Viral Genomics and Bioinformatics, 24 - 28 June 2024, University of Glasgow, Garscube Campus							
	Monday	Tuesday	Wednesday	Thur	sday	Friday	
	24/06/2024	25/06/2024	26/06/2024	27/06/2024		28/06/2024	
09:00	Registration (0900-0930)						
	Tea & Coffee in the Barn						
09:30	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)	
11:30		Reference alignment	Script Practice Session	De novo assembly quality assessment		Build a pipeline:	Build a pipeline:
		& Statistics				Ref alignment	Transcriptomics
12:00	(40.47 4000)	51.1.10.4				or denovo	
10.00	(1045-1230)	Richard Orton				or metagenomics	
12:30	Lunch	(4420.4200)	S	Joseph Hughes (1130-1300)		All Instructors	
13:00	(1230-1330)	(1130-1300) Lunch	Sreenu Vattipally (1130-1300) Lunch	Lunch		All Instructors  Lunch	
13.00		(1300-1400)	(1300-1400)	(1300-1400)		(1300-1400)	
13:30	Login, Setup & Basic Linux	(1300-1400)	(1300-1400)	(1300-1400)		(1300-1400)	
13.30	Srikeerthana Kuchi						
44.00	Onicordiana raoni	Consensus & Variant Calling	1.416410.4	Intro to platting in D		Duild a single	
14:00	(4220.4420)	Consensus & Variant Canning	Introduction to MSA	Intro. to plotting in R		Build a pipeline continued	
14:30	(1330-1430) NGS data & QC	Richard Orton					
14.30	NGS data & QC	Richard Orton	Practical MSA			All Instructors	
15:00			Fractical MISA			What next on your bioinformatic journey?	
10.00	Derek Wright (1430-1530)	(1400-1530)	David Robertson (1400-1530)	Joseph Hughes (1130-1300)		What hext on your bioinformatic journey?	
15:30	Break (1530-1600)	Break (1530-1600)	Break (1530-1600)	Break (1530-1600)		Tea & Coffee (1530-1600)	
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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			
16:20				Thus discovery	Differential expression		
16:30	Sequencing facility tour  Danny Mair (1630-1700)		Tree building practicals visualization and annotation		Differential expression analysis in R		
17:00	Social event		visualization and annotation		allalysis III K		
17.00	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	Crocke Valapany (1000-1750)	Social event	Coopii ilugiles (1000-1700)	Quair Gu (1000-1700)		
17.50	Sy 5711 Start and Stadents		Course dinner				
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