

Preliminary Program

Tuesday, July 18, 2017	Wednesday, July 19, 2017	Thursday, July 20, 2017	Friday, July 21, 2017
(NG)-Sequencing at Göttingen Genomics Laboratory	An introduction to microbial genome annotation	RNA-Seq: Challenges in experimental design and analysis	Computational challenges in metagenome analysis
Tutorial: Introduction to G2L computing environment	Annotation Tutorial - Introduction to IMG	Tutorial: RNA-Seq based transcriptomics	Coffee Break
Coffee Break	Coffee Break	Coffee Break	
Tutorial: Processing of NGS raw data	Tutorial: SNP analysis, Mapping	Tutorial: RNA-Seq based transcriptomics (continued)	Tutorial: Analysis of metagenomic and metatranscriptomic data
·	Lunch	RNA-Seq: A powerful tool for (bacterial) systems biology	Closing remarks
Lunch		Lunch	Lunch snacks
NG-Sequencing: Experimental design	From genomics to comparative genomics	Metagenomics and Metatranscriptomics	Discussion / Question time
3rd Generation Sequencing on the PacBio RSII @ DSMZ	Tutorial: Comparative genomics: Phylogenomics	Tools for metagenome and metatranscriptome analysis	- End of workshop -
Coffee Break	Coffee Break	Coffee Break	
Tutorial: Assembly of prokaryotic genomes	Tutorial: Comparative genomics: Pathogenomics	Tutorial: Amplicon-based assessment of microbial community structures	
- End of 1st day -	- End of 2nd day -	- End of 3rd day -	

Social programme