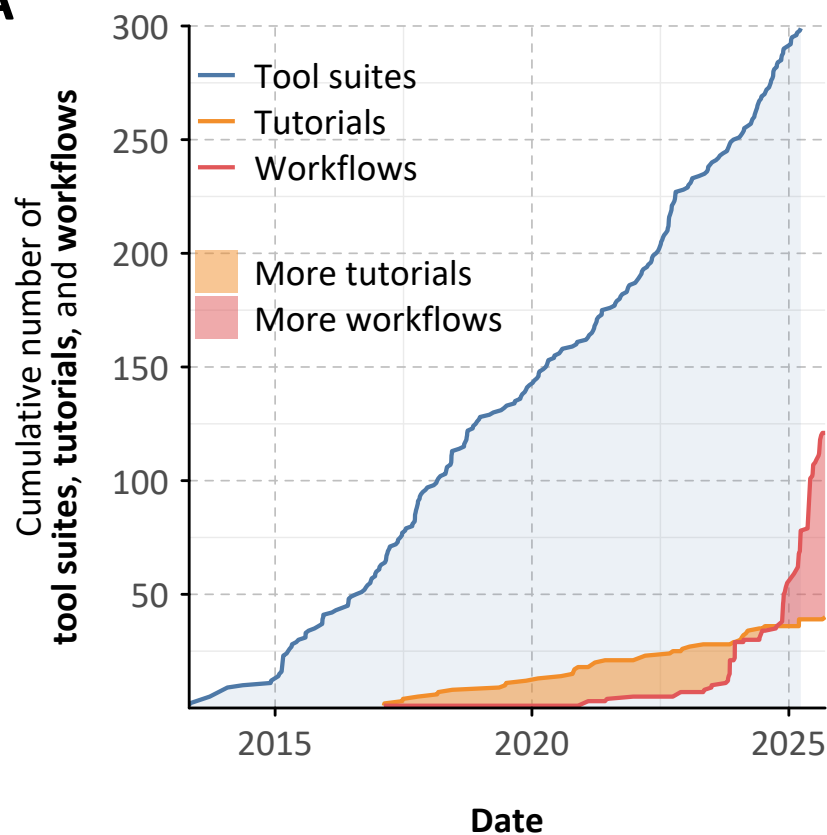
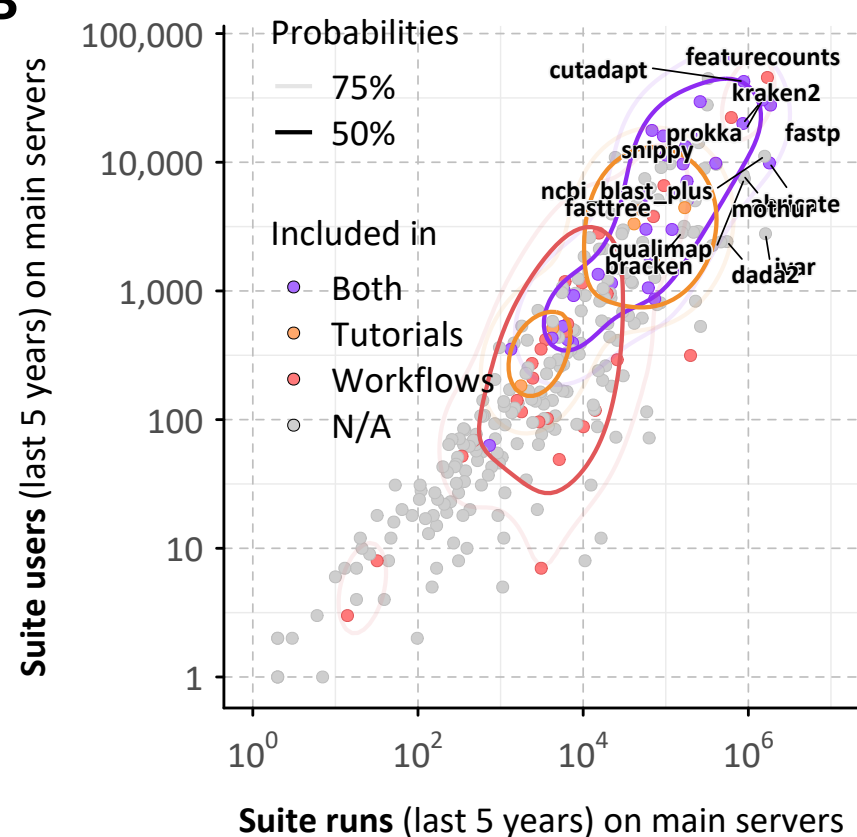
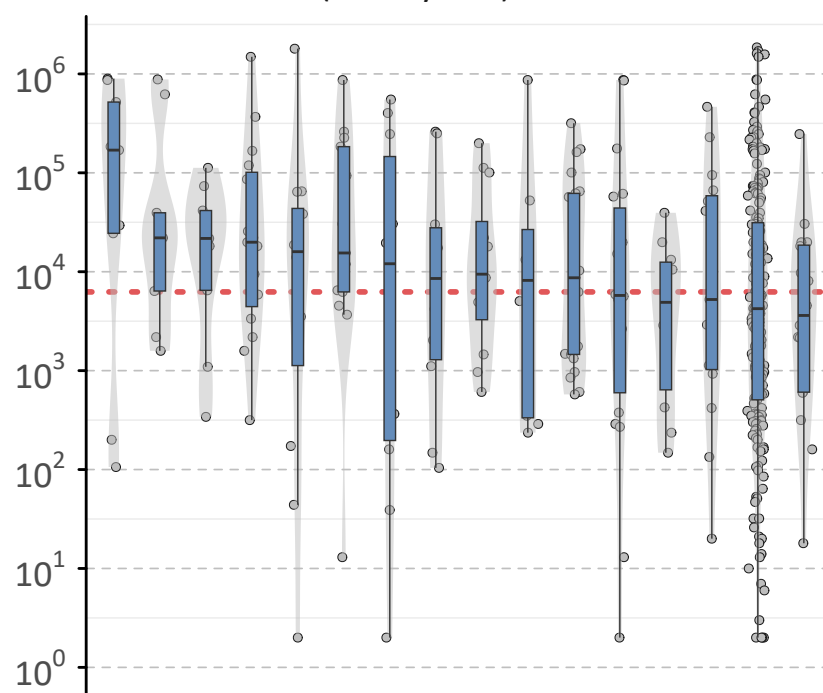
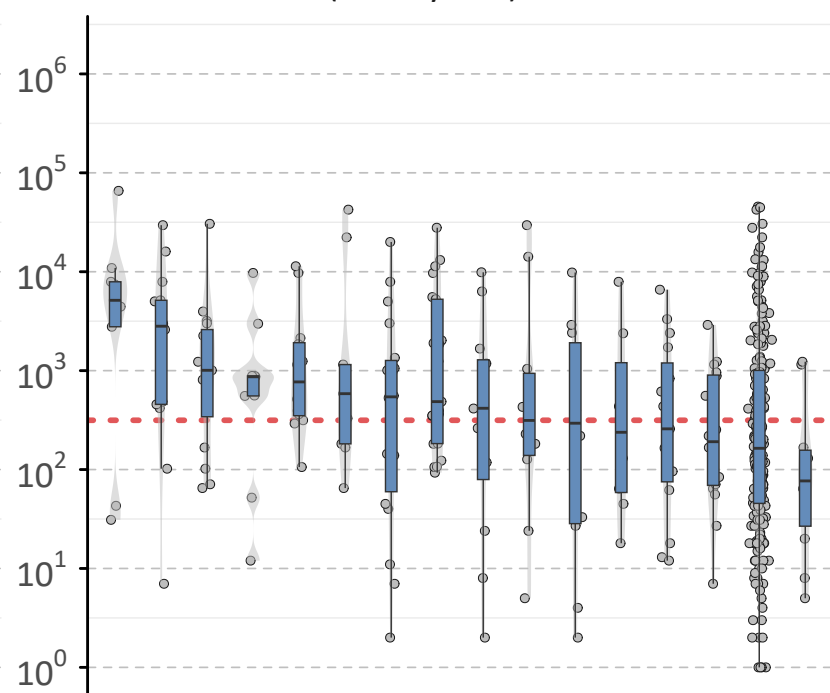


A**B****C****Suite runs (last 5 years) on main servers****Suite users (last 5 years) on main servers**

Sequencing quality control
 Sequence trimming
 Sequence formatting
 Genome annotation
 Antimicrobial resistance prediction
 Visualisation
 Variant calling
 De-novo validation
 De-novo assembly
 Sequence clustering
 Genome assembly
 Taxonomic classification
 Read binning
 Sequence analysis
 Other
 Sequence assembly

EDAM operation