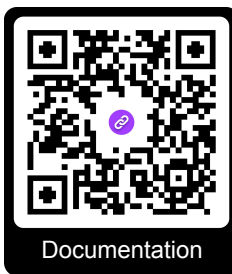




# TAXONBRIDGE

Build and analyse a custom taxonomy in four steps



## STEP 1

Load

load\_sample()



load\_population()



download\_ncbi()



load\_taxonomies()



download\_gbif()

## STEP 2

Filter

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()