Monday 27 March				
9:00 AM – 1:00 PM	Early Career Satellite Event Location: Auditorium			
1:00 PM – 2:00 PM	Lunch Location: Hall D			
2:10 PM – 2:15 PM	Welcome: Martin Howard Location: Auditorium			
2:15 PM – 3:00 PM	Keynote I: Biophysical models for 3D epithelia in vertebrate embryonic development - Lisa Manning, Syracuse University Chair: Laura Machesky   Location: Auditorium			
3:00 PM – 3:30 PM	Tea and coffee break – Sponsored by Oxford Nanopore Location: Hall D			
	CELL METABOLISM AND ENERGETICS	DIFFERENTIATION AND DEVELOPMENT	COMPLEX SYSTEMS: CLIMATE, ECOLOGY AND EPIDEMIOLOGY	
	Chair: Laura Machesky   Location: Auditorium	Chair: Martin Howard   Location: Queens Suite	Chair: Andrew Turberfield   Location: Kings Suite	
3:30 PM – 4:00 PM	(Invited) Mitochondrial dynamics as a new mechanics-to-biology transduction module Sirio Dupont, University of Padua	(Invited) Navigating the Waddington Landscape: Geometric models of cell fate decisions James Briscoe, The Francis Crick Institute	(Invited) Using Critical Slowing Down to suggest statistical indicators of disease emergence and elimination Louise Dyson, The University of Warwick	

4:00 PM – 4:30 PM	(Invited) Mechanosensitive polyol pathway regulates biomolecular condensates Stéphanie Torrino, Université Côte d'Azur	(Invited) Spermatogenesis: a paradigm of stem cell regulation Ben Simons, University of Cambridge	(Invited) Eco-evolutionary dynamics of fluctuating populations Mauro Mobilia, University of Leeds
4:30 PM – 4:45 PM	Simultaneous intracellular nanorheology and nanothermometry using diamond quantum sensing Louise Shanahan, University of Cambridge	Spatial mechano-transcriptomics of the early mouse embryo Adrien Hallou, University of Cambridge	Modelling patterns and coexistence in large ecological communities Sandro Azaele, University of Padua
4:45 PM - 5:00 PM	FRETzel: a new software package for single- cell glucose uptake measurements within heterogeneous populations Adam Wollman, Newcastle University	Folding oneself into shape: Apical actomyosin contraction is sufficient to drive a furrow- shaped buckling in a curved embryonic epithelium Jocelyn Etienne, Liphy, CNRS - Univ Grenoble Alpes	Structure and function of MicA, a novel ClpC adaptor for metabolic shutdown Rivka Isaacson, King's College London
5:00 PM -5:15 PM	Metabolism, nonequilibrium thermodynamics and system dynamics Orkun Soyer, University of Warwick	Feedback mechanisms for morphogen scaling and their evolution Zena Hadjivasiliou, University College London and The Francis Crick Institute	
5:15 PM – 5:30 PM	Modelling oscillatory dynamics in cell energy metabolism Joe Rowland Adams, Lancaster University	YAP levels and dynamics control cell fate and proliferation Kirstin Meyer, University of California San Francisco	
5:30 PM – 7:30 PM	Poster Session 1 and Exhibition – Sponsored by Location: Hall D	Beckman Coulter	BECKMAN COULTER

Tuesday 28 March			
9:00 AM – 9:45 AM	Keynote II: Symmetry breaking and crypt morphogenesis in intestinal organoids - Prisca Liberali, Friedrich Miescher Institute for Biomedical Research Chair: James Briscoe   Location: Auditorium		
9:45 AM – 10:15 AM	Coffee/Tea Break Location: Hall D		
	CLOCKS, TIMERS AND CELL CYCLE DYNAMICS	CANCER, DISEASE AND AGEING	IMAGING ACROSS SCALES
	Chair: James Briscoe   Location: Auditorium	Chair: Kevin Chalut   Location: Queens Suite	Chair: Mark Leake   Location: Kings Suite
10:15 AM – 10:45 AM	(Invited) Dissecting a slow cold temperature dependent timer Martin Howard, John Innes Centre	(Invited) Cancers Exploit Diverse Mechanical Features of the Stroma Erik Sahai, The Francis Crick Institute	(Invited) Community-driven super-resolution microscopy Dylan Owen, University of Birmingham
10:45 AM – 11:15 AM	(Invited) On the role of cell rearrangements in pattern formation Berta Verd, University of Oxford	(Invited) Mechanobiology of ageing Kevin Chalut, Altos Labs	(Invited) Cell shape control: from molecules to cellular forces Ruby Peters, University of Cambridge
11:15 AM – 11:30 AM	Patterns of interdivision time correlations reveal hidden cell cycle factors Fern Hughes, Imperial College London	Insights into cancer from machine learning: translation to clinic Peter Weightman, University of Liverpool	Molecular rotors as tools to image non- classical mechanical behaviour of lipid membranes under pathogenic stress Miguel Paez Perez, Imperial College London
11:30 AM – 11:45 AM	The alternative sigma factor RpoD4 pulses at division, linking the clock and the cell cycle in cyanobacteria Chao Ye, University of Warwick	Fluctuating methylations clocks allow for the quantification of the evolutionary dynamics in blood cancers Calum Gabbutt, Institute of Cancer Research	A label free method to measure lipid membrane dynamics of giant unilamellar vesicles Freya Turley, Cardiff University

11:45 AM – 12:00 PM	Optical synchronization of synthetic genetic clocks Roberto Di Leonardo, Sapienza University of Rome	Far from equilibrium approach to complex neurovascular dynamics in healthy ageing and dementia Juliane Bjerkan, Lancaster University	PolyScope: a 3D-printed minimal microscopic system for microswimmer tracking Wesley Shao, University of Bristol
12:00 PM – 12:15 PM	Bilateral feedback in oscillator model is required to explain the coupling dynamics of Hes1 with the cell cycle Andrew Rowntree, University of Manchester	<b>Direct observation of Shelterin dynamics and</b> <b>T-loop formation at telomeres</b> Matthew Newton, The Francis Crick Institute	Correlative atomic force microscopy with structured illumination microscopy for the investigation of nanoscale features of tuneable bacterial outer membrane models loanna Mela, University of Cambridge
12:15 PM – 1:15 PM	Lunch and Exhibition Location: Hall D		
12:30 PM – 1:15 PM	BBS AGM Location: Auditorium		
12:40 PM – 1:00 PM	Sponsored Industry Session I Location: Hall D Tweezers and single molecules: How to visualize and manipulate single biomolecules in real-time (Emma Verver, LUMICKS) Enabling Life Science Research via the Application of Photonics Technologies (Andy Keatin, HORIBA)		
1:15 PM – 2:00 PM	Keynote III: Epithelial mechanobiology from the bottom up - Xavier Trepat, Institute for Bioengineering of Catalonia Chair: Kevin Chalut   Location: Auditorium		

	CHROMATIN AND GENE REGULATION	PATTERNS, WAVES, TRANSPORT AND FLOWS	ENGINEERING BIOMOLECULES, CELLS AND ORGANOIDS
	Chair: Martin Howard   Location: Auditorium	Chair: Pietro Cicuta   Location: Queens Suite	<b>Chair:</b> Andrew Turberfield   <b>Location:</b> King's Suite
2:15 PM – 2:45 PM	(Invited) Understanding the link between 3D gene structure and transcription by computer simulations Davide Marenduzzo, University of Edinburgh	(Invited) Algal phototaxis and the evolution of multicellularity Ray Goldstein, University of Cambridge	(Invited) Patterning DNA-based artificial cells with reaction-diffusion Lorenzo Di Michele, University of Cambridge
2:45 PM – 3:15 PM	(Invited) Mathematical modelling of transcription Jane Mellor, University of Oxford	(Invited) Towards controlling bacterial cell behaviours using electricity and light Munehiro Asally, University of Warwick	(Invited) Life-like motility in droplet systems Nathalie Katsonis, University of Groningen
3:15 PM – 3:30 PM	Gene silencing regulation by heterochromatin compaction Ander Movilla Miangolarra, John Innes Centre	<b>Odd dynamics of living chiral crystals</b> Alexander Mietke, University of Bristol	Membrane transport processes control the kinetics of enzymatic pH clock reactions confined within lipid compartments Paul Beales, University of Leeds
3:30 PM – 3:45 PM	Simulating 3D chromatin structure at genomic rearrangements in cancer Chris Brackley, University of Edinburgh	<b>Tuneable metachronal coordination in</b> <b>Platynereis dumerilii larvae</b> Rebecca Poon, University of Exeter	A Universal Method for Analysing Copolymer Growth Benjamin Qureshi, Imperial College London
3:45 PM – 4:00 PM	Atomistic simulations of supercoiled DNA under tension Matthew Burman, University of York	The interplay of photosynthesis and phototaxis in a model microswimmer Stephen Williams, University of Warwick	<b>Bioinspired cytoskeletal active networks</b> Isabella Guido, University of Surrey

4:00 PM – 4:15 PM	Single-molecule imaging of nucleosome dynamics during DNA replication Dominika Gruszka, University of Oxford	How precise is patterning with noisy morphogen gradients? Roman Vetter, ETH Zürich	Two-dimensional positioning and patterning with a molecular printer made from DNA Rafael Carrascosa Marzo, University of Oxford
4:15 PM – 4:45 PM	Tea/Coffee Break - Sponsored by Oxford Nanopore Location: Hall D		
4:45 PM – 5:30 PM	Keynote IV: Designing Biological Circuits for Multicullularity - Michael Elowitz, California Institute of Technology Chair: Pietro Cicuta   Location: Auditorium		
5:30 PM – 7:30 PM	<b>Poster Session 2 and Exhibition – Sponsored by</b> Location: Hall D	Lumicks	LUWŠCK2

## Wednesday 29 March

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9:00 AM – 9:45 AM	The Tom McLeish Lecture - Keynote V: How would a biophysicist design a cell division machinery? - Petra Schwille, Max Planck Institute of Biochemistry Chair: Olwyn Byron   Location: Auditorium		
9:45 AM – 10:15 AM	Tea/Coffee Break Location: Hall D		
	BIOMOLECULAR STRUCTURE, DYNAMICS AND INTERACTIONS (SPONSORED BY HORIBA)	TISSUE FUNCTION, MECHANICS AND MECHANOSENSING	BIOLOGICAL CIRCUITS: FROM GENES TO NEURONS
	Chair: Olwyn Byron   Location: The Auditorium	Chair: Ewa Paluch   Location: Queens Suite	Chair: James Briscoe   Location: King's Suite
10:15 AM – 10:45 AM	(Invited) How do Biomolecules Walk? Sarah Harris, University of Leeds	(Invited) Emergence of cell tissue shape during development Tim Saunders, University of Warwick	(Invited) Genetic Circuits that Adapt: Design Principles and Genetic Implementation Mustafa Khammash, ETH Zürich
10:45 AM – 11:15 AM	(Invited) Minutes-long single molecule tracking in live bacteria reveals that molecular motor tug-of-war regulates elongasome dynamics and bacterial cell shape in Bacillus subtilis Stuart Middlemiss, Warwick University	(Invited) Mechanical impact of cell delamination on tissue dynamics, in developmental and tumoral contexts Magali Suzanne, University of Toulouse IIIul	(Invited) Mechanisms for functional noise in gene regulation James Locke, University of Cambridge
11:15 AM – 11:30 AM	Mechanisms of action of the DNA binding protein HU towards DNA repair and biofilm stability Elliot Chan, University of York	Patterns of contractility within a tissue modulate force-balance dynamics John Robert Davis, University of Manchester	Multiscale, dynamic and spatially-periodic coordination of gene expression in the developing spinal cord Veronica Biga, University of Manchester

11:30 AM – 11:45 AM	Complementary approaches to obtaining thermodynamic parameters from protein ligand systems: challenges and opportunities and a case for neutrons Mona Sarter, ISIS Neutron and Muon Facility- STFC	<b>The physics of root bending under gravity</b> Sujit Kumar Nath, University of Leeds	Effect propagation through gene regulatory networks Natalia Ruzickova, Institute of Science and Technology Austria
11:45 AM – 12:00 PM	A new twist on drug design: AdhE spirosomes as cross species anti-virulence targets Ester Serrano, University of Glasgow	<b>Modelling forces in confluent cell layers</b> Julia M Yeomans, University of Oxford	A dynamical systems approach to design function of gene regulatory networks Ruben Perez-Carrasco, Imperial College London
12:00 PM – 12:15 PM	Self-assembly and hydration of a β-hairpin through integrated small and wide-angle neutron scattering Harrison Laurent, University of Leeds	Revealing the missing parameters in tissue folding - Out-of-plane stresses and patterned elasticity Steph Höhn, University of Cambridge	Perception and propagation of activity through the cortical hierarchy is determined by neural variability Matthias Loidolt, University College London
12:15 PM – 1:30 PM	Lunch Location: Hall D		
1:00 PM – 1:20 PM	Sponsored Industry Session II Location: Hall D DGE-AUC: Adapting the Power of Density Gradient Separations for In-Solution Characerization (Lutz Erhardt, Beckman Coulter) Presentation by Institute of Physics, PoLNET		
1:30 PM – 3:30 PM	Physics of Life Roadmap Session Chair: Mark Leake   Location: Auditorium		

3:30 PM – 4:00 PM	Tea and Coffee Break - Sponsored by Oxford Nanopore Location: Hall D			
	BIOMOLECULAR ASSEMBLIES AND CONDENSATES (SPONSORED BY HORIBA)	EVOLUTION	MACHINE LEARNING AT THE PHYSICS/BIOLOGY INTERFACE	
	Chair: Mark Leake   Location: The Auditorium	Chair: Laura Machesky   Location: Queens Suite	Chair: Olwyn Byron   Location: King's Suite	
4:00 PM – 4:30 PM	(Invited) Location and concentration of aromatic-rich segments dictates the percolating inter-molecular network and viscoelastic properties of ageing condensates Jorge Rene Espinosa, University of Cambridge	(Invited) Does evolution have an inbuilt Occam's razor? Ard Louis, University of Oxford	(Invited) Challenges and advantages in the integration of computational biology and deep learning Pietro Lio, University of Cambridge	
4:30 PM – 5:00 PM	(Invited) Multiscale modelling of liquid-like chromatin organisation Rosana Collepardo Guevara, University of Cambridge	(Invited) Quantitatively understand microbial community traits during evolution Wenying Shou, University College London	4:30 PM – 4:45 PM Combinatorial approaches for understanding morphogenesis in 3D embryos Salvish Goomanee, Collège De France & CNRS	
5:00 PM – 5:15 PM	Counting the cold: do flowering plants encode winter memory with large nuclear assemblies? Alex Payne-Dwyer, University of York	A theory of the evolution of multi-site population resistance/rescue and how the genetic diversity of the population significantly increases its probability Bhavin Khatri, Imperial College London	4:45 PM – 5:00 PM CellPhe: a toolkit for cell phenotyping using time-lapse imaging and pattern recognition Laura Wiggins, University of York	
5:15 PM <i>–</i> 5:30 PM	A model system reveals how different linker histones direct chromatin to liquid-like or fibre-like condensed states Katherine Stott, University of Cambridge	Phenotypic noise as an evolutionary trait in bacteriophage populations Diana Fusco, University of Cambridge	5:00 PM – 5:15 PM Probing the rules of cell coordination in live tissues by interpretable machine learning based on graph neural networks Takaki Yamamoto, RIKEN Center for Biosystems Dynamics Research	

			5:15 PM – 5:30 PM	
			Topic Modeling to analyze spatial	
			transcriptomic data from Allen Human Brain	
			Alids	
			Letizia Pizzini, Oniversity of Turin and INFN	
	How the condensation behaviour of the RNA	Is the variation of intragenic DNA methylation	Physics-informed neural networks for solving	
5:30 PM – 5:45 PM	binding protein TDP-43 influences RNA binding	in Arabidopsis natural populations governed	the inverse Turing problem	
	and regulation	by genetic or epigenetic inheritance?	Antonio Matas Gil, Imperial College London	
	Martina Hallegger, The Francis Crick Institute	Amy Briffa, John Innes Centre		
	Coarse-grained simulations of biomolecular	Mechanical interactions affect the growth and	Decoding cellular identities from single-cell	
5:45 PM – 6:00 PM	condensates with the Martini forcefield	evolution of bacterial colonies	data	
	Chris Brasnett. University of Groningen	Bartlomiei Waclaw. The University of	Zoe Piran. The Hebrew University of Jerusalem	
	, , , .	Edinburgh/Dioscuri Centre IPC Warsaw		
6:15 PM – 7: 00 PM	Keynote VI: Super-resolution imaging of tran	nscription in living cells - Ibrahim Cisse, Max Planc	k Institute of Immunobiology and Epigenetics	
7.00 014 7.20 014		Drinke Recention		
7:00 PIVI - 7:30 PIVI				
	Location: Hall D			
7.00 004 44.00 004				
7:30 PM – 11:00 PM		Conterence Dinner		
		Location: Studio 1		

## Thursday 30 March

	CELL MEMBRANES, CELL ARCHITECTURE AND FORCES	IMMUNITY, RESISTANCE AND HOST/PATHOGEN DYNAMICS	SINGLE MOLECULE BIOLOGY
	Chair: Ewa Paluch   Location: Auditorium	Chair: Pietro Cicuta   Location: Queens Suite	Chair: Mark Wallace   Location: King's Suite
9:00 AM – 9:30 AM	(Invited) Control of cortical mechanics by signalling Guillaume Charras, University College London	(Invited) How do bacteria interact with cells? Clare Bryant, University of Cambridge	(Invited) Tackling Topology using TopoStats Alice Pyne, University of Sheffield
9:30 AM -10:00 AM	(Invited) Mechanical control of immune synapse organisation and function Katelyn Spillane, King's College London	(Invited) Environmental conditions define the energetics of bacterial dormancy and its antibiotic susceptibility Teuta Plizota, University of Edinburgh	(Invited) The NEOtrap, DyeCycling, NEO-FRET & Co: expanding the single-molecule toolbox Sonja Schmid, Wageningen University
10:00 AM – 10:15 AM	Friction forces determine cytoplasmic reorganization and shape changes of ascidian oocytes upon fertilization Silvia Caballero Mancebo, University College London	Learning the differences: a transfer-learning approach to predict antigen immunogenicity and T-cell receptor specificity Barbara Bravi, Imperial College London	Watching bacterial cell division one molecule at a time using vertical cells Kevin Whitley, Newcastle University
10:15 AM – 10:30 AM	Study of E-cadherin and actin dynamics during cell-cell adhesion using a cell - lipid bilayer assay Sayantika Ghosh, University of Warwick	New tools to study cell-cell interactions: Using optical tweezers and microfluidics to study malaria parasites invading human erythrocytes Emma Jones, University of Cambridge	Dimers are forever: single-molecule techniques shine light on single-protein exchange within synaptic complexes – a potential mechanism for long-term memory storage Katie Morris, University of Edinburgh

10:30 AM – 10:45 AM	Using AFM and automated image analysis to study the correlation between bacterial shape and Peptidoglycan orientation in B. subtilis Laia Pasquina Lemonche, University of Sheffield	Physiological dynamics of individual microbes under complex changing conditions Somenath Bakshi, University of Cambridge	Super-resolution single-molecule detection of biomolecules using a nanopore Kaikai Chen, University of Chinese Academy of Sciences
10:45 AM – 11:00 AM	Model of inverse bleb growth explains giant vacuole dynamics during cell mechanoadaptation Andrea Cairoli, University of Cambridge	Use of Shigella-septin interactions to explore biophysical determinants in cell-autonomous immunity Gizem Özbaykal-Güler, The London School of Hygiene and Tropical Medicine	Revealing the mechanism of supercoil relaxation by the Human Topoisomerase3α- RMI1-RM2 complex Graeme King, University College London
11:00 AM – 11:30 AM	Tea/Coffee Break - Sponsored by Oxford Nanopore Location: Hall D		
11:30 AM – 12:15 PM	Keynote VII: Our archaeal ancestry - Buzz Baum, Chair: Ewa Paluch   Location: Auditorium	Laboratory for Molecular Cell Biology	
12:15 PM – 12:30 PM	Conclusions and close Chair: Martin Howard   Location: Auditorium		
12:30 PM – 1:30 PM	Lunch and depart Location: Hall D		