Taxonbridge

Build and analyze a custom taxonomy in four steps

STEP 1

- load_sample()
- ⇕
- load_population()
- ⇕
- download_ncbi()
- ⇓
- load_taxonomies()
- ⇑
- download_gbif()

STEP 2

- get_lineages()
- ⇕
- get_taxa()
- ⇕
- get_validity()
- ⇕
- get_status()
- ⇕
- term_conversion()
- ⇕
- dedupe()

STEP 3

- get_inconsistencies()
- ⇓
- annotate()
- ⇑
- fuzzy_search()

STEP 4

- prepare_rank_dist()
- ⇓
- plot_mdb()
- ⇑
- prepare_comparable_rank_dist()