

Metagenome

- +sequences
- +orfs
- +kmer_freqs
- +taxonomy_info

- +length_filter()

Dependencies

- +executables
- +databases

- +check_executables()
- +check_databases()
- +update_databases()

Bin

- +parent_metagenome
- +contig_names
- +status
- +marker_set
- +core_markers

- +split_taxonomy()
- +split_nucleotide()

Markers

- +sequences
- +orfs
- +taxid
- +marker_set
- +core_markers

- +get_marker_set()
- +get_core_markers()

Prodigal

- +sequences
- +orfs
- +n_orfs

- +get_orfs()