



Clinical Genomic Data Analysis: A Practical Course

Objectives To explore and examine clinical genomic data in depth;
To develop an understanding of processes that underpin clinical genomic data analysis; and
To obtain hands on experience in identifying variants and deducing their impact.

Times	Monday	Tuesday	Wednesday	Thursday	Friday
Morning (9.30am start)	Talk: Genomics and genome structure	Lecture: Genome bioinformatics for clinicians; <i>Seave</i>	Talk: Homozygosity mapping and recessive disorders	Talk: Gene networks, protein modelling and clinical databases	Talks: Molecular biology of cancer; Cancer genomics (practicality & pathology)
Morning	Talk: How do you interpret a variant?	Workshop: Integrated Genome Viewer and assessing variant quality	Workshop: Genome filtering and homozygosity mapping	Workshop: Additional analysis tools when a result is not clear	Workshop: Cancer genomic data and analysis
Afternoon	Workshop: Interpreting variants	Talk: Genome filtering	Talk: Copy Number Variants	Talk: How to write a genomic report	Talk: Cancer genomics in research and WGS Evaluation
Afternoon (finish time will vary between 3.30 to 5pm)		Workshop: Genome filtering	Workshop: Copy Number Variants	Talk/mini-workshop Quality metrics; Clinical capture and Patient Archive	Task: Solving a family [self-assessment opportunity with individual feedback]
Afternoon (4pm)	Tour: KCCG genomics facility			Evaluation for 4-day course participants	

Please note this outline may be subject to change in line with course applications and the composition of the attendees. Specific timings and objectives for each day will be provided directly to participants.

Course Convenor: Dr Tony Roscioli. Faculty members/speakers for 18-22 July include: Prof Michael Buckley, Dr Mark Cowley, Prof Michael Field, Dr Andre Minoche, Prof Sandra O’Toole, Prof Neil Watkins. Education coordinator: Bronwyn Terrill (b.terrill@garvan.org.au).