			Wednesday 9 th July		
11:45		Check in, registration and sandwiches			
			Stream 1	Stream 2	
12:45	13:45	Plenary	Plenary The respiratory enzyme chain: a triumph for biophysics Professor Sir John Walker FRS (MRC, Cambridge)		
13:45	15:15		PROBE MICROSCOPY/AFM	MEMBRANE PROTEINS / MAGNETIC RESONANCE	
			Chairs: Stephanie Allen, Dr Phil Williams	Chairs: Antony Watts, Ann Dixon	
13:45	14:15		Vertical Probe Force Microscopy: High-Speed Non-Contact To 4π Holo Tweezer AFM	Activation and Inactivation of a Potassium Channel	
			Mervyn Miles (University of Bristol)	Ann McDermott (Columbia University)	
14:15	14:35		Nanomechanics and interfaces of biological systems with dynamic AFM: from native membranes to live cells. Sonia Contera (University of Oxford)	Daniel Nietlispach (University of Cambridge)	
14:35	14:55	Session 1	Unravelling mechano-allostery using atomic force microscopy David Brockwell (University of Leeds)	Interaction of the allosteric ligand chlorin e6 with the G protein coupled receptor rhodopsin Judith Klein-Seetharaman (University of Warwick)	
14:55	15:05		High Resolution Imaging of Membrane Protein Structure using Scanning Probe Microscopy Khizar Sheikh (University of Leeds)	Investigating how lipids modulate G protein coupled receptor (GPCR) function. Juan H. Bolivar (University of Oxford)	
15:05	15:15		Nanomechanical Investigation of Soft Biological Cell Adhesion using Atomic Force Microscopy Lefteris Siamantouras (University of Warwick)	Connecting cell signalling, alternative splicing and apoptosis: a structural approach Cyril Dominguez (University of Leicester)	
15:15	15:45		Coffee & posters		

15:45	17:15		MODELLING AND THEORY	FIBROUS STRUCTURES
			Chairs: Mark Rodger, Jonathan Essex	Chairs: Louise Serpell, Adam Squires
15:45	16:15		The sequence-dependence of DNA structure	Multicomponent Dipeptide Hydrogels
			and flexibility: lessons from the ABC3	Dave Adams (University of Liverpool)
			dataset.	
			Charles Laughton (University of Nottingham)	
16:15	16:35		Can we a priori predict the conductances of	The structure of cross-ß tapes and tubes formed by a
			potassium ion channels?	fragment of alpha-synuclein
			Phil Fowler (University of Oxford)	Kyle Morris (University of Warwick)
16:35	16:55	Session 2	Fluctuating Finite Element Analysis:	Protein-lego: The self-assembly of linear repeat
		36331011 2	Continuum modelling of Biomacromolecules.	proteins in fibrous nanostructures.
			Robin Richardson (University of Leeds)	Ewan Main (Queen Mary College, London)
16:55	17:05		Modelling chemotactic motion of cells in	Cystatin C and Alzheimer's Disease
			biological tissues	Abi Williams (University of Sheffield)
			Bakhtier Vasiev (University of Liverpool)	
17:05	17:15		Mutual Modulation between Embedded	A de novo designed self assembling peptide nanotube
			Receptor Clustering and Ligand Binding in	system
			Lipid Membranes	Natasha Burgess (University of Bristol)
			Salvador Tomas (ISMB and Birbeck, University	
			of London)	
17:15	18:15		Poster Flash Presentation	ons
18:15	18:45	Wine		
19:15	20:15	Dinner		

Thursday 10 th July					
10:00	· · · · · · · · · · · · · · · · · · ·				
		Professor Ernst Bambe	rg,		
	(Max Planck Institute of Biophysics, Frankfurt				
11:45		SINGLE MOLECULE IMAGING	SCATTERING / HYDRODYNAMICS		
		Chairs: Mark Leake, Rob Cross	Chairs: David Scott, Olwyn Byron		
10:45		New single-molecule imaging methods for	A SAXS Perspective of Large Amplitude Motions in		
		studying DNA processing	Proteins		
		Achillefs Kapanidis (University of Oxford)	Pau Bernado (CBS, Montpelier)		
11:05		Interferometric scattering microscopy: From	The solution structures of MASP (mannose-binding		
			lectin-associated serine protease) and MBL (mannose-		
			binding lectin) provides insight on the activation of the		
		·	lectin pathway of complement		
			Ruodan Nan (University College, London)		
11:25	Session 3		Utilising neutrons for the study of a membrane protein		
		•	complex Tim Knowles (University of Birmingham)		
		• •	Tilli Kilowies (Offiversity of Birthingham)		
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11:35		· ·	Oral 1		
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11:45		,	Oral 2		
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	11:45 10:45	11:45 10:45 11:05 Session 3	10:00 Microbial Rhodopsins: Molecular Mech Professor Ernst Bambe (Max Planck Institute of Biophysi 11:45 SINGLE MOLECULE IMAGING Chairs: Mark Leake, Rob Cross New single-molecule imaging methods for studying DNA processing Achillefs Kapanidis (University of Oxford) Interferometric scattering microscopy: From high-speed single particle tracking to label- free detection, imaging and tracking of single proteins Philipp Kukura (University of Oxford) Single-molecule visualization of transcription initiation by human mitochondrial RNA polymerase Andrey Revyarkin (University of Leicester) Single molecule TIRFM reveals the dynamic behaviour of G-protein coupled receptors and potassium ion channels in live cultured cells and tissue explants Gregory Mashanov (National Institute for Medical Research)		

			Alice Pyne (UCL)	
11:45	12:15		Coffee & Posters	
12:15	13:45		MODELLING AND THEORY	EM / TOMOGRAPHY
			Chairs: Mark Rodger, Jonathan Essex	Chairs: Elena Orlova and Robert Ford
12:15	12:45		Biomolecular simulation: interactions,	Why direct electron detection makes such a
			dynamics and spectroscopy	difference in electron cryo-microscopy.
			Jonathan Hirst (University of Nottingham)	Greg McMullan (MRC LMB Cambridge)
12:45	13:05		Multicell theory to calculate the entropy of	New achievements in techniques - Achievements in
			biomolecular hydration	structural EM
			John Higham (University of Manchester)	Elena Orlova (Birkbeck College, London)
13:05	13:25		Multi-scale simulations of bacterial	Uncoating of clathrin-coated vesicles is mediated
		Session 4	membranes: successes and outlook for the	through the action of the molecular chaperone, Hsc70,
			future	together with its co-factor, auxilin.
			Syma Khalid (University of Southampton)	Corinne Smith (University of Warwick)
13:25	13:35		Functional dynamics of hexameric helicases	Structure of Mitochondrial Ribosome, Large Subunit
			probed by hydrogen exchange and simulation	Alexey Amunts (MRC Laboratory of Molecular Biology)
			Gael Radou (University of Leeds)	
13:35	13:45		Global Low Frequency Protein Motions in the	
			Allosteric Regulation of CRP/FNR family	
			transcription factors	
42.45	44.45		Tom McLeish (Durham University)	:
13:45	14:45	Lunch & British Biophysical Society AGM		
14:45	15:45	Structure and molecular biology of secretion nanomachines in Gram-negative bacterial		machines in Gram-negative bacterial
			pathogens	
			Plenary: Professor Gabriel Wak	sman FRS
			(Birkbeck College, Lond	
15:45	17:15		SINGLE MOLECULE IMAGING	FIBROUS STRUCTURES
			Chairs: Mark Leake, Rob Cross	Chairs: Louise Serpell, Adam Squires
15:45	16:15	Session 5	Membrane mechanisms as revealed by	Protein-based Fibres and Tubes by Design
			single-molecule tracking in living cells	
			Akihiro Kusumi (Kyoto University)	Dek Woolfson (University of Bristol)

16:15	16:35		Investigating the Molecular Mechanism of Adaptive Immunity in T-cell Triggering with Single-molecule Imaging Steven Lee (University of Cambridge)	The mechanism of amyloid fibril fragmentation and the stability of amyloid fibrils toward breakage Wei Feng Xue (University of Kent)
16:35	16:55		Molecular membrane dynamics studied by super-resolution STED microscopy Christian Eggeling (University of Oxford)	Structural analysis of fibrillar biomaterials using polarised Raman and infrared spectroscopy on aligned samples Adam Squires (University of Reading)
16:55	17:05		Dynamics of the bound head of myosin 5 during processive motion along actin revealed by interferometric scattering microscopy Joanna Andrecka (University of Oxford)	James Carr-Smith (University of Birmingham)
17:05	17:15		One (patchy) ring to rule them all: investigating the structure and function of the bacterial cell division machinery using 3D super-resolution microscopy Seamus Holden(EPFL)	
17:15	17:45		Coffee & Posters	
17:15 17:45	17:45 19:15		SELF ASSEMBLY Chair: Jennifer Potts & Dek Woolfson	SCATTERING / HYDRODYNAMICS Chairs: <i>David Scott, Olwyn Byron</i>
			SELF ASSEMBLY	•
17:45	19:15	Session 6	SELF ASSEMBLY Chair: Jennifer Potts & Dek Woolfson Assembly of membrane pore-forming complexes	Chairs: David Scott, Olwyn Byron Optimizing utilizable information from small angle solution scattering of biological macromolecules
17:45	19:15 18:15	Session 6	SELF ASSEMBLY Chair: Jennifer Potts & Dek Woolfson Assembly of membrane pore-forming complexes Helen Saibil (Birkbeck College, London) SasG forms a single chain protein nanorod of tunable length	Chairs: David Scott, Olwyn Byron Optimizing utilizable information from small angle solution scattering of biological macromolecules Emre Brookes (UTHSCSA) Formation of bacterial translocon: a view from the end of scattered beam

		protocell model Dora Tang (University of Bristol)	
19:05	19:15	Sugars self-assembling like a proteins Steve Harding (University of Nottingham	Oral 2
19:30	20:15	Conference Gala	Dinner

			Friday 11 th July	
09:15	10:45	Session 7	SELF ASSEMBLY	EM / TOMOGRAPHY
			Chairs: Jennifer Potts & Dek Woolfson	Chairs: Elena Orlova and Ford
09:15	09:45		The Design of Highly Symmetric Self- Assembling Protein Materials Todd Yeates (UCLA)	Investigating the mechanisms of the proteasome by cryo-EM Paula da Fonseca (MRC-LMB, Cambridge)
09:45	10:05		Structural and evolutionary versatility in protein complexes with uneven stoichiometry Joseph Marsh (EMBL, EBI)	Cryo-EM study of a polysaccharide chain length- determining membrane protein, Wzz. Robert Ford (University of Manchester)
10:05	10:25		Guiding the folding pathway of DNA origami. Jon Bath (University of Oxford)	Cryo-EM analysis of the PilF ATP-ase components of the Thermus Type IV pilus assembly system Richard Collins (University of Manchester)
10:25	10:35			Structure and Mechanism of Action of BRCA2 Breast Cancer Tumour Suppressor Taha Shahid (Imperial College London)
10:35	10:45			, , , , ,
10:45	11:15		Coffee & Posters	
11:15	12:15	Britis	sh Biophysical Society, 2014 Young Investigato Tuomas Knowles (Cambridge)	
12:45		Electron Cryomicroscopy: From Molecules to Cells Professor Wolfgang Baumeister (Max Planck Institute of Biochemistry, Martinsried)		er