



XLV ANNUAL MEETING Chilean Society for Biochemistry and Molecular Biology

2022

Program

Conferences

Opening conference:
“Running from the law: how to evade the genome sentinel Rho during stress” - **Irina Arsttimovitch**, Department of Microbiology and The Center for RNA Biology, The Ohio State University, Columbus, Ohio, USA

Osvaldo Cori Conference:
“Epigenetic control of gene transcription during cell lineage commitment” - **Martín Montecino**, Institute of Biomedical Sciences and Millennium Institute Center for Genome Regulation CRG, Faculty of Medicine amnd Faculty of Life Sciences, Universidad Andres Bello, Santiago, Chile

Severo Ochoa Conference:
“Cell senescence in development, regeneration and cancer” - **Manuel Collado Rodríguez**, Laboratory of Cell Senescence, Cancer and Aging, Health Research Institute of Santiago de Compostela, IDIS, Santiago de Compostela, Spain



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Symposia

Symposium “Plant Defense”

Chair: Lorena Pizarro, Universidad de O’Higgins,
San Fernando, Chile

1. “Uncovering new roles for cytokinin in plant immunity”

Maya Bar, Department of Plant Pathology and Weed Research, Plant Protection Institute, Agricultural Research Organization, Volcani Institute, Rishon LeZion, Israel

2. “Incidence of bacterial canker in sweet cherry orchards of the O’Higgins Region of Chile: Varietal susceptibility and genetic characteristics of the causal agent”-

Boris Sagredo, Centro Regional Rayentué, Instituto de Investigaciones Agropecuarias (INIA), Rengo, Chile

3. “Identification of defense-responsive genes in sweet cherry against Pss strains with different virulence levels”-

Alan Zamorano, Laboratorio de Fitovirología, Departamento de Sanidad Vegetal, Facultad de Ciencias Agronómicas, Universidad de Chile, Santiago, Chile

4. “Involvement of the cell wall and its dynamic in plant defense against aphids”- **María Francisca Blanco-Herrera,**

Centro de Biotecnología Vegetal, Facultad de Ciencias de la Vida, Universidad Andres Bello, Santiago, Chile



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Symposium “Unveiling the complex world of RNA”

Chair: Marcelo López-Lastra, Pontificia Universidad Católica de Chile, Santiago, Chile

1. “Evaluation of the impact of m6A methylation on RNA structure”- **Bruno Sargueil**, Université Paris Cité, CNRS, Paris, France

2. “Modified RNAs as Molecular Tools for Structural and Functional Studies of aminoacyl transferases and methyltransferases enzymes”- **Mélanie Etheve-Quelquejeu**, Laboratoire de Chimie et Biochimie Pharmacologiques et Toxicologiques, Université Paris Cité, Paris, France

3. “Mechanochemistry of the HIV-1 reverse transcriptase on an RNA substrate, one molecule at a time”- **Andrés Bustamante**, Laboratorio de Virología Molecular y Celular, Instituto de Ciencias Biomédicas, Facultad de Medicina, Universidad de Chile, Santiago, Chile

4. “Regulation of retroviral non-canonical messenger RNA translation by RNA-binding proteins”- **Marcelo López-Lastra**, Laboratorio de Virología Molecular, Departamento de Enfermedades Infecciosas e Inmunología Pediátrica, Escuela de Medicina, Pontificia Universidad Católica de Chile, Santiago, Chile



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Symposium “DNA viruses and cancer: signaling pathways involved in cancer progression”

Chair: Francisco Aguayo, Universidad de Chile,
Santiago, Chile

1. “Molecular pathways altered by early expressed proteins from Human Papillomavirus”- **Marcela Lizano Soberón**, *Unit of Biomedical Research in Cancer, Instituto Nacional de Cancerología and Biomedical Research Institute, Universidad Nacional Autónoma de México*

2. “Regulation of extracellular matrix components during human papillomaviruses mediated carcinogenesis” -
Enrique Boccardo Pierulivo, *Laboratory of Oncovirology, Department of Microbiology, Instituto de Ciencias Biomédicas, Universidade de São Paulo, São Paulo, Brazil*

3. “High-risk human papillomavirus (HR-HPV) and Epstein Barr virus (EBV) co-presence in oral cancer” - **Julio Cesar Osorio Patiño**, *Universidad del Valle, Cali, Colombia*

4. “Crosstalk between persistent viruses and tobacco smoke in carcinogenesis”- **Francisco Aguayo González**, *Laboratory of Oncovirology, Virology Program, Institute of Biomedical Sciences, Faculty of Medicine, Universidad de Chile, Santiago, Chile*



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Symposia

Symposium “Yeast functional genomics”

Chair: Francisco Salinas, Universidad Austral de Chile,
Valdivia, Chile

- 1. “The fitness landscape of telomere variation”-** **Gianni Liti**, Université Côte d'Azur, CNRS, INSERM, IRCCAN Nice, France
- 2. “Deciphering the molecular bases of wort adaptation in novel lager yeast hybrids”-** **Jennifer Molinet**, Laboratorio de Genética Molecular, Departamento de Biología, Facultad de Química y Biología, Universidad de Santiago de Chile, Santiago, Chile
- 3. “A cohesive panel of 142 telomere-to-telomere genome assemblies unfolds the structural genetic diversity at the species-scale in *Saccharomyces cerevisiae*”-** **Gilles Fischer**, Sorbonne Université, Francia
- 4. “The metabolic and genomic making of yeast biodiversity”-** **Chris Todd Hittinger**, Laboratory of Genetics, DOE Great Lakes Bioenergy Research Center, Center for Genomic Science Innovation, J. F. Crow Institute for the Study of Evolution, Wisconsin Energy Institute, University of Wisconsin-Madison, Madison, WI, USA



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Symposium “Chaos, panic and (dis)order in protein evolution”

Chair: César A. Ramírez-Sarmiento & Pablo Galaz-Davison, Pontificia Universidad Católica de Chile, Santiago, Chile

1. “Computational approaches to study protein-protein interactions regulating the expression and function of membrane proteins”- **Ariela Vergara**, Center for Bioinformatics, Simulation and Modeling, Faculty of Engineering, Universidad de Talca, Talca, Chile

2. “Alpaca-derived nanobody against viral infections” - **Naphak Modhiran**, School of Chemistry and Molecular Biosciences, University of Queensland; Australian Institute for Bioengineering and Nanotechnology; Brisbane, Australia

3. “Metamorphic proteins don’t just wiggle wiggle, they (re)fold” - **César A. Ramírez-Sarmiento**, Institute for Biological and Medical Engineering (IIBM), Pontificia Universidad Católica de Chile, Santiago, Chile

4. “Conformational sampling with AlphaFold2 by phylogenetic amnesia”- **Matías R. Machado**, Protein Engineering Unit, Institut Pasteur de Montevideo, Montevideo, Uruguay

5. “FrustraEvo: Assessing protein families divergence in the light of sequence and energetic constraints”- **María Inés Freiberger**, Protein Physiology Lab, Departamento de Química Biológica, Facultad de Ciencias Exactas y Naturales, Universidad de Buenos Aires-CONICET-IQUIBICEN, Buenos Aires, Argentina



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Symposia

Symposium “New insights in non-coding RNAs in prokaryotic and eukaryotic cells”

Chair: Valentina Parra, Universidad de Chile, Santiago, Chile; Mauricio Latorre, Universidad de O'Higgins, Rancagua, Chile

1. “Computational tools for identifying and characterizing non-coding RNAs in prokaryotic and eukaryotic organisms”

- **Raúl Arias-Carrasco**, Programa Institucional de Fomento a la Investigación, Desarrollo e Innovación (PIDi), Universidad Tecnológica Metropolitana, Santiago, Chile

2. “Non-coding RNA global transcriptional regulatory network activated by iron in *Enterococcus faecalis*” - **Victor Aliaga-Tobar**

Bioengineering Laboratory, Institute of Engineering Sciences, Universidad de O'Higgins, Rancagua, Chile

3. “Studying the non-coding transcriptome of endocrine pancreatic cells from zebrafish” - **Estefanía Tarifeño-Saldivia**, Gene Expression and Regulation laboratory (GEaRLab), Departamento de Bioquímica y Biología Molecular, Facultad de Ciencias Biológicas, Universidad de Concepción, Chile

4. “RNA at the Epicenter of Genetic Information: from epigenetic to epitranscriptomic regulation in development and cancer” - **Paulo P. Amaral**, Insper Institute of Education and Research, Brazil



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New members session

“Mitochondrial Ca²⁺ overload in the neurodegeneration associated with early-onset familial Alzheimer’s disease”

Marioly Müller, Departamento de Tecnología Médica,
Facultad de Medicina, Universidad de Chile, Santiago, Chile

“Epigenetics effects of obesity in the neurons that regulate feeding behavior” - Estefanía Tarifeño-Saldivia, Departamento de Bioquímica y Biología Molecular, Facultad de Ciencias Biológicas, Universidad de Concepción, Concepción, Chile

“SYSTEMIX: Systems biology center for the study of extremophile communities from mining tailings”- Mauricio Latorre, Universidad de O’Higgins, Rancagua, Chile

“Evolution of enzyme regulation and the mechanistic drift hypothesis: The case of the activation by AMP in archaeal ADP-dependent Sugar Kinases”- Gabriel Vallejos-Baccelliere, Laboratorio de Bioquímica y Biología Molecular, Departamento de Biología, Facultad de Ciencias, Universidad de Chile, Santiago, Chile