

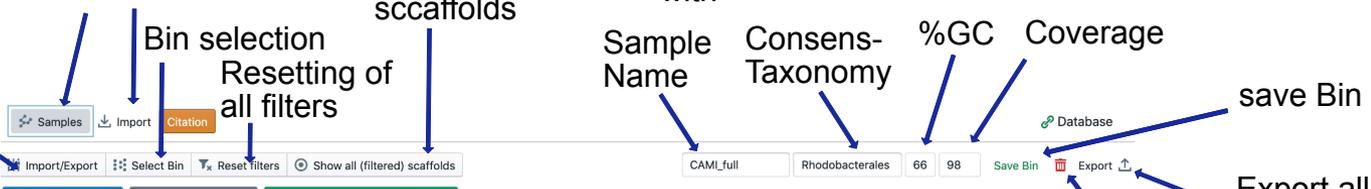
Sample selection and deletion

Switch between the current Sample tab and the Import Tab

Bin selection
Resetting of all filters

Show only binned scaffolds (default) or show all scaffolds

Bin name if its saved right now with



save Bin

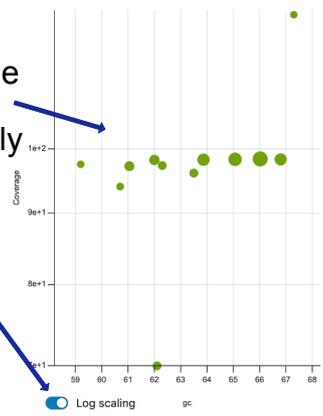
Export all Bins, optionally as fasta files

delete Bin

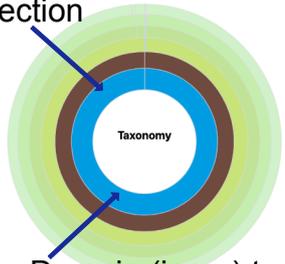
universal bacterial single copy genes

universal archaeal single copy genes

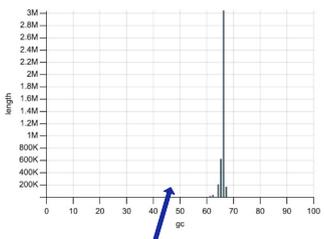
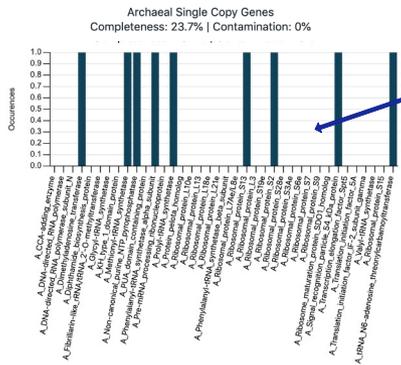
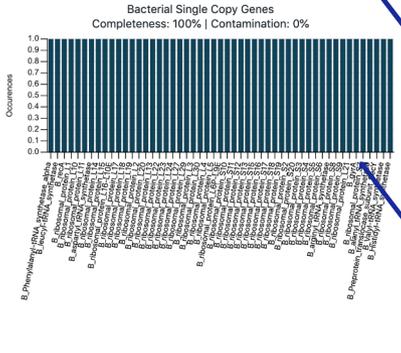
GC vs coverage plot, optionally log10 scaled



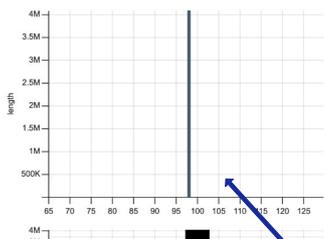
Consens-taxonomy distribution across selection



from Domain (inner) to species (outer) circle



GC-barplot, with the summed length of scaffolds on the Y-axis



Zooming window for the cov-barplot

Cov-barplot, with teh summed length of scaffolds on the Y-axis