

EuPA NEWSLETTER

2026
JANUARY

ISSUE 1

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MESSAGE FROM THE EDITORS

Dear Colleagues and Members of the European Proteomic Society (EuPA),

Welcome to the first issue of the EuPA Newsletter for 2026. As we begin a new year, we are pleased to share with you a collection of articles that reflect both the scientific excellence and the strong sense of community that define EuPA.

This issue opens with a heartfelt tribute to Amos Bairoch, whose visionary work laid essential foundations for modern proteomics and whose influence continues to resonate across our field. We also report on key outcomes of the EuPA General Council Meeting held in December 2025, highlighting new initiatives, upcoming elections, and preparations for EuPA 2026 in Würzburg.

Scientific inspiration continues with reflections on HUPO 2025, but we are also looking ahead as we announce the 3rd Annual Proteomics Gathering organized by YPIC, an event dedicated to empowering early-career researchers, as well as the opening of applications for the EuPA Awards 2026.

We hope this issue informs, inspires, and encourages you to engage with EuPA's activities throughout the year. As always, we warmly welcome your feedback and contributions to future newsletters.

Best regards,

On behalf of the EuPA Conference and Communication Committee
the Editor of the EuPA Newsletter: Eleni Zografos, and
the Co-Editors of the EuPA Newsletter: Éva Csősz and Fernando Corrales

IN MEMORIAM: AMOS BAIROCH, THE ARCHITECT OF SWISS-PROT

Past November 29, **Amos Bairoch** passed away at the age of 68. He was one of the most outstanding scientists in laying the foundations of our knowledge of proteins, always with the commitment to make this knowledge accessible to the entire scientific community. As leader of the **CALIPHO** group at the *Swiss Institute of Bioinformatics* of the University of Geneva, Amos specialized in protein sequence analysis and their meticulous annotation in databases under strict quality control. This work culminated in the release of the first version of the **Swiss-Prot** database in 1986, which later evolved into today's **UniProt**, whose most recent release was launched last April. He also promoted the development of other highly valuable resources such as **PROSITE**, **Cellosaurus**, and **ExPASy**.

A tireless researcher, together with **Lydie Lane** he drove the creation of **neXtProt**, a curated draft of the human proteome that has become a cornerstone of the **Human Proteome Project**. His role in the consolidation and advancement of proteomics was essential, particularly through international initiatives, and most notably within the **EuPA**. EuPA had the privilege of benefiting from his advice and support, which undoubtedly played a fundamental role in its consolidation.

We will miss his enthusiasm and wisdom.

Farewell, dear friend.

Contribution by: Fernando Corrales

HIGHLIGHTS FROM THE EUPA GENERAL COUNCIL MEETING – DECEMBER 18, 2025

At the end of last year, on December 18th there was an online General Council meeting. 17 members out of 21 with voting rights were present. During the meeting, the YPIC President and the EuPA committee chairs presented updates on their ongoing activities.

Two new initiatives were officially approved: MS-based Histone Epigenetics (MSHE) and Palaeoproteomics and Archaeology: Society for Techniques and Advances (PAASTA). At the same time, the Food and Nutrition Initiative was terminated due to inactivity.

Council members emphasized the importance of strengthening the EuPA map by including more laboratories and encouraged National Societies and institutions to consider linking the map on their own websites.

The EuPA award campaign was launched (more info on the EuPA website and social media), and the presidents of the National Societies committed to sharing the call widely and encouraging applications from their members.

Preparations for the EuPA 2026 Conference in Würzburg are progressing well. Plenary speakers have already been invited, the Industry Working Group will organize an industry-focused session, and YPIC will organize several sessions dedicated to supporting early-career researchers.

Increasing the visibility of EuPA and its National Societies remains a key priority. In March, EuPA and the Canadian National Proteomics Network will jointly host a social media workshop focused on LinkedIn.

Looking ahead, EuPA elections will take place in 2026, with all seven positions open for renewal. More details will be announced soon on the EuPA website.

Contribution by: Éva Csósz

REFLECTIONS ON HUPO 2025: PROTEOFORMS AND STRUCTURAL PROTEOMICS

Submitted by Di Tang, Lund University

HUPO has always been an inspiring and enjoyable event for me as an active researcher in mass spectrometry, proteomics, and structural biology. This year I was invited for an oral presentation in the Structural Proteomics theme of the main program, proudly representing our research group [Malmström Lab](#) from Lund University, Sweden.

This year's congress showcased a clear paradigm shift in two particular directions: a heightened emphasis on proteoforms and a broader deployment of structural proteomics to explore protein dynamics and function. Notable examples included Neil Kelleher's team (US), who successfully linked specific proteoforms to disease and elaborated the concept of $p_{(PTM)}QTL$, and Albert Heck's group (Netherlands), who directly profiled circulating IgG clonality to reveal new biological insights. These studies remind us that the mainstream of the proteomics community still routinely infers protein functionalities from peptides analysed by the bottom-up approach rather than by measuring proteins directly. Emerging evidence has indicated the value of understudied proteoforms, and there's more to explore.

Structural proteomics was the second major theme. Two plenary speakers working with structural MS methods demonstrated its huge versatility and potential: covalent painting in John Yates III (US)'s talk and limited-proteolysis MS in Paola Picotti (Switzerland)'s presentation. Both approaches capable of scaling to the proteome level and complementing with other MS modalities to probe solution-phase protein conformational dynamics across molecular events. Coupled with advances in protein folding and interaction prediction-modelling algorithms (e.g. AlphaFold, RoseTTAFold), the link between protein structure and function is being underscored as never before. In my presentation during this panel session, I discussed how structural MS can generate molecular and biological insights that conventional biophysics methods are incapable of providing, and outlined the challenges that remain in data-driven modelling with MS data. Moreover, I strongly believe that community efforts, such as the Symposium of Structural Proteomics, will provide us with more molecular insights and knowledge through structural mass spectrometry.

I particularly valued multiple pre-congress workshops and ECR mentoring sessions, which offered refreshing perspectives from both academia and industry and allowed me to reflect on my supervision skills and project management, topics that resonate strongly with the guidance I received during my PhD training. The rest of the HUPO program was equally exciting: high-

throughput sample processing for MS, the democratisation of deep plasma proteomics, novel instrument modalities from various vendors, dedicated search algorithms to illuminate the “dark proteome”, and the integration of multi-omics data to study physiology/disease in patients and model systems. One can’t help but feel rejuvenated when witnessing mass spectrometry, proteomics, and their neighbouring fields flourishing at such a pace and scale, all while maintaining a commendable level of transparency, openness and collectiveness. On the other hand, ongoing discussions also emerged regarding the handling, processing, and sharing of potentially sensitive proteomics data, particularly derived from human patient samples, during the Day of Research Initiatives session.



**Di presented in the Structural Proteomics Theme panel during Day 1 of HUPO 2025 (photo credit: Dr. Anahita Bakochi)*

3RD ANNUAL PROTEOMICS GATHERING, 16TH APRIL 2026, MADRID, SPAIN

The Young Proteomics Investigators Club (YPIC) Annual Proteomics Gathering will take place on the 16th of April at the Facultad de Farmacia, Universidad Complutense de Madrid, Spain, bringing together early-career researchers from across Europe for a dynamic and interactive proteomics meeting.

This annual event provides a welcoming platform for PhD students, postdoctoral researchers, and young scientists to present their work, exchange ideas, and discuss emerging trends and technologies across all areas of proteomics. The scientific program will feature presentations selected from submitted abstracts, with a strong emphasis on interaction and discussion.

A highlight of this year's meeting will be the presentations by the Students Proteomic Fund (SPF) finalists, who will showcase their proposed research projects and discuss their scientific objectives, methodologies, and expected impact. This session offers a unique opportunity to support and engage with promising young researchers at an early stage of their careers.

Abstract submission deadline: 7 February 2025.

Registration opens: 15 February 2025.



Submitted by: Leonie Hexamer

2026 YPIC ANNUAL PROTEOMICS GATHERING



MADRID, SPAIN
FACULTAD DE FARMACIA
UNIVERSIDAD COMPLUTENSE DE MADRID



**APRIL
16**

**ABSTRACT SUBMISSION
07.01.2026 - 07.02.2026**

**REGISTRATION OPENS
15.02.2026**



ONLY 100 PARTICIPANTS
STAY TUNED TO SECURE YOUR SPOT!

EUPA AWARDS 2026 – APPLICATIONS ARE NOW OPEN

The European Proteomics Association (EuPA) is pleased to announce that applications for the **EuPA Awards 2026** are now open. These awards recognise excellence, innovation, and commitment within the proteomics community and aim to highlight outstanding scientific achievements as well as contributions to technology, bioinformatics, outreach, and networking. The application period runs from **18 December 2025 to 14 February 2026**.

The **EuPA Best Doctoral Thesis Award** is intended to honour a young PhD researcher who has presented an outstanding doctoral thesis in any field of proteomics within the last two calendar years. Applicants must be members of a EuPA member society and must not have previously applied for this award. This prize celebrates exceptional early-career research and the next generation of leaders in proteomics.

With the **EuPA Bioinformatics for Mass Spectrometry Award**, EuPA recognises a researcher who has made an outstanding contribution to the development of bioinformatics tools or methods for mass spectrometry. Applicants must be members of a EuPA member society and must not have received the EuPA Breakthrough in Proteomics Award within the previous three years.

The **EuPA Breakthrough in Technology and Proteomics Award** aims to reward a discovery that has significantly transformed the field of proteomics and expanded the frontiers of knowledge. Eligible breakthroughs may include widely adopted workflows, technological developments in mass spectrometry, or outstanding research with a major impact on science, technology, or society. Applicants must be members of a EuPA member society and must not have been awarded this prize within the last three years.

EuPA also invites applications for the **Science Outreach Award**, which supports and acknowledges individuals or groups engaged in science outreach activities. This award is intended to recognise past engagement and enable future initiatives that communicate scientific topics to non-scientific audiences, such as children, patients, or the general public.

Finally, the **EuPA Vision and Commitment Award (VACA)** honours an early career researcher who has demonstrated exceptional commitment to networking and supporting young European scientists in proteomics. Eligible applicants must be members of a EuPA member society, have no more than seven years of work experience since completing their MSc or PhD degree, and must not have previously received the VACA award.

Researchers are warmly encouraged to apply or to share this opportunity with eligible colleagues. Full details on the awards, eligibility criteria, and application procedures can be found on the EuPA Awards website.

The applicant must have had no more than seven (7) years of work experience since completing their MSc or PhD degree.

The applicant must not have previously been awarded the EuPA VACA award.



2026 EuPA Awards Applications

Application opening: 18th of December 2025
Application deadline: 14th of February 2026

1. Best Doctoral Thesis Award
2. Bioinformatics for Mass Spectrometry
3. Vision and Commitment Award
4. Breakthrough in Technology and Proteomics Award
5. Science Outreach Award

find more information
<https://eupa.org/eupa-awards-campaign/>



Submitted by: Luisa Schmidt

INTERNATIONAL DAY OF WOMEN AND GIRLS IN SCIENCE – FEBRUARY 11TH

In honor of the International Day of Women and Girls in Science on February 11, the European Proteomics Association (EuPA) is proudly launching a special digital showcase to celebrate the women shaping the future of our field.

Starting February 11, EuPA will feature one standout contribution per day across our social media platforms and a brief highlight of the research's significance. From groundbreaking methodologies to clinical breakthroughs, we are highlighting the vital impact of women-led research in proteomics and the life sciences.

On the big day, we will unveil a special presentation on EuPA's social media platforms. The slide will feature the publication titles and the names of the First and Last authors.

We invite our members to contribute to this initiative. If you would like to see a specific publication or researcher featured in our digital tribute, please send to education@eupa.org the publication title (or a link to the paper) and a brief sentence on why this work inspires you.

Stay tuned to our social media channels for the daily reveals!

Submitted by: Maike Langini

UPCOMING EVENTS

EUPA 2026 Congress, November 29-December 3, Würzburg, Germany



**Proteomic Forum
EuPA 2026 Würzburg** XV Annual Congress
of the European Proteomics
Association

29 November – 3 December 2026

Congress Centrum Würzburg | Germany
www.proteomic-forum.com

 Deutsche Gesellschaft für
Proteomforschung e.V.

 **EUPA**
EUROPEAN PROTEOMICS ASSOCIATION

2026 Finnish Proteomics Symposium, 12-13 March, Espoo, Finland



2026 Finnish Proteomics Symposium 

 **Hotel Nuuksio**
Espoo, Finland

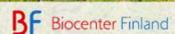
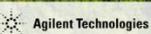
 **12-13 March 2026**
Early registration opens: 16.12.2025
Abstract submission opens: 1.1.2026

Keynote Speakers

-  **Lotta Happonen**,
Lund University
-  **Sajjalisa Kangasjärvi**,
University of Helsinki
-  **Kirsi Savijoki**,
University of Helsinki

- Experts and enthusiasts from Finland and beyond.
- Keynotes on human, plant, and microbial proteomics.
- Insights from Finnish core facilities.
- Oral talks, posters, and networking.

More info  **Register**  **Submit abstract** 

YPIC Annual Proteomics Gathering, April 16, Madrid, Spain

2026 YPIC ANNUAL PROTEOMICS GATHERING



MADRID, SPAIN
FACULTAD DE FARMACIA
UNIVERSIDAD COMPLUTENSE DE MADRID



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ProteoVilamoura, May 20-22, Vilamoura, Portugal

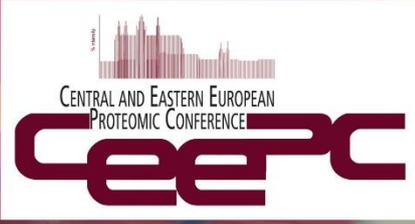
Proteo
Vilamoura

**6th Joint Meeting of Spanish, French
and Portuguese Proteomics Societies**
20-22 May 2026, Portugal

Organized by



20th Central and Eastern European Proteomics Conference, June 14-17, Zagreb, Croatia



**20th Anniversary
CEEPC**

Twenty Years of Proteomics, Genomics and Metabolomics

MSBM Summer School, July 5-10, Dubrovnik, Croatia



Fundamentals & Applications in Mass Spectrometry

Summer School - Dubrovnik

5th - 10th July 2026

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| Mass Spectrometry Basics <ul style="list-style-type: none">Mass AnalyzersIonization SourcesIon MobilitySeparations MethodsMass Spec Imaging | Tandem MS <ul style="list-style-type: none">Mass Spectrometry SystemsMS Data ProcessingProteomicsLipidomicsGlycomics |
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Learn More and Sign Up Here:





SEMM WORKSHOP
The Proteomics Revolution
Advancing Science and Society

June 29th- July 1st 2026 Quark Hotel, Milan

Proteomics is revolutionizing our understanding of life and disease. This workshop will showcase the multifaceted applications of proteomics, illustrating the latest breakthroughs in deciphering the intricate world of proteins and their crucial roles across diverse biological systems.

Speakers

Angela Bachi IFOM Institute of Molecular Oncology, IT
Petra Beli Institute of Molecular Biology, DE
Isabel Teresa Cabruja Thermo Fisher Scientific, IT
Annamaria Cattaneo University of Milan, IT
Fabio Conforti Humanitas University, IT
Fabian Coscia Max Delbrück Center, DE
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Sara Zanivan MD Anderson Cancer Center, US

Sessions

PROTEOMICS IN HEALTH AND DISEASES:
from Biomarkers to Mechanisms and
Therapies

SPATIAL PROTEOMICS: Unlocking Spatial
Information of Proteomes and their
Dynamics in Cells and Tissues

DECODING CELLULAR SIGNALING: Insights
from Proteomics

THE PROTEOMICS REVOLUTION:
Technological Innovation & Computational
Advancements