



PhD Day Symposium

Thursday 13 July 2017

Talks: William Harvey Lecture Theatre,
University of Cambridge School of Clinical Medicine

Poster session: Seminar Rooms,
University of Cambridge School of Clinical Medicine

Talk Schedule

Session 1		
09:40	Alisa Molotova, CSCI PhD rep Welcome	Franklin/Chalut
09:45	Dr Alessandro Serra Multiplexing done easy: Detect miRNA or proteins directly from crude samples with Fireplex	Abcam
09:55	Stanley Strawbridge Session introduction	Smith
10:00	Moosa Qureshi Cellular Model of CEBPA N321D aims to capture Gene Expression Profile in the Transition to Pre-Leukaemic Status	Göttgens
10:15	Olivia Harris Un(MaSC)ing mammary gland development: combining novel lineage tracing and 3D imaging for studying mammary stem cells	Watson
10:30	Claudia Pama Myelination during development and learning	Káradóttir
10:45	Anastasiya Sybirna Exploring transcriptional regulators of human germ cell fate	Surani
11:00	Madeline McNamara Development of potential therapeutic strategies for Marfan Syndrome using a human iPSC-derived <i>in vitro</i> disease model	Sinha
11:15	<i>Coffee break</i>	
Session 2		
11:40	Dr Rebecca Jones Public Engagement – ways to get involved	SCI Public Engagement
11:50	Eugene Park Session introduction	Ringshausen
11:55	Serena Belluschi Molecular regulation of quiescence exit within the heterogeneous human LT-HSC compartment	Laurenti
12:10	Lucia Cordero Espinoza Stromal regulation of facultative progenitors in the regenerating liver	Huch
12:25	Dr Brian Huntly Introduction to External Speaker	
12:30	Professor Margaret Goodell - External Speaker Immortal Stem Cells and their Epigenetic Regulation	Baylor College of Medicine, Texas
13:30	<i>Lunch Buffet</i>	

Session 3		
14:15	Julia Spindel Session introduction	Reik
14:20	Anna Albiero Cartilage progenitor cells: study of DACTs proteins in human articular cartilage	McCaskie
14:35	Rebecca Caesar Functional mutation screening using CRISPR in germinal centre malignancies	Hodson
14:50	Fiona Hamey Modelling the transcriptional landscape of haematopoiesis	Göttgens
15:05	Michael Segel Contextualising OPC Ageing	Franklin
15:20	<i>Coffee break</i>	
Session 4		
15:45	Cora Olpe Session introduction	Winton
15:50	Ludovica Di Canio Regulation of oligodendrocyte lineage cell progression by the RXR γ nuclear receptor	Franklin
16:05	Caroline Oedekoven Characterisation of fate choice regulators in single mouse haematopoietic stem cells	Kent
16:20	Giovanni Canu Cell-cycle dependent mechanisms controlling differentiation of haematopoietic stem cells	Vallier
16:35	Amanda Collier Comprehensive Cell Surface Protein Profiling Identifies Specific Markers of Human Naive and Primed Pluripotent States	Rugg-Gunn
16:50	<i>End of talks</i>	
17:00	Closed PI meeting	
17:00	<i>Poster session, with refreshments</i>	
18:30	<i>Talk and poster winners announced</i>	



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Poster List

1	Theresa Bartels Systematic Characterisation of Astrocyte Heterogeneity in the Mouse Forebrain by RNAscope	Rowitch
2	Lawrence Bates Developing new methods to study the role of Oct4 in pluripotency	Silva
3	James Baye The thermodynamics of stem cell fate transitions: single-cell and collective dynamics	Chalut
4	Francesca Beaton The Bone Marrow Niche and its Manipulation for Cartilage and Bone Repair	McCaskie
5	Daniel Bode Characterisation of protein networks driving haematopoietic stem cell fate choice	Kent
6	Jan Botthof Loss of the homologous recombination gene rad51 leads to Fanconi anemia-like symptoms in zebrafish	Cvejic
7	Aracely Castillo Venzor Decoding the early cell state transitions during primordial germ cell specification	Surani
8	Amanda Dalby An Inducible Human iPSC Line for the Scaling up of Megakaryocyte and Platelet Production <i>in vitro</i> for Transfusion Medicine	Ghevaert
9	Charlene Fabian Polycomb-Group Proteins in Human Pancreatic Cell Differentiation	Rugg-Gunn
10	Sophie Frankham-Wells The interaction of a blood subpopulation and MSCs in hypoxia	McCaskie
11	Nikoletta Gkatza Post-transcriptional RNA methylation affects protein translation and response to stress	Frye
12	Myfanwy Hill Imaging Demyelination in the CNS	Franklin
13	Wajid Jawaid Deep learning models of bifurcating developmental journeys from single-cell transcriptomic data can make accurate predictions of gene perturbations	Göttgens
14	Timo Kohler Cells in Gels: Multi germ layer differentiation of clonal embryonic stem cells in biomimetic microgels	Chalut
15	Tim Lohoff Mechanisms of lineage commitment during exit from pluripotency <i>in vivo</i>	Nichols/Reik
16	Ludovica Marando The role of the DNMT3A mutation in the subversion of normal Haematopoietic Stem Cell function and generation of preleukaemia	Huntly

17	Jamie McGinn Epithelial stem cell fate plasticity in health and disease; a mouse oesophageal model	Alcolea
18	Chris McMurran Remyelination in germ-free mice	Franklin
19	Alisa Molotova Dynamic interplay between ageing OPCs and their surrounding microenvironment	Franklin/Chalut
20	Sam Myers Determining the signalling pathways that govern naive human pluripotency	Smith
21	Cora Olpe Investigating the Cancer Chemopreventive Effect of Aspirin	Winton
22	Anna Osnato Transcriptional networks variations during cell cycle progression in human embryonic stem cells	Vallier
23	Luca Peruzzotti-Jametti Transplanted induced neural stem cells ameliorate experimental autoimmune encephalomyelitis by metabolic reprogramming of mononuclear phagocytes	Pluchino
24	Blanca Pijuan-Sala eRNAs: a potential proxy for enhancer state during blood differentiation at the single-cell level	Göttgens
25	Alexander Ross Development of gut location and developmental stage specific iPSC derived human intestinal epithelial organoids for use in disease modelling, drug discovery and autologous cell therapy	Vallier
26	Antonella Santoro Hematopoietic niche remodelling in B cell malignancies	Ringshausen
27	Vijitha Sathiaselvan Characterization of ZAP70 mediated signaling pathways in distinct cellular compartments in Chronic Lymphocytic Leukaemia.	Ringshausen
28	Kirsten Scott B lymphocytes in Parkinson's Disease	Barker
29	Mairi Shepherd Single cell approaches identify the molecular network driving malignant haematopoietic stem cell self-renewal	Kent
30	Elsa Sousa Investigating the link between naïve cell identity and X-inactivation	Silva
31	Julia Spindel The role and regulation of UHRF1 during epigenetic reprogramming	Reik
32	Samantha Tilson Induced Pluripotent Stem Cell Derived Model for the Study of Non-alcoholic Steatohepatitis and Hepatitis C Virus	Vallier

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