

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