

Tuesday 30th August

14:00 to 17:00	Registration in EXCHANGE BUILDING, JUBILEE CAMPUS, UNIVERSITY OF NOTTINGHAM
	Exhibition stands set-up in C3 and C33, The Exchange Building
15:00	Check-in at Southwell Hall, Jubilee Campus
16:00 to 17:25	Session 1, Chairperson: Professor Neil Hall
	Lecture Theatre 3, Exchange Building
16:00-16:10	Opening remarks
16:10-16:40	Prof. Neil Hall (Centre for Genomic Research, University of Liverpool)
	Genomics of Neglected Diseases
16:40-17:00	Mr. Mick Watson (ARK-Genomics, Roslin Institute)
	21 - 23 facts; the devil in the detail of microRNA-Seq
17:00-17:25	Dr. Xuemin Liu (Roche)
	GS FLX+ System
17:25-18:00	Coffee/Tea break in C3 and C33, Exchange Building
18:00 to 19:15	Session 2, Chairperson: Dr. Nick Loman
	Lecture Theatre 3, Exchange Building
18:00-18:30	Dr. Nick Loman (Centre for Systems Biology, University of Birmingham)
	Benchtop Sequencers and Bacterial Outbreaks
18:30-18:50	Dr. Andreas Heger (CGAT, University of Oxford)
	The MRC CGAT programme - Computational Genomics: Analysis and Training
18:50-19:15	Dr. Martin Seifert (Genomatix)
	ChIPseq in Disease Relevant Environment

19:30 onwards Social in Nottingham City Centre with postgraduate student guides



Wednesday 31st August

07:30 to 9:00	Breakfast in The Atrium
08:30 to 9:30	Registration and poster set-up in Exchange Building
09.00 to 10.35	Session 3, Chairperson: Professor Mark Blaxter
	Lecture Theatre 3, Exchange Building
09:00-09:30	Prof. Mark Blaxter (The GenePool, University of Edinburgh)
	Assembling environmental genomes from Illumina data
09:30-09:50	Dr. Rachel Brenchley (The University of Liverpool)
	Sequencing the genome of hexaploid wheat for SNP discovery
09:50-10:15	Dr. Raimo Tanzi (Life Technologies)
	Whole Exome Sequencing on the SOLiD™ 5500 System
10:15-10:35	Ms. Carolin Müller (University of Nottingham)
	Deep Sequencing to Study Genome Replication
10:35-11:00	Coffee/Tea break in C3 and C33, Exchange Building
11:00 to 12:15	Session 4, Chairperson: Dr. Paul Coupland
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11:00-11:30	Lecture Theatre 3, Exchange Building Dr. Paul Coupland (Wellcome Trust Sanger Institute) A Year of PacBio at the Sanger: The Story so Far
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13:45 to 15:45 Session 5, Chairperson: Dr. Guy Cochrane **Lecture Theatre 3, Exchange Building** 13:45-14:15 Dr. Guy Cochrane (EBI, Wellcome Trust Genome Campus) European Nucleotide Archive: Life on the log-scale 14:15-14:35 Mr. Thomas Hancocks & Dr. Angela Davies (Nowgen, Manchester) Professional training in bioinformatics and next generation sequencing 14:35-15:00 Dr. Holger Karas (CLCbio) CLC Genomics Gateway as Platform for Deep Sequence Analysis 15:00-15:20 Dr. Gerton Lunter (Wellcome Trust Centre for Human Genetics, Oxford) Mapping and variant calling from Illumina data 15:20-15:45 Dr. Hannes Arnold (Caliper Life Sciences) Dr. Melanie Febrer (The Genome Analysis Centre) Integrated Solutions for Automated NGS Sample Preparation 15:45-16:20 Coffee/Tea break in C3 and C33, Exchange Building 16:20 to 17:35 Session 6, Chairperson: Professor Chris Ponting **Lecture Theatre 3, Exchange Building** 16:20-16:50 Prof. Chris Ponting (MRC FGU, University of Oxford) A Transcriptomic Atlas of Mouse Neocortical Layers 16:50-17:10 Damian Kao (University of Nottingham) Use of client-side web development for presenting bioinformatics data

The Biomek Continuum - Automating Next Generation Sequencing

Poster session & Drinks reception in C3 and C33 Exchange Building

Buses depart for Conference Dinner at Trent Bridge, Nottinghamshire County Cricket

Dr. Michelle Hiscutt (Beckman Coulter)

17:10-17:35

17:35-19:00

19:15, 19:30

Club



Thursday 1st September

07:30 to 09:00 Breakfast in The Atrium

Check out of rooms in Southwell Hall

09:00 to 10:15	Session 7, Chairperson: Professor Ed Louis
	Lecture Theatre 3, Exchange Building
09:00-09:30	Prof. Ed Louis (Centre for Genetics and Genomics, University of Nottingham)
	A 'Complete' de novo Yeast Genome using Mixed NGS Platforms (454 and SOLiD)
09:30-09:50	Dr. Laura Kelly (Royal Botanic Gardens Kew, Queen Mary University of London)
	Sequencing Giant Genomes: Second Generation Sequencing Analysis of Fritillaria (Liliaceae)
09:50-10:15	Neil Ward - (Illumina)
	Illumina Sequencing - Whole genomes to amplicons
10:15 to 11:00	Coffee/Tea break in C3 and C33, Exchange Building
Before 11:30	Delegates remove posters from poster boards
	Sponsors remove exhibition materials

11:00 to 13:00 Session 8, Chairperson: Dr. Mario Caccamo Lecture Theatre 3, Exchange Building 11:00-11:30 Dr. Mario Caccamo (The Genome Analysis Centre, Norwich Research Park) Bread Wheat Chromosome-based Survey Sequencing Initiative 11:30-11:55 Dr. Scott Brouillette (Partek) Whole Transcriptome Analysis – a Start-to-Finish Solution using Partek Genomics Suite 11:55-12:15 Dr. Ian Adams (Food and Environment Research Agency) Invasive insect surveillance using 454 amplicon sequencing



12:15-12:40 Dr. Eric van der Walt (Kappa Biosystems)

An Engineered High-Fidelity DNA Polymerase Minimizes Library Amplification Bias and Improves Coverage Uniformity in Illumina Sequencing

12:40-13:00 Dr. Miriam Smith (Wellcome Trust Sanger Institute, Cambridge)

Evaluation of the Ion Torrent Personal Genome Machine Sequencer at the Wellcome Trust Sanger Institute

13:00 END