

## AUC2019 Symposium programme

	Gender		Level		
	Female	Male	Student	Emerging	Established
Registrations	26% (19)	74% (54)			
Session chairs	38% (6)	62% (10)	0	3	13
Talks	31% (13)	69% (28)	5	13	23
Full talks	42% (8)	58% (11)	1	5	13
Short talk	20% (5)	80% (17)	4	8	10

## Tuesday 27 August

1600 - 1900	Symposium registration
1900 - 1910 E5 Engineering	<b>Welcome address</b> Ren Dobson, Chair AUC2019
1910 - 2000 E5 Engineering	<b>Session 1: Svedberg Lecture</b> Chair Ren Dobson

Wednesday 28 August - E5 Engineering Core.

0930-1055	<b>Session 2: Complementary methods: Scattering techniques (room E5)</b> <b>Chairs: Karen Fleming &amp; Grant Pearce</b>
0930-0950	<b>Jill Trehwella</b> - <i>Reliable biomolecular structural modelling with small-angle scattering</i>
0950-1005	<b>Christine Ebel</b> - <i>Development of a SANS strategy for the study of membrane proteins: application to a prokaryotic NADPH oxidase homolog</i>
1005-1020	<b>Ivo Nischang</b> - <i>Hydrodynamic and light scattering study of very disperse and non-ideal macromolecule populations - Example of sodium carboxymethyl celluloses</i>
1020-1035	<b>Elizabeth Rodriguez</b> - <i>Higher order structure and conformational changes in biotherapeutics</i>
1035-1055	<b>Olwyn Byron</b> - <i>The spins: bacterial aldehyde-alcohol dehydrogenase forms spiral complexes critical for activity</i>
1055-1125	Morning Break
1125-1315	<b>Session 3: Method, hardware &amp; software development (room E5)</b> <b>Sponsored by Beckman Coulter</b> <b>Chairs: Borries Demeler &amp; Akash Bhattacharya</b>
1125-1145	<b>Tom Laue</b> - <i>High concentration IgG solutions</i>
1145-1205	<b>Walter Stafford</b> - <i>SEDANAL v7.1; Non-ideality at high concentrations, new features, and improvements</i>
1205-1225	<b>John Correia</b> - <i>Analysis of nonideality: Insights into fitting of high concentration therapeutic proteins in human serum</i>
1225-1240	<b>John Philo</b> - <i>Dealing with strong non-ideality in self-association: lessons learned from teduglutide</i>
1240-1255	<b>Wenqi Li</b> - <i>Analytical ultracentrifugation reveals poly acidic or basic amino acids sequence alters protein's hydrodynamic properties</i>
1255-1315	<b>Karen Fleming</b> - <i>HullRad: a fast, new tool for prediction of hydrodynamic parameters from structure</i>
1315-1405	Lunch
1325-1400	<b>Izon Workshop (room E6)</b>
1405-1530	<b>Session 4: Nanoparticles (room E5)</b> <b>Chairs: Helmut Cölfen &amp; Johannes Walter</b>
1405-1425	<b>Helmut Cölfen</b> - <i>Emulsification of ultra-flexible microemulsions (UFME) triggered by soft Centrifugation</i>
1425-1440	<b>Guillermo González-Rubio</b> - <i>Analytical ultracentrifugation for understanding the role of cluster nature in the seed-mediated growth of gold nanocrystals</i>
1440-1500	<b>Johannes Walter</b> - <i>Accessing core-shell properties and functionalization of nanoparticles by means of analytical ultracentrifugation</i>
1500-1515	<b>Louis Weber</b> - <i>Analysis of Ti/NbCN nanoparticles extracted from microalloyed steels by SV-AUC</i>
1515-1530	<b>Xufeng Xu</b> - <i>Concentrated colloidal nanoparticles in analytical ultracentrifugation</i>
1530-1555	Afternoon Break (in the atrium)
1555-1730	<b>Session 5: Biological applications: Protein-ligand/nucleic acid interactions (room E5)</b> <b>Sponsored by Izon</b> <b>Chairs: Grant Pearce &amp; Philipp Koldewey</b>
1555-1615	<b>Luitgard Nagel-Steger</b> - <i>Ligand interactions of amyloid-<math>\beta</math> protein studied by analytical ultracentrifugation in concert with microscale thermophoresis</i>
1615-1630	<b>Fareeda Barzak</b> - <i>Modelling APOBEC3-ssDNA bound complexes using small-angle X-ray scattering</i>
1630-1645	<b>Richard Kingston</b> - <i>Models for the site-specific treatment of cooperative ligand binding by proteins</i>
1645-1700	<b>Vlad Dinu</b> - <i>Sedimentation velocity studies of two salivary glycoproteins: possible structure-function effects of food flavour</i>
1700-1715	<b>Chris Horne</b> - <i>The repressor, depressor of sialic acid catabolism</i>
1715-1730	<b>Sarah Atkinson</b> - <i>Biophysical characterisation of novel Flavivirus antivirals that target host nuclear transport</i>
1730-1900	Mixer & Poster session
1900-2300	Symposium dinner – Riccarton House (buses from UC)

Thursday 28 August

<b>0900-1020</b>	<b>Session 6: Polymers (room E5)</b> <b>Chairs: Trushar Patel &amp; Walter Stafford</b>
0900-0925s	<b>Simon Wawra</b> - <i>Determination of two-dimensional size distributions of plasmonic nanoparticles via the optical back coupling method</i>
0925-0950	<b>Igor Perevyazko</b> - <i>Hydrodynamic analysis of co-polymers with a gradient structure: solution properties, molar mass and conformation</i>
0950-1005	<b>Xiaodong Ye</b> - <i>Characterization of linear and cyclic polystyrene sulfonates using analytical ultracentrifugation</i>
1005-1020	<b>Jennifer Wakefield</b> - <i>AUC analysis of chitosan and aminocellulose for use in archaeological wood conservation</i>
1020-1050	Morning Break (in the atrium)
<b>1050-1300</b>	<b>Session 7: Biological Applications: Protein complexes (room E5)</b> <b>Chairs: Tatiana Soares da Costa &amp; Sarah Atkinson</b>
1050-1110	<b>Davide Comoletti</b> - <i>A proteomic screen of neuronal cell surface molecules reveals IgLONs as structurally conserved interaction modules at the synapse</i>
1110-1130	<b>Trushar Patel</b> - <i>Towards studying nano-scale structures of Flaviviral non-coding RNAs-host protein complexes.</i>
1130-1150	<b>Ute Curth</b> - <i>E. coli primase and DNA polymerase III holoenzyme are able to bind concurrently to a primed template during DNA replication</i>
1150-1210	<b>Yee Mok</b> - <i>Tracking polyglutamine and polyalanine aggregates in cells using fluorescence-detected analytical ultracentrifugation</i>
1210-1230	<b>Grant Pearce</b> - <i>How does Rubisco activase activate Rubisco?</i>
1230-1245	<b>Akash Bhattacharya</b> - <i>What happened to the hiv capsid? A tale of armored viruses, goldilocks zones, molecular staples and traitorous co-factors</i>
1245-1400	Lunch (in the atrium)
<b>1300-1355</b>	<b>Beckman Coulter workshop (room E6)</b>
<b>1400-1530</b>	<b>Session 8: Novel uses of AUC (room E5)</b> <b>Chairs: Olwyn Byron &amp; Jill Trehwella</b>
1400-1420	<b>Alexander Bepperling</b> - <i>High concentration AUC analysis of biopharmaceuticals</i>
1420-1435	<b>Tess Malcolm</b> - <i>Metal-dependent dynamic equilibrium: A mechanism for regulation of the Plasmodium M17 aminopeptidases?</i>
1435-1450	<b>Arthur Gadon</b> - <i>Characterisation of high molecular weight hop proanthocyanidins using analytical ultracentrifugation</i>
1450-1505	<b>Grzegorz Piszczek</b> - <i>Single molecule versus bulk detection: Interferometric light scattering (iSCAMS) and AUC applied to study protein-protein interactions</i>
1505-1525	<b>Chad Brautigam</b> - <i>Modern approaches to the detection of ligand-induced protein conformational changes using AUC</i>
1525-1545	Afternoon Break (in the atrium)
<b>1544-1800</b>	<b>Session 9: Biological Applications: Protein structures (room E5)</b> <b>Chairs: Ren Dobson &amp; Luitgard Nagel-Steger</b>
1545-1605	<b>David Goldstone</b> - <i>Structural and biophysical characterisation of the master regulator TRIM28</i>
1605-1625	<b>Beata Wielgus-Kutrowska</b> - <i>Adenylosuccinate synthetase from Helicobacter pylori: biochemical and biophysical characteristics including comparison of C- and N-terminal His-tag variants</i>
1625-1645	<b>Stephen Perkins</b> - <i>The solution structure of the human IgG2 subclass is distinct from those for human IgG1 and IgG4, providing an explanation for their discrete functions</i>
1645-1700	<b>Valentina Spiteri</b> - <i>Unravelling the solution structures and stabilities of therapeutic antibodies with and without glycans</i>
1700-1720	<b>Mike Griffin</b> - TBA
1720-1800	<b>Closing comments and business meeting</b>