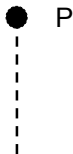


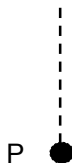
names

id
tax_id
name_txt
unique_name
name_class



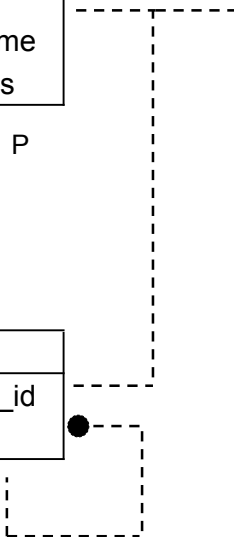
precalc_lineages

species_name
tax_id
lineage
num_terms



nodes

tax_id
parent_tax_id
rank



db_seq_ids

full_id
gi_num
alt_id
tax_id
alt_tax_id
species_name
seq_length
annotation