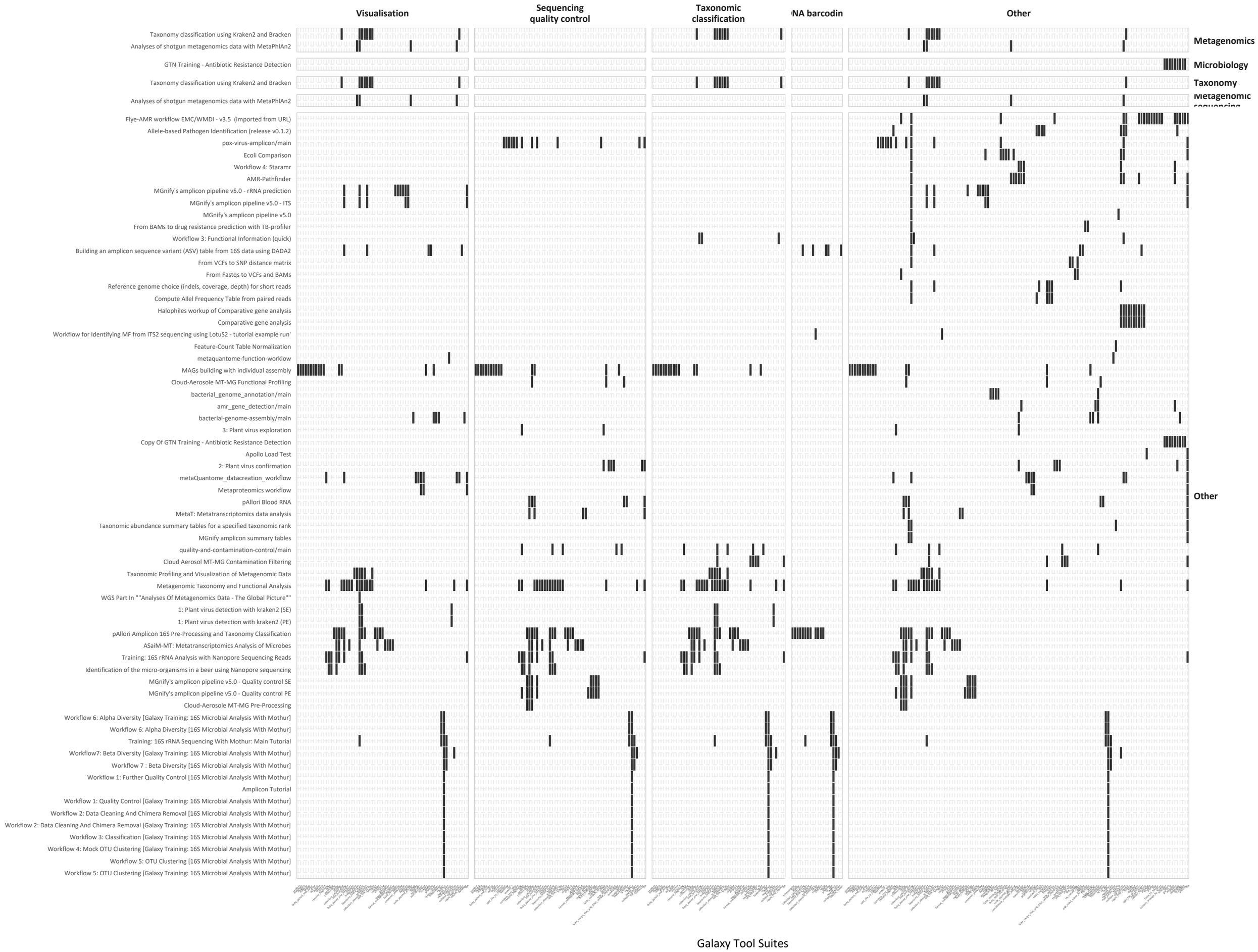
Visualisation
Sequencing quality control
Taxonomic classification
NA barcodin
Other
Metagenomics
Microbiology
Taxonomy
Metagenomic sequencing
Other
Taxonomy classification using Kraken2 and Bracken
Analyses of shotgun metagenomics data with MetaPhlAn2
GTN Training - Antibiotic Resistance Detection
Taxonomy classification using Kraken2 and Bracken
Analyses of shotgun metagenomics data with MetaPhlAn2
Flye-AMR workflow EMC/WMDI - v3.5 (imported from URL)
Allele-based Pathogen Identification (release v0.1.2)
pox-virus-amplicon/main
Ecoli Comparison
Workflow 4: Staramr
AMR-Pathfinder
MGnify's amplicon pipeline v5.0 - rRNA prediction
MGnify's amplicon pipeline v5.0 - ITS
MGnify's amplicon pipeline v5.0
From BAMs to drug resistance prediction with TB-profiler
Workflow 3: Functional Information (quick)
Building an amplicon sequence variant (ASV) table from 16S data using DADA2
From VCFs to SNP distance matrix
From Fastqs to VCFs and BAMs
Reference genome choice (indels, coverage, depth) for short reads
Compute Allel Frequency Table from paired reads
Halophiles workup of Comparative gene analysis
Comparative gene analysis
Workflow for Identifying MF from ITS2 sequencing using LotuS2 - tutorial example run'
Feature-Count Table Normalization
metaquantome-function-worklow
MAGs building with individual assembly
Cloud-Aerosole MT-MG Functional Profiling
bacterial_genome_annotation/main
amr_gene_detection/main
bacterial-genome-assembly/main
3: Plant virus exploration
Copy Of GTN Training - Antibiotic Resistance Detection
Apollo Load Test
2: Plant virus confirmation
metaQuantome_datacreation_workflow
Metaproteomics workflow
pAllori Blood RNA
MetaT: Metatranscriptomics data analysis
Taxonomic abundance summary tables for a specified taxonomic rank
MGnify amplicon summary tables
quality-and-contamination-control/main
Cloud Aerosol MT-MG Contamination Filtering
Taxonomic Profiling and Visualization of Metagenomic Data
Metagenomic Taxonomy and Functional Analysis
WGS Part In ""Analyses Of Metagenomics Data - The Global Picture""
1: Plant virus detection with kraken2 (SE)
1: Plant virus detection with kraken2 (PE)
pAllori Amplicon 16S Pre-Processing and Taxonomy Classification
ASaiM-MT: Metatranscriptomics Analysis of Microbes
Training: 16S rRNA Analysis with Nanopore Sequencing Reads
Identification of the micro-organisms in a beer using Nanopore sequencing
MGnify's amplicon pipeline v5.0 - Quality control SE
MGnify's amplicon pipeline v5.0 - Quality control PE
Cloud-Aerosole MT-MG Pre-Processing
Workflow 6: Alpha Diversity [Galaxy Training: 16S Microbial Analysis With Mothur]
Workflow 6: Alpha Diversity [16S Microbial Analysis With Mothur]
Training: 16S rRNA Sequencing With Mothur: Main Tutorial
Workflow7: Beta Diversity [Galaxy Training: 16S Microbial Analysis With Mothur]
Workflow 7 : Beta Diversity [16S Microbial Analysis With Mothur]
Workflow 1: Further Quality Control [16S Microbial Analysis With Mothur]
Amplicon Tutorial
Workflow 1: Quality Control [Galaxy Training: 16S Microbial Analysis With Mothur]
Workflow 2: Data Cleaning And Chimera Removal [16S Microbial Analysis With Mothur]
Workflow 2: Data Cleaning And Chimera Removal [Galaxy Training: 16S Microbial Analysis With Mothur]
Workflow 3: Classification [Galaxy Training: 16S Microbial Analysis With Mothur]
Workflow 4: Mock OTU Clustering [Galaxy Training: 16S Microbial Analysis With Mothur]
Workflow 5: OTU Clustering [16S Microbial Analysis With Mothur]
Workflow 5: OTU Clustering [Galaxy Training: 16S Microbial Analysis With Mothur]
Workflows
Galaxy Tool Suites