| Viral Genomics and Bioinformatics, 16 - 20 June 2025, University of Glasgow, Garscube Campus | | | | | | | |
|--|---|--------------------------------------|-------------------------|--------------------------------------|-------------------------------|--|-----------------------|
| | Monday | Tuesday | Wednesday | Thu | rsday | Friday | |
| | 16/06/2025 | 17/06/2025 | 18/06/2025 | 19/06 | 8/2025 | 20/06/2025 | |
| 09:00 | Registration (09:00-09:30) | | | | | | |
| | Tea & Coffee in the Barn | | | | | | |
| 09:30 | Welcome | Read cleaning & QC | Bash - Advanced Linux | De novo | assembly | K-mer based | Gene ontology in R |
| | Viral bioinformatics | | Conditions & Loops | | | metagenomics | Pathway analysis in R |
| 10:00 | David Robertson 09:45-10:15 | | | | | | |
| | Participants introduction | | | | | | |
| 10:30 | 10:15-11:00 | Richard Orton | Joseph Hughes | Sreenu | Vattipally | Richard Orton | Quan Gu |
| | 2min talk by each participant | 09:30-11:00 | 09:30-11:00 | 09:30-11:00 | | 09:30-11:00 | 09:30-11:00 |
| 11:00 | Break | Break | Break | Break Break | | eak | |
| | 11:00-11:30 | 11:00-11:30 | 11:00-11:30 | 11:00-11:30 | | 11:00-11:30 | |
| 11:30 | Login, Setup & Basic Linux | Reference alignment | Script Practice Session | De novo assembly quality assessment | | Build a pipeline: | |
| | | & Statistics | | | | Ref alignment, metagenomics | |
| 12:00 | | | | | | or transcriptomics | |
| | | | | | | | |
| 12:30 | Sreenu Vattipally | Richard Orton | Joseph Hughes | Sreenu Vattipally | | All Instructors | |
| | 11:30-13:00 | 11:30-13:00 | 11:30-13:00 | 11:30-13:00 | | 11:30-13:00 | |
| 13:00 | Lunch | Lunch | Lunch | Lunch | | Lunch | |
| | 13:00-14:00 | 13:00-14:00 | 13:00-14:00 | 13:00-14:00 | | 13:00-14:00 | |
| 13:30 | | | | | | | |
| | | | | | | | |
| 14:00 | Login, Setup & Basic Linux | Consensus & Variant Calling | Introduction to MSA | Introduction to plotting in R | | Build a pipeline continued | |
| | Sreenu Vattipally | | and | | | | |
| 14:30 | 14:00-14:45 | | Phylogenetics | | | All Instructors | |
| | NGS FASTQ data | | | Leavel II | | 14:00-15:00 | |
| 15:00 | Derek Wright | Richard Orton | David Robertson | Joseph Hughes | | What next on your bioinformatic journey? | |
| | 14:45-15:30 | 14:00-15:30 | 14:00-15:30 | 14:00-15:30 | | 15:00-15:30 | |
| 15:30 | Break | Break | Break | Break | | Tea & Coffee & Goodbyes | |
| | 15:30-16:00 | 15:30-16:00 | 15:30-16:00 | 15:30-16:00 | | 15:30-16:00 | |
| 16:00 | NGS technologies | Bash - File editing & | MSA | Metagenomics | Alignment for transcriptomics | | |
| | Katherine Smollet (16:00-16:30) | Text processing | and | virus discovery | Differential expression | | |
| 16:30 | Sequencing facility tour | | Tree building practical | | analysis in R | | |
| 47.00 | Katherine Smollet (16:30-17:00) Bioinformatics Hub | Jaconh Hughan | David Robertson | Croony Vottinally | Quan Gu | | |
| 17:00 | | Joseph Hughes 16:00-17:30 | | Sreenu Vattipally | Quan Gu 16:00-17:30 | | |
| 47.00 | Scottish Drinks & Snacks | Social event | 16:00-17:30 | 16:00-17:30 16:00-17:30 Social event | | | |
| 17:30 | | Social event Course dinner - Ka Pao | | CVR hot buffet | | | |
| | Mondov | | Wadnesday | Thursday | | Fu: | day |
| | Monday | Tuesday | Wednesday | 19/06/2025 | | Friday 20/06/2025 | |
| | 16/06/2025 | 17/06/2025 | 18/06/2025 | 19/00/2025 | | 20/06/2025 | |