47th LORNE CONFERENCE ON PROTEIN STRUCTURE AND FUNCTION 2022

6 - 10 February 2022

The Delegate Book Is Sponsored By: Thermo Fisher

SCIENTIFI



2022 Conference T-Shirt Design by Weixi Gu

The illustration presents a human brain, constructed from depictions of structures of the SARM1 protein in its inactive octameric state, as determined by cryo-electron microscopy. The different colours reflect brain anatomy. SARM1 is an executioner in axon degeneration. It is held in an inactive state in healthy axons, maintaining axon health. Upon injury, SARM1 is activated to initiate axon degeneration; this is depicted by white SARM1 molecules radiating from the brain. Axon degeneration leads to axon pathology in several neurodegenerative diseases, such as Alzheimer's and Parkinson's disease. Inhibiting SARM1 is a way to block axon loss, hence a therapeutic candidate to treat a wide range of neurodegenerative diseases.

Figley, M. D*., Gu, W*., Nanson, J. D*., Shi, Y*., Sasaki, Y., Cunnea, K., et al. (2021). SARMI is a Metabolic Sensor Activated by an Increased NMN/NAD+ Ratio to Trigger Axon Degeneration. Neuron 109, 1118–1136. doi:10.1016/j.neuron.2021.02.009.

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WELCOME & IMPORTANT INFORMATION

We are very excited to welcome you to the 47th Lorne Conference on Protein Structure and Function 2022.

This year's meeting is a hybrid meeting taking place at the Mantra Lorne and via Pheedloop.

Bus Transfers:

For arrival, buses will depart from Melbourne Airport and Southern Cross Station at 11:00 AM and 11:30 AM respectively on Sunday, 6th of February (AEDT).

At the conclusion of the conference, buses will depart from Lorne to Southern Cross Station and Melbourne Airport at 9:00 AM on Thursday 10th of February (AEDT).

Venue:

All posters and Sponsor exhibitions will take place in the Convention Centre.

Plenary presentations will take place in the Heritage Ballroom, with the Heritage rooms reserved as extra viewing spaces.

In-Person Posters:

There are three poster sessions over day two and three of the conference. You have been allocated a session and you will be required to stand by your poster during that session to answer questions and meet colleagues with similar research interests. At the conclusion of your allocated poster time, you will be required to pack up your poster so the next lot of presenters can use the space.

Virtual Platform:

The virtual platform is available to ALL attendees and will house all e-posters for the duration of the conference. At the conclusion of the conference, content will be available on the virtual platform for post-conference viewing, except for presentations where publishing has been withheld.

Parents Viewing Room:

The Walter and Eliza Hall Institute of Medical Research will again be supporting parents with young children attending the Lorne Proteins Conference in 2022. The room will have space for activities and play for children while parents will be able to listen to the presentations with the same AV projection. The Lorne Proteins Committee once again applauds this initiative. We are grateful to The Walter and Eliza Hall Institute of Medical Research for offering sponsorship to this initiative again that will offer a significant enhancement of access for all parents at the meeting.

Exhibitor Prize Draw:

Tickets can be earned from exhibition booths and placed in the competition box at the registration desk for the prize draw at the end of the conference.

Please note, for every FULLY filled in exhibitor prize draw card the Lorne Proteins Committee will donate \$5 to the Lorne Hospital.

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MAPS AND LAYOUTS

MANTRA LORNE





Cryo-electron tomography reveals a phase-separated protein degradation microcompartment at the ER membrane. Data courtesy of Dr. Benjamin Engel, formerly Max Plank Institute for Biochemistry, now Helmholtz Zentrum München. Data visualization with Thermo Scientific Amira" Software.

Resolve protein structures inside cells

Cryo-electron tomography allows you to visualize and study proteins in their functional cellular environments at unprecedented resolution. This 3D imaging technique provides insights into complex supramolecular structures and assemblies that cannot be achieved by conventional purification and structural imaging methods.

The Thermo Scientific[™] Aquilos[™] 2 Cryo-FIB is a cryo-DualBeam[™] (focused ion beam/scanning electron microscope) system dedicated to the preparation of frozen, thin lamella samples from biological specimens, now with automated workflow software. It maintains the vitrified cellular sample's structural integrity and ensures its accessibility for tomographic imaging in a cryo-transmission electron microscope such as the newly released Thermo Scientific Krios[™] G4 Cryo-TEM.

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Conference Program

Sunday, February 6th

2:00 PM REGISTRATION OPEN

3:30 PM OPENING REMARKS

SPARROW SESSION 3:45 PM – 5:45 PM Chairs: Lindsay Sparrow & Leann Tilley

Alastair G Stewart

The six steps of the complete F1 -ATPase rotary catalytic cycle

Onisha Patel

Structural basis for small molecule targeting of Doublecortin Like Kinase 1 with DCLK1-IN-1

Senthil Arumugam

Early Endosomal Acrobatics (EEA) of EEA1 drives an emergent time-keeping in endosomal maturations

Stephanie Gras

Spike specific T cell mediated immunity in vaccinated and COVID-19 recovered individuals

5:45 PM WELCOME RECEPTION

YOUNG INVESTIGATOR AWARD SESSION 7:45 PM - 9:00 PM

Chairs: Renae Ryan & Richard Birkinshaw

Josh Hardy

Viral metamorphoses: using cryo-EM to understand the maturation of flaviviruses

Lisanne M. Spenkelink

The E. coli helicase does not use ATP during replication

Raphael Trenker

Structures of the active HER2/HER3 receptor complex reveal dynamics at the dimerization interface induced by binding of a single ligand

Katrina A. Black

Ion currents through Kir potassium channels are gated by anionic lipids



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9:00 PM TRADE MIXER

END OF DAY 1

Monday, February 7th

7:00 AM BREAKFAST

SESSION 3 8:30 AM - 10:30 AM

Chairs: Jon Oakhill & Erinna Lee

Chuck Sanders

Super-Trafficking of an Ion Channel as a Disease Mechanism

Fiona Whelan

Periscope Proteins' at the host:microbe interface

Bernhard C. Lechtenberg

Structural basis of chain-type specificity and regulation of the Gordon-Holmes syndrome RBR E3 ubiquitin ligase RNF216

Jeanne Stachowiak

Stochastic mechanisms in membrane organization and curvature

10:30 AM MORNING TEA

SESSION 4 11:00 AM - 1:00 PM

Chairs: Fasseli Coulibaly & Sarah Piper

Jian Payandeh

Trapping and visualization of outer membrane protein folding intermediates

Andrew M. Ellisdon

Autoinhibition of the RAS master regulator neurofibromin revealed using cryo-EM

Ashleigh Davey

Improving safety and efficacy of chimeric antigen receptor (CAR)-T cell therapies

Allison Kraus

High resolution structures of infectious mammalian prions

1:00 PM LUNCH

1:15 PM SYNCHROTRON USERS GROUP MEETING

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3:00 PM THERMO FISHER WORKSHOP

4:00 PM POSTER SESSION A

6:00 PM DINNER AT LEISURE

8:00 PM POSTER SESSION B & TRADE MIXER

END OF DAY 2

Tuesday, February 8th

7:00 AM BREAKFAST

SESSION 7 (Session 1 of Satellite Meeting) 8:30 AM - 10:30 AM

Chairs: Jane Allison & Yu Heng Lau

Adrian Elcock

Toward molecular simulations of the bacterial cell

Lynn Kamerlin

Loop Dynamics and Evolution in the Protein Tyrosine Phosphatases Superfamily

Elisa Fadda

Fine-tuning the Spike: Role of the nature and topology of the glycan shield in the structure and dynamics of SARS-CoV-2 S

10:30 AM MORNING TEA

SESSION 8: LIGHTNING TALKS 11:00 AM - 1:00 PM

Chairs: Nicole De Weerd & Brett Collins

Ashleigh S. Paparella

Inhibition of Clostridium difficile TcdA and TcdB toxins with transition state analogues

Chris Horne

Membrane permeabilisation is mediated by distinct epitopes in mouse and human orthologs of the necroptosis effector, MLKL

David M Thal

Structural and dynamic mechanisms of allostery at the M4Â muscarinic acetylcholine receptor



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Protein stability & aggregates assessed by tracking changes in secondary structure motifs at particular wavenumbers.



Protein concentration quantified over a linear concentration range that extends from 0.1 to >200 mg/mL.





Higher order structure analysis quantified with the fractional content of different secondary structure motifs.

Protein structural similarity to detect < 2% Structural difference between samples.



Rhiannon Morris

Structural and functional analysis of target recognition by the lymphocyte adaptor protein LNK

Adam M. Damry

Electrodetection of small molecules by conformation-mediated signal enhancement

Melanie Deitrich

Blocking malaria transmission with nanobodies

Josh Ramsay

Structure-function studies of regulatory factors controlling quorum sensing and horizontal gene transfer in nitrogen-fixing symbiotic bacteria

Lizelle Lubbe

Insight into somatic angiotensin-1 converting enzyme structure and dynamics revealed by cryo-EM

Jacob S. Lewis

Cryo-EM studies of replication origin activation

Session 2: Computational Biology 11:00 AM – 1:00 PM Chairs: TBC

Megan O'Mara

Lipid modulation of neurotransmitter transporter function

Carlos Rodrigues

Computational Tools to Help Understand and Map Protein-Protein Interactions

Sarah J. Piper

Dynamic drug targets: Using Cryo-EM data and MD simulations to create realistic 3D animations of GPCR complexes

Albert Ardevol

structure and in-silico based protein engineering

Haibo Yu

Disruption of water networks is the cause of human/mouse species selectivity in urokinase plasminogen activator (uPA) inhibitors derived from hexamethylene amiloride (HMA)



Carus Lau

Structural Basis of Rapid and Voltage Dependent Inactivation of HERG Potassium Channels

Stephanie Portelli

Harnessing RNA polymerase protein structure to predict drug resistance in tuberculosis and COVID-19

1:00 PM	LUNCH	
1:00 PM	TWIST BIOSCIENCE WORKSHOP	
2:00 PM	COMPUTATIONAL BIOLOGY WORKSHOPS	
4:00 PM	POSTER SESSION C	
6:00 PM	DINNER AT LEISURE	

THE EUROPE SESSION 10 7:45 PM - 9:00 PM

Chairs: Charlie Bond & Begoña Heras

Adrian Mulholland

Multiscale simulation of enzyme catalytic mechanisms: dynamics, evolution and design

N. Amy Yewdall

Approximating the nucleoli with proteins and RNA: shaping condensate dynamics with ATP:Mg2+

Robert Tampé

Machineries and Supercomplexes in Cellular Quality Control and Adaptive Immunity



10:00 PM TRADE MIXER

END OF DAY 3



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Wednesday, February 9th

7:00 AM BREAKFAST

SESSION 11 8:30 AM - 10:30 AM

Chairs: James Murphy & Bostjan Kobe

Hao Wu

Inflammasomes - the next frontier

Chi-Min Ho

Structural Parasitology of Malaria Parasites

Alisa Glukhova

Understanding ligand binding to adenosine receptors using cryo-EM

Henry Maun

allosteric inhibition via exosite antibodies

10:30 AM MORNING TEA

SESSION 12 11:00 AM - 1:00 PM

Chairs: Emily Furlong & Debnath Ghosal

Elizabeth Villa

Opening Windows into the Cell: Bringing Structure into Cell Biology using Cryo-Electron Tomography

Thomas Ve

Crystal and Cryo-EM structures provide insight into how pro-neurodegenerative SARM1 is activated and cleave NAD+

Phillip Pymm

Development of Antibody-Based Therapeutics for COVID-19

Junyu Xiao

Molecular mechanism of secretory IgM and IgA

1:00 PM	ORACLE WORKSHOP
1:00 PM	LUNCH
3:00 PM	BRUKER WORKSHOP

LEACH LECTURE 4:00 PM - 5:30 PM

Chair: Melissa Call

Leann Tilley

Seeing is believing: From cellular machinery to molecular movies to malaria therapies

5:30 PM	POSTER PRIZES AND CONFERENCE CLOSING
7:30 PM	CONFERENCE DINNER
9:00 PM	SUPPER PARTY

END OF DAY 4

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Oryx protein crystallization robots from Douglas Instruments feature **a** greater choice of experiments than any other robot including

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INVITED SPEAKERS



Prof Adrian Elcock University of Iowa, USA



Dr Elisa Fadda Maynooth University, Ireland



Prof Chi-Min Ho Columbia University, USA



Prof Lynn Kamerlin Uppsala University, Sweden



Dr Allison Kraus Case Western Reserve University, USA



Prof Adrian Mulholland *University of Bristol, UK*



Dr Jian Payandeh Genetech Inc., USA



Prof Charles Sanders Vanderbilt University, USA

INVITED SPEAKERS



A/Prof Jeanne Stachowiak The University of Texas, USA



Prof Robert Tampé Goethe University Frankfurt, Germany



Prof Leann Tilley University of Melbourne, Australia



A/Prof Elizabeth Villa University of California San Diego & Howard Hughes Medical Institute, USA



Prof Hao Wu Boston Children's Hospital, USA



Dr Junyu Xiao Peking University, China

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SESSION SPEAKERS

Dr Albert Ardevol CSIRO, Australia

Dr Senthil Arumugam Monash Biomedical Discovery Institute, Monash University, Australia

Dr Katrina Black Walter and Eliza Hall Institute of Medical Research, Australia

Dr Ashleigh Davey Walter and Eliza Hall Institute of Medical Research, Australia

Dr Adam Damry Australian National University, Australia

Dr Melanie H. Dietrich Walter and Eliza Hall Institute of Medical Research

Dr Andrew M. Ellisdon Monash University, Australia

Prof Stephanie Gras La Trobe University, Australia

Dr Joshua Hardy Walter and Eliza Hall Institute of Medical Research, Australia

Dr Chris R. Horne Walter and Eliza Hall Institute of Medical Research, Australia Dr Carus Lau Victor Chang Cardiac Research Institute, Australia

Dr Bernhard Lechtenberg Walter and Eliza Hall Institute of Medical Research, Australia

Dr Jacob Lewis Francis Crick Institute, Australia

Dr Lizelle Lubbe University of Cape Town, South Africa

Dr Henry Maun Genentech Inc., USA

Dr Rhiannon Morris Walter and Eliza Hall Institute of Medical Research, Australia

Dr Ashleigh S. Paparella Albert Einstein College of Medicine, USA

Dr Onisha Patel Walter and Eliza Hall Institute of Medical Research, Australia

Dr Phillip Pymm Walter and Eliza Hall Institute of Medical Research, Australia

Dr Sarah Piper Monash Institute of Pharmaceutical Sciences, Monash University, Australia

SESSION SPEAKERS

Dr Stephanie Portelli Baker Heart and Diabetes Institute, University of Queensland, Australia

Dr Joshua P. Ramsay Curtin University, Australia

Dr Carlos Rodrigues Baker Heart and Diabetes Institute, University of Queensland, Australia

Dr Lisanne Spenkelink University of Wollongong, Australia

Dr Alastair Stewart Victor Chang Cardiac Research Institute, Australia

Dr David Thal Monash Institute of Pharmaceutical Sciences, Monash University, Australia

Dr Raphael Trenker University of California, USA

Dr Thomas Ve Griffith University, Australia Dr Fiona Whelan University of Adelaide, Australia

Dr N. Amy Yewdall Radboud University, Netherlands

Dr Haibo Yu University of Wollongong, Australia

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Booth 1



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Booth 15



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Booth 11

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Booths 13 & 14

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Booth 8

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