

Invited Speakers

Name	Institution	Expertise
Zbigniew Dauter	Macromolecular Crystallography Laboratory – National Cancer Institute – Argonne, USA	X ray data collection and processing.
Paul Emsley	MRC-LMB, Cambridge, UK	Model building and validation
Richard Garratt	IFSC, Sao Carlos, Brazil	Handling and analysis of structural data
Ronan Keegan	CCP4/STFC, Oxford, UK	Phasing, automated structure solution
Eugene Krissinel	CCP4/STFC, Oxford, UK	Macromolecular structure analysis and interpretation
Victor Lamzin	EMBL, Hamburg, Germany	Automatic electron density interpretation
Andrey Lebedev	CCP4/STFC, Oxford, UK	Phasing, twinning, refinement and model building
Andrew Leslie	MRC-LMB, Cambridge, UK	X ray data processing
Garib Murshudov	MRC/LMB, Cambridge, UK	Crystallographic refinement
Rob Nicholls	MRC/LMB, Cambridge, UK	Restraints in refinement
Raj Pannu	Leiden University, The Netherlands	Crystallographic refinement
Randy Read	University of Cambridge & Cambridge Institute for Medical Research, UK	Phasing (experimental/molecular replacement)
Andrea Thorn	University of Cambridge & Cambridge Institute for Medical Research, UK	Phasing, twinning and refinement
Clemens Vornrhein	Global Phasing, UK	Phasing; automated structure solution

Local Staff

Name	Institution	Expertise
Horacio Botti	IP Montevideo, Uruguay	Applications of crystallography in Structural Biology
Alejandro Buschiazzo	IP Montevideo, Uruguay	Applications of crystallography in Structural Biology
Nicole Larrieux	IP Montevideo, Uruguay	Crystallization of macromolecules
Felipe Trajtenberg	IP Montevideo, Uruguay	Applications of crystallography in Structural Biology