

NGS Bioinformatics course at the Ill 22-26 of January						
	Monday 22/01/2018	Tuesday 23/01/2018	Wednesday 24/01/2018	Thursday 25/01/2018	Friday 26/01/2018	
08:00						
08:30	room 515	room 515	room 515	room 515	room 515	08:30
09:00	Overview of Coures - Thomas	NGS Mapping exercise Malaria drug, CNV, SNP, databases Thomas / Kathryn	RNA-Seq I Mapping and differential expression <i>Malaria Knock out analysis</i> including GO enrichment Kathryn	Galaxy analysis	Group task	09:00
09:30	Introduction to Linux Thomas			ChipSeq Kathryn		09:30
10:00	Break	Break	Break	Break	Break	10:00
10:30	Introduction to Linux	NGS Mapping exercise continued	RNA-Seq I continued	ChipSeq continued	Group task continued	10:30
11:00	Genomic viewer - Artemis Kathryn					RNA-Seq Galaxy David
11:30	Lunch break	Lunch break	Lunch break	Lunch break	prepare presentations	11:30
12:00	Lunch break	Introduction to R Thomas	RNA-Seq II Clustering and intregative analysis <i>C. difficil dataset</i> <i>R, heatmaps, GO, String, KEGG</i> Thomas	Day to day Bioinformatics: Handling of data and downloading of data set The HPCC job submission Thomas	CTT WOLFSON MED:257 SEM 2-HUGH Group presentations	12:00
12:30	Lunch break					Lunch break
13:00	Lunch break	Introduction to R Thomas	RNA-Seq II Clustering and intregative analysis <i>C. difficil dataset</i> <i>R, heatmaps, GO, String, KEGG</i> Thomas	Day to day Bioinformatics: Handling of data and downloading of data set The HPCC job submission Thomas	CTT WOLFSON MED:257 SEM 2-HUGH Group presentations	13:00
13:30	Lunch break					Lunch break
14:00	GBRC level2 Seminar series Introduction to Bioinformtics and NGS Thomas	Introduction to R Thomas	RNA-Seq II Clustering and intregative analysis <i>C. difficil dataset</i> <i>R, heatmaps, GO, String, KEGG</i> Thomas	Day to day Bioinformatics: Handling of data and downloading of data set The HPCC job submission Thomas	CTT WOLFSON MED:257 SEM 2-HUGH Group presentations	14:00
14:30	introduction to Transcriptomics John Cole					Lunch break
15:00	Break	Introduction to R Thomas	RNA-Seq II Clustering and intregative analysis <i>C. difficil dataset</i> <i>R, heatmaps, GO, String, KEGG</i> Thomas	Day to day Bioinformatics: Handling of data and downloading of data set The HPCC job submission Thomas	CTT WOLFSON MED:257 SEM 2-HUGH Group presentations	15:00
15:30	Break					Lunch break
16:00	Virus Bioinformatics David Robertson	Introduction to R Thomas	RNA-Seq II Clustering and intregative analysis <i>C. difficil dataset</i> <i>R, heatmaps, GO, String, KEGG</i> Thomas	Day to day Bioinformatics: Handling of data and downloading of data set The HPCC job submission Thomas	CTT WOLFSON MED:257 SEM 2-HUGH Group presentations	16:00
16:30	Break					Lunch break
17:00	Break	Introduction to R Thomas	RNA-Seq II Clustering and intregative analysis <i>C. difficil dataset</i> <i>R, heatmaps, GO, String, KEGG</i> Thomas	Day to day Bioinformatics: Handling of data and downloading of data set The HPCC job submission Thomas	CTT WOLFSON MED:257 SEM 2-HUGH Group presentations	17:00
	Break					Lunch break
				Group task informal meeting	Finish	