A

Cumulative number of **tool suites, tutorials, and workflows**

- Tool suites
- Tutorials
- Workflows
- More tutorials
- More workflows

Date

B

Suite users (last 5 years) on main servers

Included in
- Both
- Tutorials
- Workflows
- N/A

prokka — cutadapturecounts
kraken2
ncbi_blast_plus — snippy — fastp
fasttree — abricate — mothur
qualimap — ivar
bracken — dada2

Probabilities
- 75%
- 50%

Suite runs (last 5 years) on main servers

C

Suite runs (last 5 years) on main servers

Suite users (last 5 years) on main servers

EDAM operation

Left panel categories: Sequencing quality control, Sequence trimming, Formatting, Genome annotation, Antimicrobial resistance prediction, Visualisation, Variant calling, Sequence assembly validation, De-novo assembly, Sequence clustering, Genome assembly, Taxonomic classification, Read binning, Sequence analysis, Other, Sequence assembly

Right panel categories: Sequencing quality control, Visualisation, Genome annotation, Formatting, De-novo assembly, Sequence trimming, Sequence classification, Genome assembly, Taxonomic prediction, Antimicrobial resistance prediction, Sequence assembly validation, Variant calling, Sequence clustering, Sequence analysis, Sequence assembly, Other, Read binning