



TAXONBRIDGE

Build and analyse a custom taxonomy in four steps



STEP 1

Load

load_sample()



load_population()



download_ncbi()



load_taxonomies()



download_gbif()

STEP 2

Parse

get_lineages()



get_taxa()



get_validity()



get_status()



term_conversion()



dedupe()

STEP 3

Annotate

get_inconsistencies()



annotate()



fuzzy_search()

STEP 4

Visualise

prepare_rank_dist()



plot_mdb()



prepare_comparable_rank_dist()