Module Planning NGS Bioinformatics - Remote classrooms

Module Title: Pathogen Variant Calling

- → Module Lead/Co-leads/Assistants Narender, Jon
- → Summary or objectives Choosing suitable reference for alignment, variant calling, variant annotation and identifying resistance conferring variants, calculating pairwise SNP difference, determining genetic relatedness and construct phylogenetic tree
- → List of learning outcomes specific for this module (Please list 3 5 Learning Outcomes here; refer to How to writing learning outcomes)
- → Identifying suitable reference for mapping
- → Perform alignment to reference
- → Variant calling from the alignment
- → Annotating the variants with SNPeff
- → Understanding the variant calls with IGV
- → Generating pseudogenomes
- → Calculating pairwise SNP difference
- → Constructing phylogenetic tree
- → Total number of hours/days to be spent on this content: 8 hours
- → Tools/software and resources to be used
 - ◆ Software: Trimmomatic, bwa, samtools,picard, bcftools, SNPeff, SNP-sites, SNP-dists, IGV, IQTree
 - Description of datasets to be used: Mycobacterium tuberculosis
 - Other resources, or readings
- → Overview of activities and exercises (list specific practical activities which participants will do)

Lecture: Mapping to reference, Variant calling and annotation, genetic relatedness and phylogeny

Activity 1: mapping of the reads to reference

Activity 2: Call variants

Activity 3: filtering and annotation

Activity 4: identify resistance variants

Activity 5: analyze genetic relatedness

Activity 6: generate phylogeny

Appendix (include any guideline documents related to content development such as breaking down content for remote classroom format, how to write LO's, designing assignments etc...)

See links to other guidelines

-Overall trainer guideline and template documents will include information on lecture formats and recordings, contact sessions schedules, checklists, assignments and assessments, quizzes, feedback, audio-video conferencing, learning management platform, teaching assistants