

	Visualisation	Sequencing quality control	Statistical calculation	Taxonomic classification	Other	
Analyses of shotgun metagenomics data with MetaPhlAn2						Metagenomics
Taxonomy classification using Kraken2 and Bracken						Taxonomy
Analyses of shotgun metagenomics data with MetaPhlAn2						Metagenomics
GTN Training - Antibiotic Resistance Detection						Microbiology
Metaproteomics_GTN						
Metaproteomics workflow						
metaQuantome_datacreation_workflow						
WF3_VERIFICATION_WORKFLOW						
WF2_Discovery-Workflow						
MGNify amplicon rRNA-prediction + ITS						
Ecoli Comparison						
Workflow 4: Staramr						
AMR-Pathfinder						
Gene-based Pathogen Identification (release v0.1)						
Gene-based Pathogen Identification						
pox-virus-amplicon/main						
Metatranscriptomics data analysis						
MetaT						
workflow-generate-dataset-for-assembly-tutorial						
Cloud-Aerosole MT-MG Pre-Processing						
Cloud-Aerosole MT-MG Functional Profiling						
Unicycler training						
pAllori ASiam Preprocessing MT						
pAllori Blood RNA						
M. tuberculosis Variant Analysis tutorial						
Training: 16S rRNA Analysis with Nanopore Sequencing Reads						
MGNify amplicon QC						
pAllori Amplicon 16S Pre-Processing and Taxonomy Classification						
Reference genome choice (indels, coverage, depth) for short reads						
Compute Allel Frequency Table from paired reads						
Calling variants in non-diploid systems						
pAllori WGS						
MetaG_extended						
MAGs workflow						
Cloud-Aerosole MT-MG Reference Gene Catalog Building						
WGS Part In "Analyses Of Metagenomics Data - The Global Picture"						
Mtb phylogeny						
Normalization						
Feature-Count Table Normalization						
metaquantome-taxonomy-workflow						
metaquantome-function-workflow						
3: Plant virus exploration						
WF1_Database_Generation_Workflow						
WF5_Data_Interpretation_Workflow						
WF4_Quantitation_Workflow						
MetaP						
PathoGFAIR						
Main Metatranscriptomics Analysis						
PathoGFAIR (imported from uploaded file)						
Genome Annotation with Prokka [Feb 2020]						
Apollo Load Test						
Microbial Variant Calling						
From VCFs to SNP distance matrix						
From Fastqs to VCFs and BAMs						
pAllori ASiam Functional Information MT						
From BAMs to drug resistance prediction with TB-profiler						
2: Plant virus confirmation						
pAllori ASiam Community Profile MT						
bacterial-genome-assembly/main						
amr_gene_detection/main						
bacterial_genome_annotation/main						
Metagenomics assembly tutorial workflow						
Workflow 1: Further Quality Control [16S Microbial Analysis With Mothur]						
Amplicon Tutorial						
Workflow 2: Data Cleaning And Chimera Removal [16S Microbial Analysis With Mothur]						
Workflow 3: Classification [Galaxy Training: 16S Microbial Analysis With Mothur]						
Workflow 5: OTU Clustering [16S Microbial Analysis With Mothur]						
Workflow 6: Alpha Diversity [16S Microbial Analysis With Mothur]						
Workflow 7 : Beta Diversity [16S Microbial Analysis With Mothur]						
Training: 16S rRNA Sequencing With Mothur: Main Tutorial						
Taxonomy Profiling and Visualization with Krona						
Taxonomy Profiling and Visualization with Kron (release v0.1)						
Identification of the micro-organisms in a beer using Nanopore sequencing						
Taxonomic Profiling and Visualization of Metagenomic Data						
1: Plant virus detection with kraken2 (SE)						
1: Plant virus detection with kraken2 (PE)						
Taxonomy Profiling and Visualization with Phinch						
Cloud-Aerosole MT-MG Taxonomic Profiling						
quality-and-contamination-control/main						
Cloud Aerosol MT-MG Contamination Filtering						
Nanopore Preprocessing (release v0.1)						
Nanopore Preprocessing						
Halophiles workup of Comparative gene analysis						
Comparative gene analysis						
Copy Of GTN Training - Antibiotic Resistance Detection						
Flye-AMR workflow EMC/WMDI - v3.5 (imported from URL)						
Allele-based Pathogen Identification (release v0.1)						
Allele-based Pathogen Identification						
Allele-based Pathogen Identification (release v0.1.1)						
Allele-based Pathogen Identification (release v0.1.2)						
Pathogen Detection PathoGFAIR Samples Aggregation and Visualisation (release v0.1)						
Pathogen Detection PathoGFAIR Samples Aggregation and Visualisation						
Pathogen-Detection-Nanopore-All-Samples-Analysis-Juyptool						