro-Extraction (HS-SPME) coupled to Gas Chromatography Mass Spectrometry (GC-MS) has been performed in the past using DOE by a number of practitioners on a number of matrices. However, in most of the aforementioned approaches such optimization is focused on specific compounds, or compound classes, known to be present and/or abundant in the corresponding matrices [3,4]. However, during untargeted analysis and marker discovery such approach may apply bias on the optimization of the extraction process in favour of the selected compounds and classes, while it may suppress the optimization of unknowns that could provide valuable information, especially for classification purposes.

Here we apply a Box–Behnken Response Surface Methodology (RSM) design for the optimization of 3 HS-SPME factors on a pooled grape marc spirit sample that is representative of a large pool of samples. Following data processing, individual features are extracted from raw chromatograms under specific settings and individual models are fitted to each feature. Following model validation criteria, features are included in the finalised, aggregate response upon which a model is also fitted and factor optima are provided. A validated model indicates that each grape marc spirit sample is analysed in an optimal manner, thus providing opportunities for sample classification marker discovery.

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Multiresidue QuEChERS analysis for the detection of pharmaceuticals and their metabolites in aquaculture sediments using UHPLC LTQ/Orbitrap MS

A. Kalogeropoulou[1], C. Kosma[1], T. Albanis[1,2]

[1]Laboratory of Analytical Chemistry, Department of Chemistry, University of Ioannina, 45110 Greece
[2]University Research Center of Ioannina (URCI), Institute of Environment and Sustainable Development, Ioannina 45110, Greece

E-mail address: aggelkaloge@yahoo.gr

Emerging pollutants (EPs) are chemical substances widely produced and consumed due to the rapid urbanization of free land, technological growth and human quality of life [1]. Agricultural and urban areas emit emerging contaminants potentially hazardous to the aquatic environment that may affect human health as well, such as Pharmaceutical Active Compounds (PhACs) [2,3]. Sediments play an important role in aquatic ecosystems, as habitats for a variety of aquatic organisms and they are regarded as a primary sink of organic pollutants. Aquacultures around the world are gaining more interest due to their fast growth food production, and as a result, it is essential to investigate the occurrence, distribution and fate of PhACs in the aquatic ecosystems [4]. Thus, the aim of the present study was to develop a multi-residue analytical method for the determination of traces of 22 multiclass pharmaceuticals in aquaculture sediments. The procedure based on the quick, easy, cheap, effective, rugged and safe (QuEChERS) extraction method, followed by ultrahigh-performance liquid chromatography-high-resolution linear ion trap/Orbitrap (LTQ Orbitrap) mass spectrometry, operated in positive ionization mode. In order to optimize process efficiency, three different methods (original, acidified and buffered) were tested, using different solvents, extraction salts, sediment amount and clean-up salts, as well. A matrix-matched calibration approach and isotopically labeled internal standards were employed for quantification and correction of matrix effects. The optimized method was validated in terms of linearity, recovery, precision, limit of detection (LOD) and quantification (LOQ) as well as matrix effects (ME). Finally, the validated method was successfully applied to real samples.

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Phthalic Esters released in Hot Beverages by Vending Machines: Analytical Protocol by Means of Ultrasound-Vortex-Assisted Liquid-Liquid Micro-Extraction and GC-IT/MS

I. Notardonato, S. Passarella, C. Di Fiore, A. Iannone, G. Ianiri, M.V. Russo, P. Avino

Department of Agriculture, Environmental and Food Sciences, University of Molise, via De Sanctis, I-86100 Campobasso, Italy

E-mail: c.difiore@studenti.unimol.it

Food distributed by vending machines is considered the main source of food available in public indoor environments such as universities, schools, and offices. In order to achieve efficient service, it is necessary that food and drinks are delivered by means of containers, more or less resistant. Plastic containers are versatile due to their polymer molecular structure, but at the same time, they can cause possible risks to the health of the consumer.

Plastic containers present several functional properties such as protection against microorganisms and exogenous insects, the prevention of loss of flavour and moisture, but at the same time it is possible that monomers, oligomers and plasticizer additives can migrate into the food and contaminate it.

Phthalic acid esters (PAEs), or better known as phthalates, are added to the polymer molecular structure with the aim of increasing its flexibility, plasticity and strength. However, these chemicals can leach, migrating from the polymeric structure to the food, resulting in chemical contamination and a risk to the health of the consumer.

In this study, a simple, fast, inexpensive, and effective analytical method is developed for the research and determination of seven phthalates (DMP, DEP, DiBP, DBP, DEHP, DOP, DDP) [1] potentially released in hot drinks from disposable containers used in vending machines.

The extraction and preconcentration technique used is the ultrasound vortex assisted dispersive liquid-liquid microextraction (UVA-DLLME) [2]. The effect of the waiting time from the dispensing of the hot drink to its actual consumption for the extraction of the molecules is also investigated, obtaining a temporal release profile slightly shifted towards higher molecular weight PAEs and vice versa for those with low molecular weight. The optimization of the proposed analytical methodology is carried out by means of GC-FID whereas gas chromatography with ion trap mass spectrometry (GC-IT/MS) has been used both for confirmation and for analysis of the real samples. By GC-IT/MS obtaining values of LODs and LOQs below 3 ng mL^{-1} and 9 ng mL^{-1}, respectively, which are lower than the FID for ultra-trace analysis purposes, were obtained as well as recoveries better than 92 % with RSD below 11 %. Using this protocol different hot beverages such as coffee, long coffee, decaffeinated coffee, espresso, ginseng coffee, tea, and chocolate, were analyzed regarding the PAE content and the effect of the temperature on the compound release.

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Detection and quantification of white wine adulteration with cider and apple juice using voltammetric fingreprinting on a screen-printed electrode and chemometrics

M. Liveri[1], A. Tsantili- Kakoulidou[2], F. Tsopelas[1]

[1] Laboratory of Inorganic and Analytical Chemistry, School of Chemical Engineering, National Technical University of Athens, Iroon Polytechniou 9, 15780 Athens.
[2] Laboratory of Pharmaceutical Chemistry, School of Pharmacy, University of Athens, Panepistimiopolis, Zografou, 157 71 Athens, Greece.

E-mail: ftsop@central.ntua.gr

Food fraud, defined as any deliberate action of businesses or individuals to deceive customers about the quality and/or content of the food, is a major problem in food market due to the loss of confidence in customers, damage to brands' reputation and possible health risks caused by the presence of unidentified allergens or hazardous materials. Wine as an alcoholic beverage always scores in the top ten list of the most often adulterated commodities [1] and its authentication constitutes a challenge in analytical chemistry. Wine authentication can be performed usually by the quantification of their components by means of chromatographic techniques coupled mainly with mass spectrometry as detector. NMR spectroscopy has been also proposed [2]. Electroanalytical techniques can contribute to wine authentication, providing a non- specific fingerprint of wines and their common adulterants in respect to redox properties of species present in the samples [3]. Voltammetric techniques possess substantial advantages, involving the low cost of instrumentation and analysis, low detection limits, high sensitivity, inherent simplicity and ability to be utilized in mobile analytical devices. However, to the best of our knowledge, voltammetric techniques have not been applied for white wine authentication.

In the present study, an analytical approach for the detection and quantification of adulteration of white wine with apple juice and cider was developed. For this purpose, Greek wines, produced from different grape varieties and geographical origins, juices and ciders were collected by local markets. The optimization in voltammetric conditions showed that the most suitable approach was the direct analysis of the samples with cyclic voltammetry on a multi-walled carbon nanotubes- gold nanoparticles modified screen-printed electrode (DRP110CNT-GNP) after addition of a small amount of electrolyte. Exported voltammetric data were subjected to chemometric analysis using Principal Component Analysis (PCA) and Class modeling analysis by means of soft independent modeling of class analogy (SIMCA). The results showed that SIMCA offers a satisfactory classification of wines and apple juices/ cider samples. Finally, the quantification of adulteration in wine was carried out by voltammetric analysis of mixtures of white wines, apple juices and ciders in different proportions and submission of the voltammetric data to Partial Least Squares (PLS) Analysis. LOD and LOQ were calculated to be 3.5 and 7.5% (v/v), respectively. An appropriate test set were employed for the external validation of the model.

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Fast GC-MS method, for the quantification of polystyrene/polyurethane oligomers in liquid food simulants

E.D. Tsochatzis[1,2], H. Gika[2,3,4], G. Theodoridis[2,3,4]

[1]Department of Food Science, Aarhus University, Agro Food Park 48. 8200 Aarhus N. Denmark
[2]FoodOmicsGR Research Infrastructure, AUTh node, , GR 57001 Thessaloniki, Greece.
[3]Biomic AUTh, Center for Interdisciplinary Research and Innovation, GR 57001, Thessaloniki, Greece.
[4] Aristotle University of Thessaloniki, Thessaloniki, Greece.

E-mail: Emmanouil.Tsochatzis@food.au.dk

The use of plastic packaging is on significant growth, which is correlated with an increasing concern about the potential harm caused to the environment and human health. Oligomers are considered as Non-intentionally Added Substances" (NIAS) and can potentially migrate from the polymeric material to the food or food simulant. Styrene, the constituting monomer of PS, is an aromatic hydrocarbon that was recently reported to have a carcinogenic potential as well as its oxidised from (styrene oxide). The International agency for Research on Cancer (IARC) classifies styrene as a group 2A carcinogen, that is, possibly carcinogenic to humans, as well as recently EFSA [1]. To the best of the author's knowledge, this is the first multi-analytical method for simultaneous quantification of all the aforementioned analytes. The method has been optimized to quantify the monomers styrene and α-methyl styrene, selected polystyrene oligomers (dimers, trimers) and polyester urethane-based oligomers (PU) cyclic oligomers, as well as cyclic NIAS originating from food packaging such as 2,6-di-tert-butylbenzoquinone and 7,9-Di-tert-butyl-1-oxaspiro(4,5)deca-6,9-26 diene-2,8-dione (Figure 1).
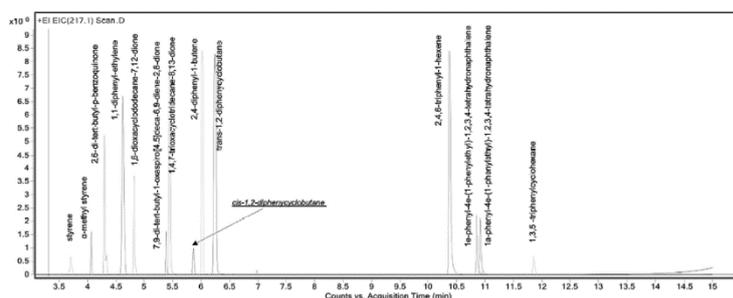
Figure 1. GC-MS chromatogram of spiked food simulant A (10 % v/v) ethanol in water

A liquid-liquid extraction (LLE) with dichloromethane, followed by gas chromatography coupled to mass spectrometry (GC-MS) was applied. Limits of detections ranging from 0.32 ng mL-1 (1,1-diphenyl-ethylene) to 14.8 ng mL-1 for 7,9-di-tert-butyl-1-oxaspiro[4.5]deca-6,9-diene-2,8-dione. Accuracy and precision results showed acceptable recoveries (80–110 %) and RSDs (<16 %), respectively at the tested concentration levels. The method was applied to seven FCM, indicating significant amounts of styrene and oligomers (dimers and trimers) and polyester urethane-based oligomers (PU) migrating into food simulant. Exposure assessment estimation was performed using EFSA's approach (Figure 2).
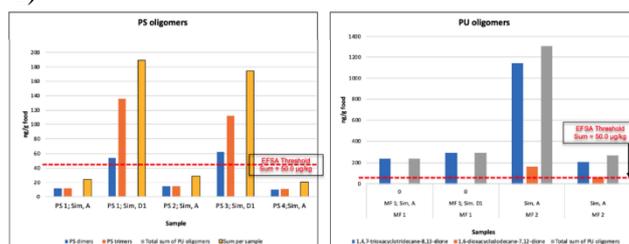

Figure 2. *In silico* exposure assessment for PS and PU oligomers resulting from testedFCM

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Wednesday 22 September 2021

POSTER SESSION 3

Materials | Sensors | Electrochemistry | Archeometry | Thermal analysis | Chemometrics

Chair: S. Girousi, K. Hatzigiannakis

IMA-2021

Novel ink-jet inks with UV-absorbers as active agents for polyester and polyamide digital printing

E. Karanikas, N. Nikolaidis

Department of Industrial Chemistry and Chemical Technology, School of Chemistry, Aristotle University of Thessaloniki, GR54124, Thessaloniki, Greece

E-mail: vagkaran@gmail.com

Materials which can provide effective protection against damage from prolonged exposure are called advanced materials for UV protection. The fabric cover factor directly decides the protection against UV radiation but indirectly depends on the type of weave, depth of shade, fabric areal density, stretchability, wetness and washing cycle of the fabric. Dianitide derivatives and various other chemicals are used to improve the UV protection factor (UPF) of textiles [1]. Two UV-absorbers, Tinuvin 477-DW and Tinuvin 5333-DW were used as active agents in this work. These were added in two different amounts (1.0% and 2.0%) in water-based disperse ink-jet inks. The surface tension, pH, viscosity and conductivity of the prepared formulations were monitored over a period of 90 days in order to estimate the ink stability and the compatibility of the UV-absorbers with the ink constituents. The inks were used for ink-jet printing on paper, transfer printing on polyester and polyamide and thermostabilization. Wash light and rub fastness were evaluated on the printed samples. Color coordinates were measured. Qualitative determination of the active agents, adsorbed on the printed samples was made by using UV-Vis Spectroscopy.

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Synthesis of Crosslinked and Short Chain Branched Polyethylenes nanocomposites reinforced with graphene nanoplatelets

D. Kourtidou[1], E. Tarani[1], D.N. Bikiaris[2], G. Vourlias[1], K. Chrissafis[1]

[1] Laboratory of Advanced Materials & Devices, School of Physics, Aristotle University of Thessaloniki, GR-54 124, Thessaloniki, Greece

[2] Laboratory of Chemistry and Technology of Polymers & Dyes, Department of Chemistry, Aristotle University of Thessaloniki, GR-541 24 Thessaloniki, Greece

E-mail: etarani@physics.auth.gr

Geothermal heat pumps are widely used as an eco-friendly and cost-effective house heating/cooling system. Additionally, there has been a growing interest toward nanocomposites applied in thermally conductive polymeric materials [1]. For this reason, the crosslinked high-density polyethylene (PEX) and the short-chain branched medium-density polyethylene (PE-RT) have been used as matrices for the synthesis of graphene nanoplatelets (GNPs) nanocomposites at various GNPs content by using the melt-mixing technique. The structural characterization of the nanocomposites was determined by X-Ray Diffraction (XRD). Thermogravimetric Analysis (TGA) was used to study the effect of the GNPs on the thermal stability of the prepared materials, respectively. Laser Flash Analysis (LFA) is used to estimate the thermal conductivity of both PEX and PE-RT/GNPs nanocomposites while tensile tests were conducted to evaluate their mechanical properties. LFA and tesnile tests results reveal that the PEX/GNPs nanocomposites increase more the thermal conductivity and elastic modulus values compared to those of PE-RT nanocomposites.
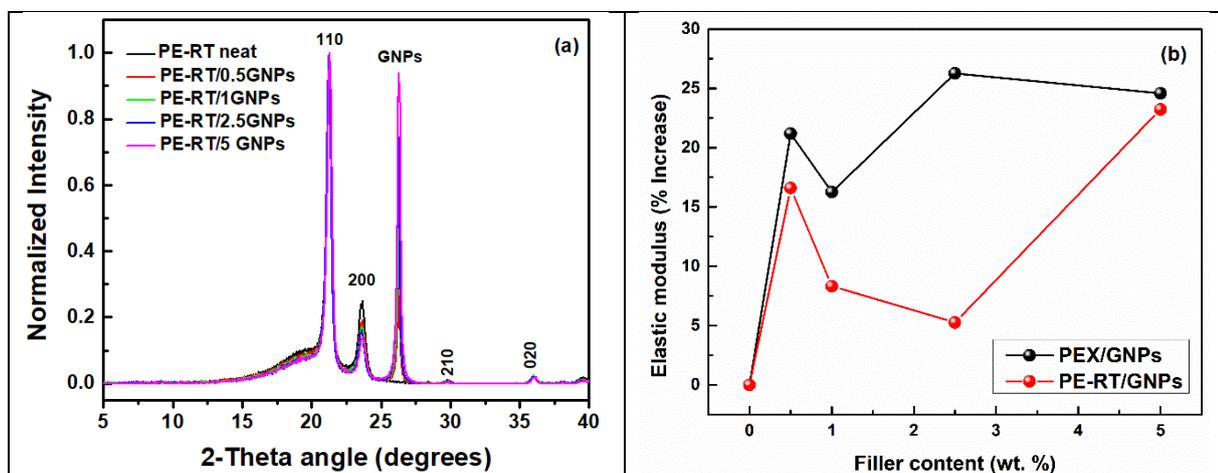
Figure 1. (a) XRD patterns of PE-RT/GNPs and (b) thermal conductivity measurements of PEX/GNPs and PE-RT/GNPs nanocomposites

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Acknowledgments

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Investigations of platinum dissolution by inductively coupled plasma mass spectrometry (ICP-MS) and transmission electron microscopy (TEM)

M. Pawlyta[1], A. Bacha-Grzechnik[2], S. Smykała[1], A. Jakóbik-Kolon[2]

[1]Silesian University of Technology, Materials Research Laboratory, Konarskiego 18A, 44-100 Gliwice, Poland
[2]Faculty of Chemistry, Silesian University of Technology, Strzody 9, 44-100 Gliwice, Poland

E-mail: miroslawa.pawlyta@polsl.pl

Polymer electrolyte fuel cells (PEFC) are a key technology for the global transition to clean and sustainable energy [1]. The high price and limited resources of platinum are currently the biggest barrier to the commercialization of fuel cells, therefore efforts are made to increase the durability of fuel cells. Durability is limited by Pt dissolution and carbon support corrosion, which additionally can initiate secondary degradation processes such as particle migration, agglomeration, Ostwald ripening, and particle detachment. The result is decrease in the electrochemically active surface area (ECSA), and further power losses in the PEMFC.

In our research, we use Transmission Electron Microscopy (TEM) to answer the question: what processes and to what extent determine the decrease in the electrochemically active surface area (ECSA) and consequently power losses in the PEMFC? For this purpose, we compare the TEM images obtained for fresh material and material after application test. Such comparisons were made for catalysts with different particle sizes and supported on carbon with varying degrees of crystallinity. The influence of the cell operating time and the range of potential change were also analyzed. Atomic scale insights on degradation process were obtained by using Identical Location TEM - microscopic techniques developed to track morphological changes of the catalysts material at the nanoscale level [2]. The obtained results were correlated with inductively coupled plasma mass spectrometry (ICP-MS), Ramman spectroscopy and Electron Energy Loss Spectroscopy.

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Preparation and characterization of biocompatible poly (lactic acid) / poly (ethylene adipate) blends

M. Lazaridou, C. Samiotaki, D. N. Bikiaris

Laboratory of Chemistry and Technology of Polymers and Dyes, Chemistry Department, Aristotle University of Thessaloniki, 54124, Thessaloniki, Greece

E-mail: christinasamiotaki@gmail.com, dbic@chem.auth.gr

Poly (lactic acid), PLA, is a highly biocompatible polyester that has concentrated a lot of interest in the last few decades. However, despite its unique properties and characteristics, this polyester requires a long time for its biodegradion to be completed and for that it tends to be combined with other biodegradable polymers. This can happen through copolymerization or with the synthesis of polymeric blends, as is this particular case. The purpose of this study is the synthesis of poly (ethylene adipate), PEAd, another aliphatic linear biodegradable polyester, and the deconstruction of the impact this polymer has on the characteristic properties of poly (lactic acid) (such as the biodegradion rate). The synthesis of poly (ethylene adipate) was performed by a two-step melt polycondensation process (esterification and polycondensation) with tributyltin as catalyst [1]. The polymeric blends were prepared by the solution casting method in five different ratios: PLA/PEAd: 90/10, 80/20, 70/30, 60/40 and 50/50. Both poly (ethylene adipate) and the mixtures were characterized by Fourier-transform infrared spectroscopy (FTIR), X-ray diffraction (XRD) and differential scanning calorimetry (DSC). In addition, viscosity, tensile strength and enzymatic hydrolysis of the blends were studied.

FTIR spectroscopy confirmed the successful synthesis of poly (ethylene adipate), while FTIR of the mixtures revealed interactions between the reactants (Fig. 1). Concerning the mixtures, DSC analysis showed reduced melting point temperature and glass transition compared to neat PLA. Viscosity study revealed a notable increase in the molecular weight and intrinsic viscosity of the polymeric blends as PLA content increases. From the tensile measurements it became clear that the absolute tensile strength increases with the increase of the PLA content of the mixtures, but we must emphasize that the tensile strength reaches up to a point and then decreases, i.e. it acquires a maximum. Finally, enzymatic hydrolysis revealed that as the content of the mixtures in PEAd increases, the films present a greater hydrolysis rate. In order to gain a better insight into the enzymatic hydrolysis process, morphological examination of the samples was performed by scanning electron microscope (SEM) (Fig. 2).
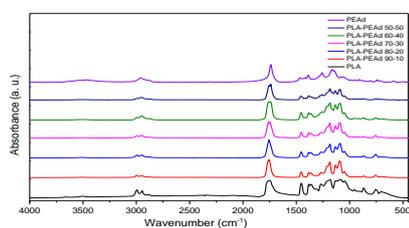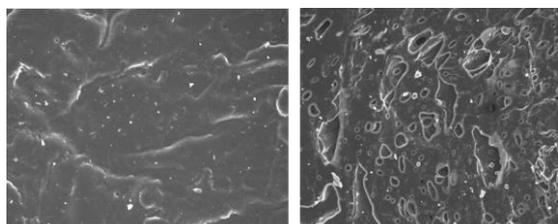

Figure 1. FTIR spectra of PLA-PEAd blends

Figure 2. PLA-PEAd 60-40 day 0 (left) and day 18 (right)

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Development of an electrochemical sensor for the detection of mephedrone on a modified carbon paste electrode with polymerized silver nanoparticles capped with saffron

G.C. Papaioannou, S. Karastogianni, S. Girousi

Aristotle University of Thessaloniki, Analytical Chemistry Laboratory, Chemistry Department, 54124 Thessaloniki, Greece

Email: girousi@chem.auth.gr,skarastogianni@hotmail.com, Geopapai.8@gmail.com

Electrochemical deposition methods provide a simple, repeatable, and green method for the formation of thin and uniform polymer film on the electrode surface [1]. Mephedrone or 4-methylmethcathinone is becoming the most popular new psychoactive substance among youth as a recreational drug. [2]. Mephedrone (1-(4-methylphenyl)-2-methylaminopropane-1-one) is one of the most potent derivatives of the cathinone. Mephedrone increase the level of lipid peroxidation and expression of antioxidant enzymes [3]. Regarding mephedrone's detection, only few attempts were made using electrochemical sensors, such as the imprinted electrochemical sensor, based on sol-gel molecular imprinted polymer polytyramine and gold nanoparticles (f-MWCNT@AuNPs) [4]. Therefore, this study describes the development of a novel, sensitive, selective, environmentally friendly and cost-effective electrochemical sensor for the detection of mephedrone. Silver nanoparticles (AgNPs) were capped with saffron (AgNPs@Sa) using a green method [5] and studied with ultraviolet-visible spectroscopy (UV-Vis), as well as with infrared spectroscopy (IR). The morphology of AgNPs@Sa was studied with electron scanning microscopy (SEM). The cyclic voltammetric deposition of poly-AgNPs@Sa on a carbon paste electrode (CPE) was performed by cycling the potential for 5 scans in phosphate buffer pH 7.0 containing silver nanoparticles. The electrode was immersed in a solution that contains the mephedrone. The sensor was evaluated under optimum condition to evaluate its analytical features such as linearity, selectivity, stability and reproducibility. Moreover, it was applied for the analysis of mephedrone in real samples. To the best of our knowledge, it is the first detection effort to electrochemically analyze mephedrone with AgNPs as electrode surface modifiers.

Keywords: electrochemically produced polymers, electrochemical sensors, saffron, silver nanoparticles, voltammetry.

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Development of a chemically modified sensor for the voltammetric determination of carbofuran

J. Barek[1,2], A.N. Solomonenko[1], E.V. Dorozhko[1], E.I. Korotkova[1]

[1] National Research Tomsk Polytechnic University, Tomsk, Russia
[2] Charles University, Prague, Czech Republic

E-mail: ans51@tpu.ru

Carbofuran (CBF) is toxic carbamate pesticide used for seed and plant protection against soil pests. Since CBF inhibits enzymatic activity and can lead to reproductive, cytotoxic, and genotoxic disorders, it is important to develop a method for CBF determination for monitoring environmental and food safety. The development of electrochemical sensors based on the chemical modification of the electrode surface for the CBF determination in recent years has aroused the active interest of scientists [1].

In this work, the CBF determination in model solutions was developed using a carbon-containing electrode (CCE) modified with carbon ink (CI) with the addition of a chromatographic sorbent (CS) based on chromaton (Ch), polyethylene glycol (PEG) and iron acetylacetonate (Fe(AA)$_3$). The surface morphology of the Ch/PEG/Fe(AA)$_3$/CI/CCE was investigated by scanning electron microscopy (SEM) in backscattered-electron (BSE) mode.

The electrochemical behavior of CBF at Ch/PEG/Fe(AA)$_3$/CI/CCE was studied by cyclic voltammetry (CV) in the potential range from -1 to 1.5 V and by electrochemical impedance spectroscopy (EIS) in a phosphate buffer solution (PBS) pH 6.86 (fig. 1).
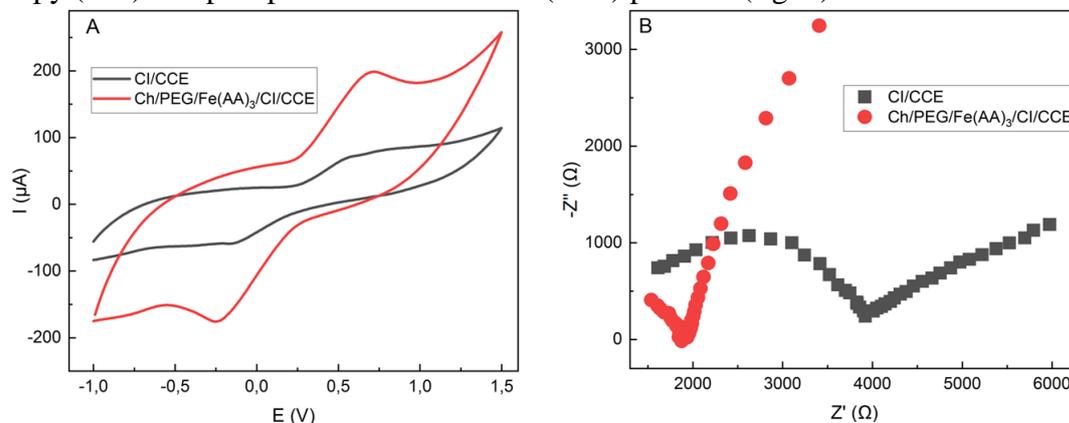

Figure 1. A) CVs of 0.1 M [Fe(CN)$_6$]$^{3-/4-}$ in 0.1 M KCl at CI/CCE (grey line) and Ch/PEG/Fe(AA)$_3$/CI/CCE (red line); B) EIS of 0.1 M [Fe(CN)$_6$]$^{3-/4-}$ in 0.1 M KCl at CI/CCE (grey) and Ch/PEG/Fe(AA)$_3$/CI/CCE (red).

The CV and EIS diagrams demonstrate the successful preparation of the Ch/PEG/Fe(AA)$_3$/CI/CCE. Under optimal experimental conditions for CBF model solutions, the detection limit (*LOD*) was 0.08 µM with a linear concentration dependence from 0.25 to 10 µM. The proposed Ch/PEG/Fe(AA)$_3$/CI/CCE demonstrated good sensitivity for CBF detection. The use of sorbents as modifying agents of electrodes for the determination of carbamate pesticides opens up the possibility of waste-free use of column sorbents in the framework of material disposal.

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IMA-2021

Electrochemical sensing of maple syrup urine disease (MSUD) biomarkers in a modified carbon paste electrode saffron coupled with silver nanoparticles

S. Karastogianni, S. Girousi

Analytical Chemistry Laboratory, Chemistry Department, Aristotle University of Thessaloniki, Thessaloniki, Greece.

E-mail: skarastogianni@hotmail.com

Branched-chain amino acids (b AAs) (leucine (Leu), isoleucine (Ile), and valine (Val)) partake in proteinic synthesis and protein catabolism [1]. A rare disease due to amino acid disorders is maple syrup urine smelling disease (MSUD). It is provoked by tremendous amounts of the L-branched-chain amino acid (b AAs), like leucine, isoleucine and valine (Leu, ILe and Val, biomarkers) [2]. In malady, these excessive elevations of b AAs provoke complications liver illness, neurological impairments and indeed death [2]. To avert these abrasions, it is exceptionally noteworthy to receive a timely and rapid diagnosis in the prime periods of life overdue to the inevitable consequence on nurslings [2, 3]. As it can be seen by the lack of literature [4-6], electrochemical sensors have been rarely used in the detection of b AAs related to MSUD clinical diagnostics. The present study describes the green synthesis of silver nanoparticles (AgNPs) obtained using saffron [7]. The proposed AgNPs coupled saffron were electropolymerized using cyclic voltammetry (CV) on a carbon paste electrode. The electrochemical behaviour of the modified electrode was performed by CV. The morphology of the electrochemical sensor was characterized by electron scanning microscopy (SEM). The constructed electrochemical sensor was used in the detection of selected biomarkers of MSUD, determining its analytical features such as linearity, selectivity, sensitivity and stability.

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Impedimetric sensor for hydroxyl radical detection in biological systems

J. Barek[1,2], A.V. Geraskevich[1], E.I. Korotkova[1]

[1]National Research Tomsk Polytechnic University, Tomsk, Russia

[2] Charles University, Faculty of Science, Department of Analytical Chemistry, UNESCO Laboratory of Environmental Electrochemistry, Albertov 6, CZ 128 43 Prague 2, Czech Republic

E-mail: avg48@tpu.ru

The hydroxyl radical (OH•) is known to be the marker of oxidative stress associated with different pathologies including neurodegenerative and cardiovascular diseases, as well as cancer [1]. There are various approaches for OH• determination in biological objects: ESR-spectroscopy [2], chromatography [3] and fluorimetry [4]. However, these techniques are rather complex and expensive. Therefore, electrochemical methods seem to have certain advantages due to their high sensitivity, short time of analysis, and simple equipment.

In these study, we have developed a new impedimetric sensors for OH• detection in living cells. The sensors are based on impregnated graphite electrode (IGE) and glassy carbon electrode (GCE) sequentially modified with gold film and 1-hexanethiol. Electrochemical impedance spectra (EIS) of bare electrodes and their modifications are presented in Figure 1.
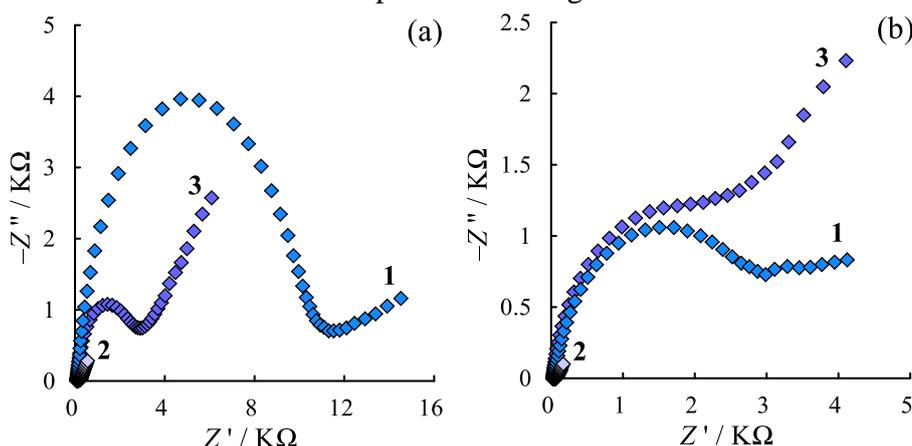

Figure 1. EIS of impregnated graphite electrode (a) and glassy carbon electrode (b); 1 – bare electrode, 2 – gold film surface, 3 – 1-hexanethiol modified gold surface

Obtained spectra fit the Randles scheme. As Figure 1 shows, bare IGE and GCE have considerable current transfer resistance (R_{ct}): 10 000 Ω and 2 744 Ω, respectively. However, gold film coating decreases R_{ct} to 45.5 Ω (IGE) and 18 Ω (GCE). Remarkable R_{ct} increase (IGE: 2 381 Ω; GCE: 2 047 Ω) after modification with 1-hexanethiol indicates the alkanethiol self-assembled monolayer (SAM) formation on sensors.

Developed sensors exhibit good analytical response to OH• (10^{-7} M) attack. R_{ct} decreased by approximately 100 Ω after conducting Fenton reaction for 15 minutes. Control experiments showed that SAM is not attacked by both hydrogen peroxide and iron (II) ions. Obtained data allow making C(OH•)–R_{ct} calibration curve and its statistical evaluation.

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Fluorescent carbon nanodots from *Dunaliella salina* biomass and their application as a 4-nitrophenol nanoprobe

T. Skolariki, T. G. Chatzimitakos, C. D. Stalikas

Laboratory of Analytical Chemistry, Department of Chemistry, University of Ioannina, 45110 Ioannina, Greece

E-mail: cstalika@uoi.gr

Dunaliella salina, an algal biomass often found in natural lakes or marine waters, is known for producing carotenoids and glycerin [1]. In the present work, a solvothermal chemistry approach was applied for the synthesis of two different kinds of fluorescent carbon nanodots (CNDs), utilizing a natural precursor i.e., biomass of halophilic microalgae *Dunaliella salina*. The strain of *Dunaliella salina* used was isolated from the Greek salt works. The synthesis produced two kinds of CNDs and after extraction with toluene the hydrophilic CNDs in the water-soluble fraction were characterized and further exploited. The fluorescent nanomaterial showed appreciable optical properties, quantum yield (9%) and a blue fluorescence emitted at 420 nm after excitation at 330 nm. Moreover, the CNDs also showed good photostability and their fluorescence intensity exhibited dependence on the pH. The CNDs proved to be a selective nanoprobe for the detection of 4-nitrophenol (4-NP), a man-made chemical widely used in the production of pharmaceuticals and pesticides, which has been listed as a priority pollutant due to the extent and severity of the environmental pollution it can cause. It has also been reported as human urinary metabolite which can be used as a marker to evaluate pesticides exposure [2]. The instant assay of 4-NP was achieved by quenching the fluorescence of CNDs via an inner filter effect mechanism [3] accompanied by a marked wavelength shift while the fluorescence intensity decreased with increasing the concentration. The developed method was applied to a variety of matrixes such as tap water, treated wastewater and human urine. The method showed a linear range from 0.8 to 53 μM with regression coefficient $R^2 = 0.99$, and acceptable sample recoveries (from 98 to 107%) in tap water and wastewater samples.

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3D Printed Voltammetric Sensor Modified with an Fe(III) Cluster for Nonenzymatic Determination of Glucose in Sweat

H. Koukouviti[1], F. Vasileiou[1], A. K. Plessas[2], A. Economou[1], N. Thomaidis[1], G. S. Papaefstathiou[2], C. Kokkinos[1]

[1]Laboratory of Analytical Chemistry, Department of Chemistry, National and Kapodistrian University of Athens, 15771 Athens, Greece
[2]Laboratory of Inorganic Chemistry, Department of Chemistry, National and Kapodistrian University of Athens, 15771 Athens, Greece

E-mail: christok@chem.uoa.gr

The skin is the largest human organ offering excellent opportunities for noninvasive access, especially via perspiration. Among the different biomarkers measured in sweat, glucose is the most important. The core devices for its determination are the electrochemical enzymatic biosensors [1,2]. However, the use of an enzyme entails several drawbacks as the activity of glucose oxidase is critically dependent on the immobilization procedure and on temperature, pH, ionic strength and other factors[2]. Nonenzymatic glucose monitoring devices can be alternatively used, but this area remains still largely unexplored, since very few works have been published in the pertinent literature [2-4]. Here, we present a 3D printed electrode modified with iron(III) basic benzoate, $[Fe_3O(PhCO_2)_6(H_2O)_3] \cdot PhCO_2$, for the nonenzymatic determination of glucose in sweat by stripping voltammetry. The electrode is 3D printed from a carbon black loaded polylactic acid filament and the trinuclear Fe(III) complex is drop-casted on the surface of the electrode, followed by its coverage with a Nafion film. The iron (III) basic benzoate was selected as a modifier thanks to its minimum water solubility, offering stable sensors to sweat conditions. The electrocatalytic determination of glucose is based on the in-situ formation of Fe_2O_3 by the reduction and oxidation of Fe(III) during the preconcentration and stripping steps, respectively. The catalytic mechanism of the Fe_2O_3 to glucose oxidation can be explained by the reaction: $Fe_2O_3 + Glucose \rightarrow 2FeO + Gluconolactone + H_2O$. Under the optimum conditions (i.e. preconcentration for 240s at -1.4 V) the sensor enables noninvasive glucose detection in sweat in the physiologically relevant concentration range of 50−500 μmol L^{-1} without interference from common electroactive metabolites. The 3D printing procedure adopted in this work overcomes the fabrication disadvantages of manufacturing technologies used in existing sensors (i.e. screen printed and sputtering), while the use of catalytic sensing of glucose on an Fe(III) cluster, as opposed to enzymatic sensing, endows the sensor with an attractive electroanalytical performance for sweat glucose monitoring, including high sensitivity, selectivity, reproducibility and stability.

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Electrochemical paper-based devices fabricated via programmable pen-drawing and screen-printing

D. Soulis[1], V. Pagkali[1], C. Kokkinos[1], M. Prodromidis[2], A. Economou[1]*

[1] Department of Chemistry, National and Kapodistrian University of Athens, Athens, 157 71 Greece
[2] Department of Chemistry, University of Ioannina, Ioannnina 45 100, Greece

E-mail: aeconomo@chem.uoa.gr

The field of paper-based analytical devices (PADs) was experienced a rapid growth over the last decade. The key features of paper as a platform for the fabrication of PADs are: a) flexibility; b) low thickness and lightness; c) absorbency; d) high surface-to-volume ratio; e) hydrophilicity and capillary action; f) chemical and biological inertness; g) disposability and biodegradability, and; h) low cost and wide worldwide availability. As a result of recent advances in this field, a host of paper-based devices has been developed for application in various fields such as clinical diagnostics and POC testing, environmental monitoring and food quality control [1,2].

The patterning of the hydrophobic barriers and electrodes are the most critical step in the fabrication of electrochemical PADs and several approaches have been proposed in the literature [3]. Pen-on-paper (PoP) strategies involve the use of a suitable pen or pencil to deposit functional materials on paper substrates with the view to create hydrophobic patterns on paper [4]. Compared to the traditional methods for fabricating PADs, the advantages of PoP approaches include simplicity, low cost, wide availability of materials, scope for rapid prototyping and flexibility. On the other hand, screen-printing technology has facilitated enormously the development of low-cost and portable electrochemical paper-based devices [5].

The aim of this work was the development of disposable electrochemical PADs using a dual fabrication approach which combines programmable pen-drawing and screen-printing. For this purpose, hydrophobic barriers were delimited by x-y plotting with a hydrophobic commercial marker pen. Electrodes were formed by depositing carbon electrodes on the PADs *via* screen-printing. Applicability of the PADs was demonstrated for electrochemical detection of organic compounds by differential pulse voltammetry and of heavy metals by stripping voltammetry.

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Development of a flexible electrochemical biosensor for fish stress monitoring

H. Wu[1], Y. Ube[1], H. Ohnuki[2], H. Endo[1]

[1] Department of Ocean Sciences, Tokyo University of Marine Science and Technology,
4-5-7, Konan, Minato-ku, Tokyo 108-8477, Japan

[2] Department of Marine Electronics and Mechanical Engineering, Tokyo University of
Marine Science and Technology,
2-1-6, Etchujima, Koto-ku, Tokyo 135-8533, Japan

E-mail: endo@kaiyodai.ac.jp

Measuring the stress response in fish is essential toward maintaining and monitoring fish health, and preventing the outbreak of fish diseases in the aquaculture industry. To achieve this goal, some wearable biosensor systems that can be easily implanted into a living fish body have been developed to monitor real-time blood glucose levels as a stress indicator in swimming fish. Because these sensors are constructed of hard material, however, they have the potential to damage the fish body. As a result, it is difficult to achieve "actual" stress monitoring in free-swimming fish. In recent years, attention has been focused on the use of thinner, more flexible materials for the construction of smaller and lighter biosensors using advanced metal nano inks. Here we introduce a flexible biosensor for stress monitoring that is friendly for living test fish.

First, an inkjet printer was filled with conductive gold and silver nano inks, printed on the surface of a PET film, and moved to high-temperature sintering to prepare flexible gold and silver electrodes. Next, to confirm the functionality of both electrodes, the electrochemical characteristics of each electrode were evaluated. The silver electrode was treated with chloride to stabilize its performance and glucose oxidase (GOx) was immobilized on the surface of the gold electrode to prepare a flexible biosensor with the gold electrode as the working electrode and the silver electrode as the counter electrode. We next attempted to quantify glucose by analyzing the oxidation current generated by the reaction between GOx and glucose by amperometry both before and after bending. Furthermore, as an application of this sensor to actual samples, we attempted to quantify glucose in the plasma of test fish (*Oreochromis niloticus*).

A sufficient redox current was obtained for the gold electrode, and we confirmed that the sensor could be used as a good working electrode. The silver electrode exhibited stable polarization potential by chlorination treatment, and could be used as the counter electrode for the sensor. These results clarified that the newly constructed electrode could be applied as a flexible biosensor. Next, when the standard glucose solution was measured using this biosensor, an increase in the output current value correlated with an increase in the glucose concentration at 10 to 400 mg dL^{-1} (Fig. 1), providing good coverage for fish glucose measurements. The signal trend was similar when measuring the actual sample. The output current of the sensor was decreased, however, compared with the *in vitro* results, which may be due to some biologic components in the fish. To address this issue, we will try to improve the sensor by coating it with biocompatible membranes in our future work. Overcoming this problem will make it possible to develop sensors with different morphologies by taking advantage of the characteristics of the flexible sensor, and it is expected that such a sensor could be applied to various attachment sites on fish.
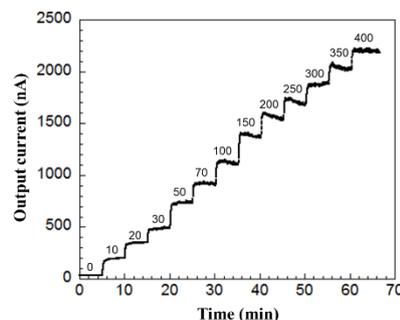

Fig. 1 The response curve of the proposed sensor in glucose solution with different concentrations.

Study of the influence of MWCNTs addition to the ion-selective membrane on the analytical parameters of electrodes sensitive to nitrates

K. Pietrzak, C. Wardak

Maria Curie-Sklodowska University, Faculty of Chemistry, Institute of Chemical Sciences, Department of Analytical Chemistry, Maria Curie-Sklodowska Sq. 3, 20-031 Lublin, Poland

E-mail: karolina.pietrzak@poczta.umcs.lublin.pl

Potentiometry is one of the simplest electrochemical techniques that allows to determine the concentration of inorganic and organic ions in liquid samples. The most popular potentiometric sensors are ion-selective electrodes, the potential of which depends on the activity of ions in the sample. In combination with a reference electrode (with a constant potential), they form a cell whose measurements of the electromotive force enables to determine the concentration of ions in the solution [1].

In order to improve the analytical parameters of sensors, new chemicals are constantly searched for, which act as an active substance in the membrane, enabling the achievement of good selectivity of the sensors. Various materials are also developed that can provide good stability and reversibility of the potential (ion-electron transducers) in ion-selective electrodes with solid contact which do not contain an internal solution. Conductive polymers were the first substances used as solid contact, but nowadays, due to the great popularity of all nano-sized materials, nanomaterials and nanoparticles are widely used for this purpose. Nanomaterials are characterized by a high surface to volume ratio, very good electrocatalytic activity as well as high electrical conductivity and mechanical strength [2]. Their use as an intermediate layer or a component of the ion-selective membrane improves the stability of the electrode capacitance and reduces the membrane resistance [3]. So far, carbon nanomaterials such as single-walled carbon nanotubes (SWCNTs) and multi-walled carbon nanotubes (MWCNTs), graphene, fullerenes, carbon nanofibers and carbon black have been used as a solid contact.

Initial studies of ion-selective electrodes for the determination of nitrate(V) ions were performed, for the construction of which various types of MWCNTs were used. The effect of modifying the composition of the ion-selective membrane of the obtained potentiometric sensors was tested by determining the basic analytical parameters, which include: slope of calibration curve, detection limit, measuring range, pH range, response time and potential stability. As a result of the research, ion-selective nitrate electrodes were obtained, which are simple both in construction and operation, and have good analytical parameters. The electrodes obtained as a result of the modification, regardless of the type of MWCNTs, show lower detection limits and wider measuring ranges compared to unmodified electrodes.

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Identification of degradation products and new colouring components in shellfish purple and indigo by UHPLC-MS/MS

A. Vasileiadou[1], I. Sampsonidis[1,2], S. Kalogiannis[2], G. Theodoridis[1], I. Karapanagiotis[3], A. Zotou[1]

[1]Laboratory of Analytical Chemistry, Department of Chemistry, Aristotle University of Thessaloniki, 54124 Thessaloniki, Greece
[2]Department of Nutritional Sciences and Dietetics, International Hellenic University (IHU), 57400 Thessaloniki, Greece
[3]University Ecclesiastical Academy of Thessaloniki, Department of Management and Conservation of Ecclesiastical Cultural Heritage Objects, 54250 Thessaloniki, Greece

E-mail: azotou@chem.auth.gr

Degradation products of indigo [1] and shellfish purple and four colourants of shellfish purple which are not often detected [2] were identified (i) in standard solutions of indigotin and 6-monobromo-indigotin, (ii) pigments and (iii) aged textile samples using a UHPLC-MS/MS method. Chromatographic separation in UHPLC was carried out on ACQUITY UPLC BEH C18 (1.7μm, 2.1 x 1.8 mm) column thermostated at 40°C. The mobile phase consisted of solvent A: water – 0.1% (*v/v*) formic acid and solvent B: ACN – 0.1% (*v/v*) formic acid using a gradient elution program during a time course of 38 min. The flow rate was 300μL min^{-1}. MS was performed using an electrospray ionization (ESI) Ion Max source with a HESI-II probe. The ESI was operated in positive mode (ESI+) for the detection of the colouring components and in negative mode (ESI-) for isatin (IS) and the degradation products of indigotin and 6-monobromoindigotin. In order to achieve adequate ionization in ESI for direct infusion analysis of the standard compounds, a novel solution preparation approach had to be developed since DMSO, the solvent into which the compounds are readily diluted in, is not fully compatible with ESI direct infusion analysis due to its high viscosity and high boiling point. The primary solutions of the colouring components, indigotin and 6-monobromoindigotin,were prepared by diluting with NaOH aqueous solutions for optimisation of detection parameters. To sum up, the four colourants were identified in all shellfish purple samples in small quantities. These components belong to the same structural class with the mono-, di- and non-brominated indigoids and have one unit lower masses than indigotin and its mono- and di-brominated derivatives. Two degradation products, isatin and isatoic anhydrite (DP3) were detected in indigotin and 6-monobromoindigotin standard solutions. Isatin and some other indigoid degradation products were identified in both shellfish purple and indigo samples, as well as, in the standard solutions of indigotin and 6-monobromoindigotin.

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Characterization of ancient burial pottery at Ban Muang Bua archaeological site (northeastern Thailand) using X-ray spectroscopy

C. Boonruang[1,2], K. Won-in[3*], N. Nimsuwan[3], P. Pakawanit[4], P. Dararutana[5]

[1] Department of Physics and Materials Science, Faculty of Science, Chiang Mai University, Chiang Mai 50200 Thailand
[2] Center of Excellence in Materials Science and Technology, Chiang Mai University, Chiang Mai 50200 Thailand
[3] Department of Earth Sciences, Faculty of Science, Kasetsart University, Bangkok 10900 Thailand
[4] Synchrotron Light Research Institute, Nakhon Ratchasima 30000 Thailand
[5] Retired Army Officer at the Royal Thai Army, Lopburi 15000 Thailand

E-mail: kritwonin@gmail.com

It was revealed that the ancient potteries found at Ban Muang Bua located in northeastern Thailand, were one of the Thung Kula Ronghai cultural groups [1]. Most of the productions were daily-used ceramics and burial goods. The characterization of potsherds was examined using techniques based on X-ray spectroscopy. Elemental composition and morphology were analyzed using a scanning electron microscope coupling with energy-dispersive X-ray fluorescent spectroscopy (SEM-EDS). Three-dimensional (3-D) structure was evaluated under X-ray tomographic microscopy based on synchrotron radiation (SR-XTM). The morphology contained variety in sizes irregularly plate-like granules of composite materials. The elements, such as C, Si, Al, P, S, Ca, K, Ti, Fe, Na, Mg, and Zn which present as major, minor, and trace elements were identified as the fingerprint origin. The 3-D tomographic images showed the internal configuration of these samples. It was found that the combination of various analytical methods was a powerful tool with which to answer the questions posed by archaeology.

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Development of a portable LED-IF instrument for the characterisation of biodeterioration on monuments

A. Giakoumaki[1], V. Piñon[1], K. Melessanaki[1], K. Hatzigiannakis[1], P. Pouli[1], D. Anglos[1,2]

[1] Foundation for Research and Technology, Institute of Electronic Structure and Laser, Nikolaou Plastira 100, 70013, Vassilika Vouton, Heraklion, Crete, Greece

[2] University of Crete, Department of Chemistry, Voutes University Campus, P.O. Box 2208, 70013, Heraklion, Crete, Greece

E-mail: agiakoumaki@iesl.forth.gr

Biological deterioration is one of the main factors of decline of the physical, chemical and mechanical properties of materials on a historic building as well as of the overall aesthetic appearance of the monument. Moreover, a monument might have many different types of biological deterioration even on adjacent areas depending on their orientation and place on the monument (that leads to different sunlight, temperature, way of rain attack, wind and humidity conditions among others), composition and microstructure of the materials etc. The different types of microorganisms/organisms that can colonise a historic building include cyanobacteria, fungi, algae, lichens, moss etc.[1]

A portable fluorometer, using an economical Light Emitting Diode (LED) source and a compact spectrometer, has been developed with the aim to characterize the biodeterioration of a monument. Since the conventional techniques used for the analysis of biological attack require sampling and are usually costly and time-consuming, a completely non-destructive method that can be used in-situ, as is LED-IF (LED-Induced Fluorescence), could be proven vital for the protection of a monument.

The potential of the system is examined on several artefacts composed of different substrates presenting diverse types of biodeterioration and a first classification is achieved. Various aspects are also discussed, such as the selection of the proper LED wavelength for this type of compounds, while a methodology proposed for its use in the lab and/or in-situ will be presented. The LED-IF system is developed within the project CALLOS: **Conservation of Athens antiquities with Laser and Lidar technologies Open to Science and public** (MIS- 5056208), an "**open innovation in culture**" action co-funded by the European Regional Development Fund (ERDF) and national resources through the Operational Program "Competitiveness, Entrepreneurship & Innovation" (NSRF -EPAnEK).

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Paper-based device for the determination chloride as a biomarker of cystic fibrosis based on the photochemical reduction of silver halide nanocrystals

C. Matiaki, A.G. Vlessidis, D.L. Giokas

Department of Chemistry, University of Ioannina, 45110, Ioannina, Greece

E-mail: pch01380@uoi.gr

A simple method for the determination of chloride anions in sweat as a biomarker of cystic fibrosis is described. The method relies on the different colorimetric response of silver ions and silver chloride crystals when exposed to UV light. Silver ions change from colourless to dark grey-brown due to the formation of nanosized Ag while AgCl exhibits a less intense colour change from white to slightly grey. The analytical signal is obtained as mean grey area intensity on the paper surface with a simple camera and is used to directly quantify the concentration of chloride anions in the sample. The method requires only three steps to perform (addition of sample, irradiation and measurement) and is performed with minimum sample volume (2 μL). The working range of the method covers the entire diagnostic range of chloride levels related to cystic fibrosis (10-140 mM) offers low detection limits (2.7 mM Cl$^-$) and satisfactory recoveries (95.2-108.7%) and reproducibility.
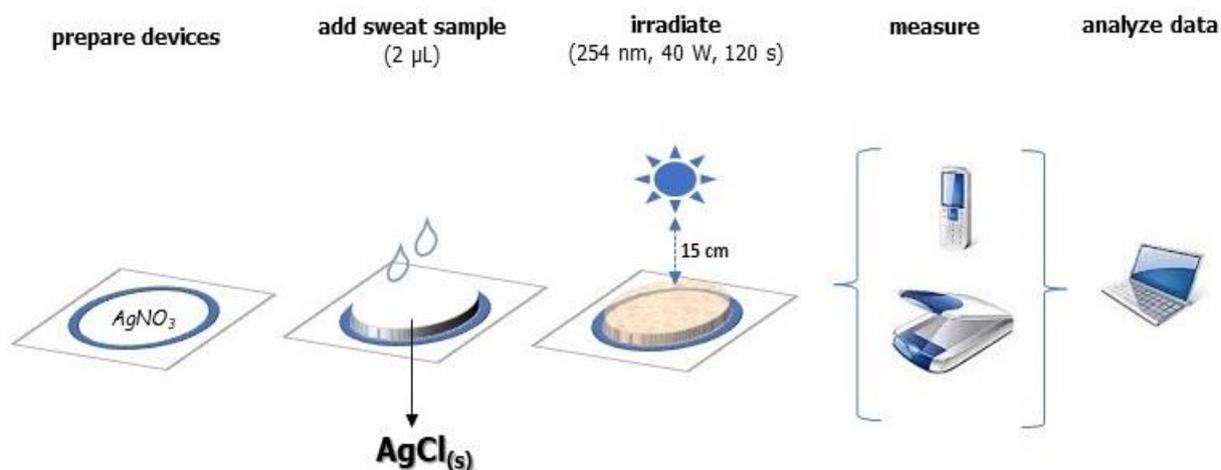
Voltammetric determination of 2,3-dichloro-1,4-naphthoquinone

J. Barek[1,2], N.V Aseeva[1], E.I. Korotkova[1]

[1] National Research Tomsk Polytechnic University, Tomsk, Russia
[2] Charles University, Faculty of Science, Department of Analytical Chemistry, UNESCO Laboratory of Environmental Electrochemistry, Albertov 6, CZ 128 43 Prague 2, Czech Republic

E-mail: natali.shkuratova@mail.ru

Naphthoquinones, as well as their derivatives, have a variety of pharmacological properties, namely, they have antimicrobial, anti-inflammatory, antiviral and antitumor effects [1]. 2,3-Dichloro-1,4-naphthoquinone (DCNQ) was chosen as a starting material for studying the electrochemical properties of naphthoquinones because it is sufficiently stable, readily available, and known as a key synthetic intermediate in organic, medicinal, and industrial chemistry. In this work, the electrochemical properties of DCNQ were investigated by cyclic voltammetry (CV) in the potential range from -1 to + 1.5 V on an impregnated graphite electrode (IMGE). A stock solution of 0.1 M DCNQ was prepared in dimethylformamide (DMF). Ethanolic solution of 0.1 M $NaClO_4$ acidified with 0.1 M HCl to pH 2.0 was used as a supporting electrolyte. IMGE was used as a working electrode, and Ag|AgCl (1M KCl) was used both as an auxiliary and a reference electrode. The electrodes were connected to TA-Lab voltammetric analyzer (Tomskanalyt, Tomsk, Russia). To determine the hydrogen index (pH), a pH meter/ionomer "ITAN" (Tomskanalyt, Tomsk, Russia), was used.

DCNQ gave one oxidation peak at 0.3 ± 0.05 V and a corresponding reduction peak at -0.1 ± 0.05 V. The calibration curve was obtained by cathodic voltammetry in the first derivative mode (**Figure 1**).
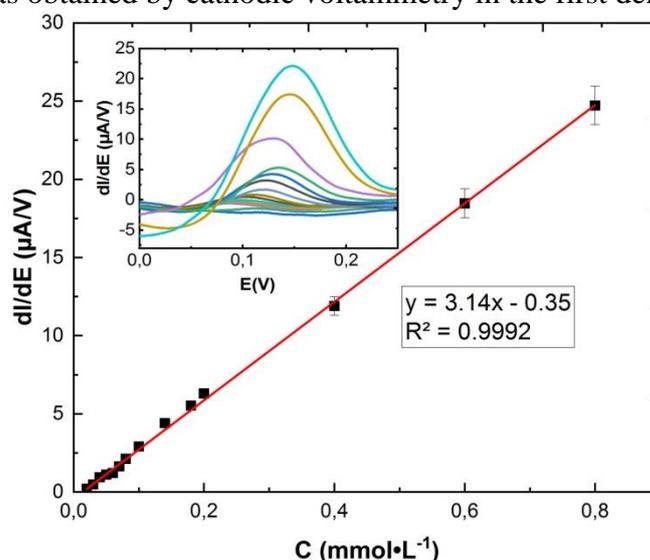

Figure 1. Dependence of the cathodic current on IMGE in the first derivative mode of 2,3-dichloro-1,4-naphthoquinone on its concentration in ethanolic solution of 0.1 M $NaClO_4$ (pH 2.0. potential scan rate 100 mV·s^{-1}.

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$CeO_2 \cdot Fe_2O_3$ nanoparticles modified electrode for the sensitive and selective quantification of lipoic acid

G. Ziyatdinova, L. Gimadutdinova

Analytical Chemistry Department, Kazan Federal University, Kremleyevskaya, 18, Kazan, 420008, Russian Federation

E-mail: Ziyatdinovag@mail.ru

Metal oxide nanoparticles are widely applied as electrode surface modifiers in modern organic electroanalysis. Among them inert metal oxides nanoparticles (TiO_2, In_2O_3, CeO_2, ZnO, Fe_3O_4 etc.) are of interest as far as provide high effective area of the electrodes, electrocatalytic activity, high selectivity and sensitivity of target analyte determination as shown on example of phenolic compounds [1, 2], neuromediators [3, 4], some pharmaceuticals [5, 6]. Combination of several metal oxide nanoparticles can provide improvement of analytical characteristics. Among the analytes, a group of S-containing compounds is less investigated. In particular, lipoic acid is almost out of consideration.

Novel modified electrode based on $CeO_2 \cdot Fe_2O_3$ nanoparticles has been developed for the lipoic acid quantification. Electrode surface modification provides 5.6-fold increase of the lipoic acid oxidation currents and 20 mV anodic shift of the oxidation potential. The best voltammetric parameters have been obtained for the 0.5 mg mL^{-1} dispersion of $CeO_2 \cdot Fe_2O_3$ nanoparticles. Scanning electron microscopy confirms the presence of spherical nanoparticles of 25-60 nm and their aggregates evenly distributed on the electrode surface and formed porous coverage. This leads to the 4.4-fold increase of the effective surface area *vs*. bare glassy carbon electrode. Electrode modification provides significant increase of the electron transfer rate as electrochemical impedance spectroscopy data indicate.

Electrooxidation of lipoic acid on $CeO_2 \cdot Fe_2O_3$ nanoparticles modified GCE is irreversible diffusion-controlled pH independent process occurring with participation of two electrons.

Direct differential pulse voltammetric method of lipoic acid determination in phosphate buffer pH 7.0 medium has been developed. The linear dynamic ranges of 0.075-7.5 and 7.5-100 μM with the detection limit of 0.053 μM have been obtained. These characteristics are the best ones reported for electrochemical methods including other modified electrodes. Another advantage of the method is high selectivity of the electrode response towards lipoic acid in the presence of inorganic ions, ascorbic acid, saccharides as well as other S-containing compounds (cysteine, cystine and methionine).

Approach developed has been applied on the pharmaceutical dosage forms of lipoic acid (tablets and concentrate for the infusion preparation). Voltammetric results have been validated with coulometric titration data.

Thus, simple, sensitive and selective, cost-effective and reliable approach has been developed for the lipoic acid quantification that can be applied for the pharmaceutical quality control.

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Adapting Surface Enhanced Raman Spectroscopy for organic residue analysis in archaeological artifacts

M.E. Konstantinou[1,2], A. Philippidis[1], D. Anglos[1]

[1]Institute of Electronic Structure and Laser, Foundation for Research and
Technology-Hellas (IESL-FORTH), 700 13 Heraklion, Crete, Greece
[2]Department of Materials Science and Technology, University of Crete, 700 13 Heraklion, Crete, Greece

E-mail: mek@iesl.forth.gr

Organic archaeological residues analysis is a high challenging field in archaeometry. Determining organic remains in archaeological objects provides information about human activities in certain period or site, including food preparation, storage or even rituals. Detection and chemical identification of organic biomarkers require the use of high-sensitivity techniques to overcome the complexity limitations of the samples, in terms of composition and low concentrations of constituent compounds. So far, the techniques that have been mainly reported to be used for this type of analysis are extraction methods followed by the use of chromatographic separation techniques (Gas Chromatography - GC, Liquid Chromatography - LC, High Performance Liquid Chromatography - HPLC) conjugated with mass spectrometry providing precise determination of the structure and molecular weight of compounds even at low concentrations. Another approach, in residues analysis, are spectroscopic methods which are able to overcome limitations of these methods arising from sample preparation requirements and measurement duration along with the difficulties in performing in situ analysis. In this study we focus on the development of Surface Enhanced Raman Spectroscopy (SERS), using Ag nanoparticles, for the identification and further study of syringic acid [1,2] (4-hydroxy-3,5-dimethoxybenzoic acid), a naturally occurring phenolic compound derived from edible plants and fruits and also a grape-related organic compound in archaeological remains. Syringic acid solutions in different concentrations mixed with the Ag nanoparticle solution were examined resulting a strong enhancement of Raman scattering. In order to simulate the detection of an organic compound in storage vessels we merged ceramic objects in syringic acid solution followed by an extraction method with polar solvents and the extraction solutions were examined with SERS technique.

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Laser-induced breakdown spectroscopy (LIBS) combined with machine learning models and neural networks for the classification of hard tissue remains

L. Spanos[1,2], P. Siozos[2], A. Koropoulis[3], N. Hausmann[5,6], M. Holst[6,7], P. Pavlidis[4], D. Anglos[1,2]

[1] Department of Chemistry, University of Crete, P.O. Box 2208, GR 710 03 Heraklion, Crete, Greece

[2] Institute of Electronic Structure and Laser, Foundation for Research and Technology-Hellas (IESL-FORTH), P.O. Box 1385, GR 711 10 Heraklion, Crete, Greece

[3] Institute of Molecular Biology and Biotechnology, Foundation for Research and Technology-Hellas (IMBB-FORTH), P.O. Box 1385, GR 711 10 Heraklion, Crete, Greece

[4] Institute of Computer Science, Foundation for Research and Technology-Hellas (ICS-FORTH), P.O. Box 1385, GR 700 13 Heraklion, Crete, Greece

[5] Niedersächsisches Institut für historische Küstenforschung, Viktoriastr, 26/28, 26382 Wilhelmshaven, Germany

[6] BioArCh, Department of Archaeology, University of York, Kings Manor, Exhibition Square, YO1 7ep York, UK

[7] York Osteoarchaeology Ltd, 75 Main Street, Bishop Wilton, York, YO42 1SR

Email: lspanos95@gmail.com

Archaeological and forensic excavations often reveal large numbers of hard tissue remains from multiple individuals. The discrimination among individuals and the classification of their remains is vital for revealing the identity, as well as information about the life and death of those individuals. There exist several well-established methods for the analysis of disarticulated or commingled skeletal assemblages. However, these methods are difficult to be applied in-situ, are time-consuming and have increased costs. Therefore, there is a strong interest in rapid and cost-effective methods for the analysis of hard tissue remains that can be applied in large numbers of skeletal findings and ideally on-site. [1, 2]

In the current work, several bone fragments and teeth were studied using a Laser Induced Breakdown Spectroscopy (LIBS) setup with nearly microscopic resolution. [2] The main objective of the investigation was to test various Machine Learning algorithms such as K-Nearest Neighbors, Random Forest and Support Vector Machine, as well as a simple Artificial Neural Network algorithm concerning their ability to facilitate discrimination and classification of hard tissue remains. Despite strong similarities among spectra the selection of specific spectral intervals with spectral lines corresponding to biogenic elements (such as Ca, Mg, Sr, Na and P) was found to be a determining factor that increased the resulting classification accuracy.

The use of the Artificial Neural Network algorithm produced significantly better results concerning classification compared to the Machine Learning models. The degree of classification achieved, varied from decent to excellent, providing an overall good sorting accuracy regardless of the type of tissue analyzed.

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Physicochemical and adsorption characteristics of thiomethacrylate-*co*-divinylbenzene microspheres

A. Bosacka, M. Wasilewska, A. Chrzanowska, D. Sternik, A. Derylo-Marczewska, B. Podskoscielna

Maria Curie-Skodowska University, Institute of Chemical Sciences, Maria Curie-Skoldowska Square 3, 20-031 Lublin, Poland

E-mail: alicja.bosacka@poczta.umcs.lublin.pl

Water pollution is a growing problem over the world due to the developing of industry and globalization. Organic compounds are particularly difficult to remove from water due to the difficulty of their decomposition [1-2]. The adsorption efficiency is influenced by many factors: adsorbent textural properties, adsorbent surface chemistry (surface functional groups) and also adsorbate properties (molecular weight, ionization strength (pKa), functional groups or water solubility (hydrophobicity)) [3-4]. New or improved adsorbents are widely sought, therefore, novel two-component materials consisting of thiomethacrylate and divinylbenzene phases are proposed.

In this work, thiomethacrylate-*co*-divinylbenzene microspheres were synthesized in a polymerization reaction at three different molar ratio phase compositions: 3:1, 2:1 and 1:1.

The obtained materials were analyzed using several analytical techniques. The textural properties of the samples were investigated using scanning electron microscopy (SEM) and low-temperature adsorption-desorption sorption. Fourier transform infrared-attenuated total reflection (FT-IR-ATR) spectroscopy was used to assess the binding efficiency of both phases. Thermal stability of thiomethacrylate-divinylbenzene materials was examined by thermal analysis (TG/DTG/DSC). Moreover, studies of adsorption kinetics were carried out in order to evaluate the adsorption efficiency of the obtained adsorbents towards selected aromatic adsorbates.

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Development of modern analytical methods employing innovative sample preparation techniques in combination with chemometric tools for the investigation of food authenticity

N. P. Kalogiouri, V. F. Samanidou

[1]Laboratory of Analytical Chemistry, Department of Chemistry, School of Sciences, Aristotle University of Thessaloniki, 54124, Thessaloniki, Greece

E-mail: kalogiourin@chem.auth.gr

Modern research in the field of food science and nutrition is moving rapidly towards the development of advanced analytical techniques to ensuring the safety, quality and traceability of food. As far as agricultural products are concerned, it is essential to establish analytical methodologies to guarantee the geographical origin and the type of farming (organic or conventional) in international trade. Nuts are a valuable source of nutrients with great economic importance for Greece. Among Greek agricultural products, almonds, walnuts, pistachios, hazelnuts and chestnuts have not been studied in depth yet [1]. The objective of this research was to develop analytical methodologies for the characterization of nut species and the determination of elements, rare earths, volatiles, fatty acids, phenols and tocopherols, employing novel sample preparation techniques, in accordance with the guidelines of Green Analytical Chemistry (GAC). For this reason, metals were determined in nuts using an Inductively Coupled Plasma – Atomic Emission Spectroscopy (ICP-OES) methodology [2]. Rare earths were determined by Inductively Coupled Plasma – Mass Spectrometry (ICP-MS) [3]. The volatile profile of nuts was investigated by Solid-Phase Microextraction (SPME) coupled with Gas Chromatography-Mass Spectrometry (GC-MS) [4]. The fatty acid profile was assessed by gas chromatography coupled with flame ionization detection (GC-FID). For the determination of tocopherols by HPLC-PDA, a fabric-phase sorptive extraction protocol was developed and validated. Finally, an HPLC-PDA methodology was developed for the simultaneous determination of phenolic acids and flavonoids. The quantification and semi-quantification results were analyzed using unsupervised and supervised chemometric tools to discriminate between Greek and imported nuts, and between nuts produced with conventional and organic type of farming.

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Thermal behavior and pyrolysis kinetics of olive stone residue

K.Chrissafis, T.Asimakidou

Advanced Materials and Devices Lab, School of Physics, Aristotle University of Thessaloniki, 541 24 Thessaloniki, Greece

E-mail: tasimaki@physics.auth.gr

In the last decades, one of the most ordinary challenges for countries of Southern Europe face is the management of wastes and by-products of olive oil production activities which are disposed into the environment and contribute to the enhancement of major ecological issues due to their organic loads. Exploitation of these olive residues wastes by thermochemical treatment can generate a wide range of valuable products. In this work, the thermal decomposition process of olive stone sample was performed under inert atmosphere at 5, 10, 15 and 20°C/min heating rates using thermogravimetric analysis (TGA), while the kinetic analysis performed via model free and model based methods. The changes in the crystal structure for various pyrolysis temperatures of the sample where examined by X-Ray Diffraction Analysis (XRD), while the morphological characteristics by Scanning Electron Microscopy (SEM). Results indicated that thermal behavior of olive stone sample was the typical one for lignocellulosic materials with the first mass loss being attributed to moisture removal while the following stages were assigned to lignocellulosic degradation substances. Ozawa-Flynn-Wall (OFW) and Vyazovkin (VYA) isoconversional methods were performed for Effective Activation Energy (E_a) and Pre-exponential factor estimation in model-free methods, while in model based method the best fitting results was obtained by a three independent parallel reactions model, obeying the n-th order with Fn code and described by the $f(\alpha)=(1-\alpha)^n$ equation.

Thermal and Spectroscopic Study of a Commercial Resin

C. Kyriakou Tziamtzi, L. Malletzidou, K. Chrissafis

Advanced Materials and Devices Laboratory, School of Physics, Aristotle University of Thessaloniki, GR-54124, Thessaloniki, Greece

E-mail: labrinim@auth.gr

Thermosetting resins have been rigorously studied due to their extensive utilization in numerous industrial applications, because of their low cost and high toughness, with their main advantage being their wide range of processing temperatures [1]. In this study, a commercial epoxy is studied by means of Thermogravimetry (TGA) and Fourier transform infrared spectroscopy (FTIR), for the evaluation of its thermal stability, curing process and structural characterization.

It is estimated that neat components of catalyst and epoxy begin to decompose at approximately 50 °C and 200 °C, respectively. Thermal degradation of epoxy/catalyst mixtures occurs in two steps under air atmosphere, and reveals lower thermal stability, compared to the decomposition of the resin under nitrogen, which is conducted in a single step. This is attributed to the higher cross-linking density accomplished during heating under inert atmosphere, while a higher mass residue is observed for the specimen heated under nitrogen, due to char formation and absence of oxygen promoted gasification of mixture fragments. Dynamic DSC ramp at a heating rate of 10 °C/min reveals the total heat of curing reaction, which is found equal to 406 J/g, while glass transition temperature (T_g) of the cross-linked structure is found equal to 66 °C. Furthermore, FTIR spectra of neat epoxy display a structure identical to that of DGEBA. Finally, the presence of amine groups is confirmed in the catalyst spectra, in addition to the effective polymerization of the epoxy/catalyst mixture, which is validated through the elimination of epoxy groups.
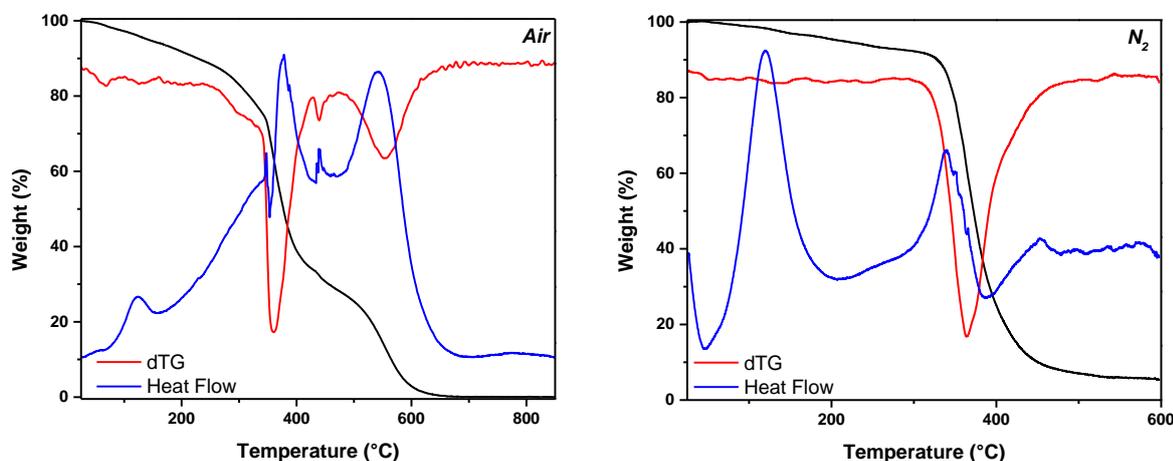

Figure 1: Thermogravimetric curves (weight loss, dTG and heat flow) of the epoxy/catalyst mixture, under air (left) and nitrogen (right) atmosphere.

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