		Viral Genomics ar	nd Bioinformatics, 24 - 2	28 June 2024, Universit	y of Glasgow, Garscub	e Campus	
	Monday	Tuesday	Wednesday	Thur	rsday	Friday	
	6/24/24	6/25/24	6/26/24	6/27/24		6/28/24	
9:00	Registration (0900-0930)						
	Tea & Coffee in the Barn						
9:30	Course and Instructor Intro	Read cleaning alignment	Bash - Advanced Linux	De novo	assembly	K-mer based	Gene ontology in R
	Participants introduction	Derek Wright	Conditions & Loops			metagenomics	Pathway analysis in R
10:00	2min talk by each participant						
	Overview of viral bioinformatic					Richard Orton	
10:30	Joseph Hughes (1015-1045)						Srikeerthana Kuchi
	Login, Setup & Basic Linux	(0930-1100)	Sreenu Vattipally (0930-1100)	Sreenu Vattipally (0930-1100)		(0930-1100)	(0930-1100)
11:00	Srikeerthana Kuchi	Break (1100-1130)	Break (1100-1130)	Break (1100-1130)		Break (1100-1130)	
14.00		D. (Out of Direction County	De mana accomplia de la companya del companya de la companya del companya de la c			
11:30		Reference alignment	Script Practice Session	De novo assembly quality assessment		Build a pipeline:	Build a pipeline:
10.00		& Statistics				Ref alignment	Transcriptomics
12:00	(1045-1230)	Richard Orton				or denovo	
12:30	(1045-1230) Lunch	Richard Orton				or metagenomics	
12.50	(1230-1330)	(1130-1300)	Sreenu Vattipally (1130-1300)	Joseph Hughes (1130-1300)		All Instructors	
13:00	(1230-1330)	Lunch	Lunch	Lunch		Lunch	
10.00		(1300-1400)	(1300-1400)	(1300-1400)		(1300-1400)	
13:30	Login, Setup & Basic Linux	(1000 1100)	(1333 1333)	(1000 1000)		(1222)	
	Srikeerthana Kuchi						
14:00		Consensus & Variant Calling	Introduction to MSA	Intro. to plotting in R		Build a pipeline continued	
14.00	(1330-1430)		ma oddedon to MOA	muot to ploung in it		Bund a pipenne continued	
14:30	NGS data & QC	Richard Orton					
			Practical MSA			All Instructors	
15:00						What next on your bioinformatic journey?	
	Derek Wright (1430-1530)	(1400-1530)	David Robertson (1400-1530)	Joseph Hughes (1400-1530)		•	
15:30	Break (1530-1600)	Break (1530-1600)	Break (1530-1600)	Break (1530-1600)		Tea & Coffee (1530-1600)	
						Goodbyes	
16:00	NGS technologies	Bash - File editing &	Intro. to phylogenetic	Metagenomics	Alignment for transcriptomics		
	Kathy S. & Will R. (1600-1630)	Text processing	methods	virus discovery			
16:30	Sequencing facility tour		Tree building practicals		Differential expression		
10.00	Danny Mair (1630-1700)		visualization and annotation		analysis in R		
17:00	Social event						
	BBQ & Poster presentation	Sreenu Vattipally (1600-1730)	David Robertson (1600-1730)	Joseph Hughes (1600-1730)	Quan Gu (1600-1730)		
17:30	by CVR staff and students	, , , , , , , , , , , , , , , , , , , ,	Social event				
			Course dinner				
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	6/24/24	6/25/24	6/26/24	6/27/24		6/28/24	