VegCore

< place | occurrence >

vegpath.org/VegCore/ERD.pdf

categories

source | event | occurrence | party | method | individual | community | specimen | place | taxon

relationships

..
id
HAS-A
base_class
id
referenced_class
inherits from record
IS-A
derived_class
id

validatable_place
id
coordinates
path

place_path
id
continent
country
state_province
county
municipality
ranks

coordinates
id
latitude_deg
longitude_deg

geovalidation
id
geovalid
lat_long_domain_valid
lat_long_in_ranks

place
id
parent
coordinates
path
locality

plot
id
name
area_m2
bounding_box

subplot
id
x_m
y_m

event
id
parent
name
date_range
place
method

event_participant
event
party
sort_order

project
id
name
info

method
id
parent
info

stratum
id
name
info

record
id
source
source_id_scope
source_record_id
info

source
id
parent
name
first_publisher
owner
info

relationship
id
record
related_record
info

party
id
info

organization
id
info

geological_context
id
name
info

community
id
name
info

place_observation
id
place
elevation_m
slope_incline_deg
slope_direction_deg_N
geological_context
community
observations

soil_observation
id
observations

taxon_observation
id
taxon_occurrence
collector
collector_number
voucher
growth_form
cultivated
traits

taxon_occurrence
id
current_determination
original_determination

taxon_presence
id
taxon_concept
traits

aggregate_observation
id
taxon_concept
traits

individual_observation
id
individual
code
traits

individual
id
tag

stem_observation
id
individual_observation
stem
traits

stem
id
individual

specimen
id
individual
code_in_individual
collection_event
orig_collection
barcode
accession_number
current_collection
owner_collection

collection
id
institution
name

specimen_observation
id
specimen
traits

specimen_holder_institution
specimen
institution
sort_order

taxon_determination
id
taxon_assertion
identified_by
fit_info

TNRS taxonomic scrubbing steps >

1a. verbatim v

taxon_string
string
parsed_taxon_assertion

taxon_assertion
id
string
taxon
cf_aff
annotations

3. parsed v

^ 2. concatenated

1b. verbatim >

taxon_name
id
unique_name
formal_name
taxon_name
author
common_name
rank

taxon_path
id
family
genus
specific_epithet
ranks

parsed_taxon_assertion
id
matched_taxon_concept
match_score
match_info

4. matched >

taxon_concept
id
according_to
parent
accepted_taxon_concept

< 5. accepted