

NERC-MDIBL Environmental Genomics and Metabolomics

4-10 March 2018

Nicholson Building, University of Birmingham, UK

Course Programme*

*Please note this is a provisional course programme and the times of some events may change

Date	Time	Event		
Sunday March 4	4.00-5.00	Registration (Biosciences Building, Undercroft)		
	5.00-6.00	Plenary lecture (Lecture Theatre NG08, Biosciences Building) Professor Susan Celniker (Lawrence Berkley National Laboratory)		
	6.00-7.00	Welcome reception (Undercroft, Bioscience Building)		
Monday March 5	8.30-9.00	Registration (The Nicolson Building)		
	Combined Track (Nettlefold Room)			
	9.00-9.45	Lecture: Introduction to Environmental Genomics, Professor John Colbourne		
	9.45-10.30	Lecture: Introduction to Environmental Metabolomics, Professor Mark Viant		
	10.30-10.45	Coffee Break		
	10.45-11.45	Lecture: The challenges of working with big data and performing multi-omics studies, Professor Ben Brown		
	11.45-12.00	Attendees logon to wireless network and computer server		
	12.00-13.00	Lunch		
	13.00-13.45	Ice-breaker session , Cate Winder		
	Genomics Track (Nettlefold Room)		Metabolomics Track (Guest Room)	
	13.45-14.00	Introduction to genomics track: Joe Shaw & John Colbourne	13.45-14.00	Introduction to metabolomics track: Cate Winder
	14.00-15.00	Workshop Presentation: Library construction methods and QC, Steve Kissane	14.00-15.00	Workshop Presentation: Experimental design in environmental metabolomics, Rick Dunn
	15.00-15.55	Workshop Presentation: Introduction to automation systems, Steve Kissane	15.00-15.50	Workshop Presentation: Quality assurance and quality control, Rick Dunn
	15.55-16.05	Coffee Break	15.50-16.00	Coffee Break
16.05-17.00	Workshop Presentation: Introduction to the sequence data workflow, Jana Asselman	16.00-17.00	Workshop Presentation: Sample preparation in metabolomics, Cate Winder	
18.00-19.00	Evening Lecture (Lecture Theatre NG08, Biosciences Building) Professor Daniel Jacobson (Oak Ridge National Laboratory)			
Tuesday March 6	Genomics Track (Nettlefold Room)		Metabolomics Track (Guest Room)	
	9.00-10.00	Bioinformatics Training: Introduction to R, Vignesh Dhandapani	9.00-10.00	Workshop Presentation: Analytical technologies – mass spectrometry, Rick Dunn
	10.00-10.45	Bioinformatics Training: Visualising of sequence data for quality, Craig Jackson	10.00-11.00	Workshop Presentation: Analytical technologies – NMR, Mark Viant
	10.45-11.00	Coffee break	11.00-11.15	Coffee Break

	11.00-12.00	Bioinformatics Training: Visualising of sequence data for quality Craig Jackson	11.15-12.00	Interactive session: Sample preparation and analytical technologies, Cate Winder
	12.00-13.00	Lunch		
	Genomics Track (Nettlefold Room)		Metabolomics Track (401A, Biosciences building)	
	13.00-14.15	Workshop presentation: Review of the sequencing technology, it's strengths and weaknesses, Professor Peter Kille (University of Cardiff)	13.00-14.00	Guest Presentation: TBC, Dr Scott Haywood
	14.15-16.00	Bioinformatics Training: Visualizing complex data, Daniel Jacobson	14.00-16.00	Group activity: Experimental design, Cate Winder
	16.00-16.15	Coffee Break	16.00-16.15	Coffee Break
	16.05-17.00	Bioinformatics Training: Navigating the command line and more advanced R, Vignesh Dhandapani	16.15-17.00	Group presentations: Experimental design, Cate Winder
	18.00-19.00	Evening Lecture (Lecture Theatre NG08, Biosciences Building) Professor Laurent Keller (University of Lausanne)		
Wednesday March 7	Genomics Track (Nettlefold Room)		Metabolomics Track (Guest Room)	
	9.00-10.45	Bioinformatics Training: Data analysis using R (Part 1), Craig Jackson, with Jana Asselman	9.00-10.30	Workshop Presentation: Data processing, Ralf Weber
	10.45-11.00	Coffee Break	10.30-10.45	Coffee Break
	11.00-12.00	Bioinformatics Training: Data analysis using R (Part 2), Craig Jackson, with Jana Asselman	10.45-12.00	Workshop Presentation: Introduction to analysing metabolomics data, Jasper Engel
	12.00-13.00	Lunch		
	13.00-14.30	Workshop presentation: Computational tools to study genetic diversity in natural populations, Benjamin Peter	13.00-13.45	Workshop Presentation: Data analysis I, Introduction to data analysis, Jasper Engel
	14.30-15.30	Bioinformatics Training: Data analysis using R (Part 3), Craig Jackson, with Jana Asselman	13.45-14.45	Computer Workshop: Hands-on data analysis, Jasper Engel
			14.45-15.30	Workshop Presentation: Data analysis II, Jasper Engel
15.30-17.00	Tour: Genomics and Metabolomics Facilities	15.30-17.00	Tour: Genomics and Metabolomics Facilities	
18.00-19.00	Evening Lecture (Lecture Theatre NG08, Biosciences Building) Dr George Loizou (Head of Computational Toxicology, Health and Safety Laboratory)			
Thursday March 8	Genomics Track (Nettlefold Room)		Metabolomics Track (Guest Room)	
	9.00-10.45	Bioinformatics Training: Gene set enrichment, Jana Asselman with Albert Zhou	9.00-9.30	Workshop Presentation: Data analysis III
			9.30-12.00	Computer Workshop: Hands-on data analysis, Jasper Engel
	10.45-11.00	Coffee Break		
	11.00-12.00	Bioinformatics Training: Pathways analysis, Jana Asselman with Albert Zhou	11.00-12.00	Computer Workshop: Hands-on data analysis, Jasper Engel
	12.00-13.00	Lunch		
	13.00-15.25	Bioinformatics Training: Exploring genome sequence variation, Benjamin Peter	13.00-14.00	Guest Presentation: Copepods and climate change, Dr Dan Mayor
14.00-14.40			Workshop Presentation: Metabolite identification and databases, Ralf Weber	
15.25-15.35	Coffee Break	14.40-15.05	Workshop Presentation: Metabolomics standards initiative,	

				reporting your results and using MetaboLights, Ralf Weber
	15.35-17.00	Synthesis session: “Why not use _____ as my model species?” John Colbourne	15.05-15.15	Coffee Break
			15.15-16.30	Computer Workshop: Hands-on metabolite identification, Ralf Weber
	17.30-18.30	Special Double Bill Evening Lecture (Lecture Theatre NG08, Biosciences Building) Dr Adam Biales (United States Environmental Protection Agency & Dr Jean-Lou Dorne (European Food Safety Authority) – Omics to Transform Environmental and Health Protection		
	18.30-19.30	Pizza and Networking		
Friday March 9	Combined Track (Nicholson Building, Nettlefold Room)			
	9.00-12.00	Synthesis session: Designing multi-omics studies, lead by Ben Brown, Albert Zhou & John Colbourne, (Jasper Engel and Cate Winder assisting metabolomics track)		
	12.00-13.00	Lunch		
	13.00-14.00	Synthesis session: How do I obtain grant funding for my project? Ipshita Ghose & Sohini Chakrabortee		
	14.00-14.30	Question and answer session, Course Trainers & Guest Faculty		
	14.30-15.00	Closing ceremony		
	15.00-16.00	Optional bioinformatics help session		