







NERC-MDIBL Environmental Genomics and Metabolomics

6-11 March 2016

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	4.00-5.00	Registration					
	5.00-6.00	Plenary lecture: optional lecture, open to the general public					
Monday	8.30-9.00	Registration					
	9.00-9.45	Lecture: Introduction to Environmental Genomics					
	9.45-10.30	Lecture: Introduction to Environmental Metabolomics					
	10.30-10.45	Break					
	10.45-12.00	Lecture & workshop: Introduction to analysing multi-omics datasets					
	12.00-13.00	Lunch					
	13.00-13.40	Ice-breaker session					
		Genomics Track	Metabolomics Track				
	13.40-15.00	Workshop Presentation: Library	13.40-14.00	Introduction to metabo	olomics track,		
		construction methods and QC			_		
	15.00-16.00	Workshop Presentation:	14.00-14.45	Lecture : Experimental of	_		
		Introduction to automation systems		environmental metabolomics			
	16.00-17.00	Bioinformatics Training:	14.45-16.15	Synthesis session : Desi	gning your		
		Introduction to R		experiments			
			16.15-17.00	Lecture: Quality assura	nce and quality		
	10.00.00			control			
	19.00-20.00 Evening Lecture						
Tuesday	0.00.10.15	Genomics Track		Metabolomics Track			
	9.00-10.15	Workshop Presentation:	9.00-10.00	· · ·			
		Introduction to the sequence data		metabolomics			
	40.45.44.00	workflow	40.00.44.00	1 /			
	10.15-11.00	Bioinformatics Training: Visualising	10.00-11.00	00 Lecture (visiting speaker)			
	11.00-11.15	of sequence data for quality Coffee break					
	11.00-11.13		11.15-12.00	Cunthosis sossion: \\/or	king with NDAF		
	11.15-12.00	Bioinformatics Training : Visualising of sequence data for quality	11.15-12.00 Synthesis session: Working with N and how to obtain funding for you		_		
		(Continued)			illig for your		
	12.00-13.00	(Continued) project					
	13.00-14.15	Workshop presentation: Review of	13.00-15.00	Group 1	Group 2		
	13.00-14.13	the sequencing technology, it's	13.00-13.00	Laboratory Session:	Synthesis		
		strengths and weaknesses		Hands-on sample	session:		
	14.15-16.00	Bioinformatics Training: Visualizing	-	preparation	selecting the		
	14.15-10.00	complex data		preparation	appropriate		
		complex data			analytical		
					method for		
					your		
					experiment		
					Схретинени		
	16.00-17.00	Bioinformatics Training: Navigating	15.00-17.00	Synthesis session:	Laboratory		
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		the command line		selecting the	Session:		
				appropriate analytical	Hands-on		
				method for your	sample		
				experiment	preparation		
	19.00-20.00	Evening Lecture					
Wednesday	Genomics Track Metabolomics Track						
weunesday	9.00-10.45	Bioinformatics Training: Data	9.00-9.55 Lecture : Analytical technologies –				
	9.00-10.43	analysis using R (Part 1)	3.00 3.33	mass spectrometry			
			9.55-10.45	Lecture: Analytical technologies –			
			3.55 10.45	NMR			
	10.45-11.00	Coffee break		IVIVIIX			
	11.00-12.00	Workshop Presentation: RNA-Seq	11.00-12.00	Lecture and software demonstration:			
		alignment to individual		Data processing (XCMS, SIMS			
		transcriptomes		stitching)			
	12.00-13.00	Lunch		5555511119)			
	13.00-14.00	Workshop presentation: Statistical	13.00-13.45	Lecture: Data analysis I, Introduction to data analysis			
		considerations for analysing					
		genome-scale data					
	14.00-15.30	5		Computer Workshop: Introduction to			
		analysis using R (Part 2)		software			
			14.45-15.30	Lecture: Data analysis II			
	15.30-17.00	Synthesis Session: Why not use	15.30-17.00				
	13.30 17.00	organism for my investigation?	13.30 17.00	analysis			
		organism for my investigation.	anarysis				
	19.00-20.00	Evening Lecture					
Thursday		Genomics Track		Metabolomics Track			
	9.00-11.00	Bioinformatics Training: Gene set	9.00-9.45	Lecture: Data analysis III			
		enrichment		·			
	11.00-12.00	Bioinformatics Training : Pathways	9.45-12.00	Computer Workshop: Hands-on data analysis			
		analysis					
	12.00-13.00	Lunch					
	13.00-15.30	Bioinformatics Training : Exploiting genome sequence variation	13.00-14.00	Lecture (Visiting Speaker):			
			14.00-14.40	Lecture: Metabolite Ide	ntification and		
				databases			
	15.30-17.00	Synthesis session: "Why not extract from my sequence data?"	14.40-15.10	Lecture: Metabolomics standards			
				initiative, reporting your results and			
				using MetaboLights			
			15.10-17.00	Computer Workshop: H			
				metabolite identification	า		
	19.00-20.00	Evening Lecture					
Friday	9.00-10.30	Synthesis session: Integrating omics data					
	10.30-12.00	Bioinformatics session: Individual help					
	12.00-13.00	Lunch					
	13.00-14.00	Synthesis session: How do I obtain grant funding for my project?					
	14.00-14.45	Question and answer session					
	14.45-15.30	Closing ceremony					