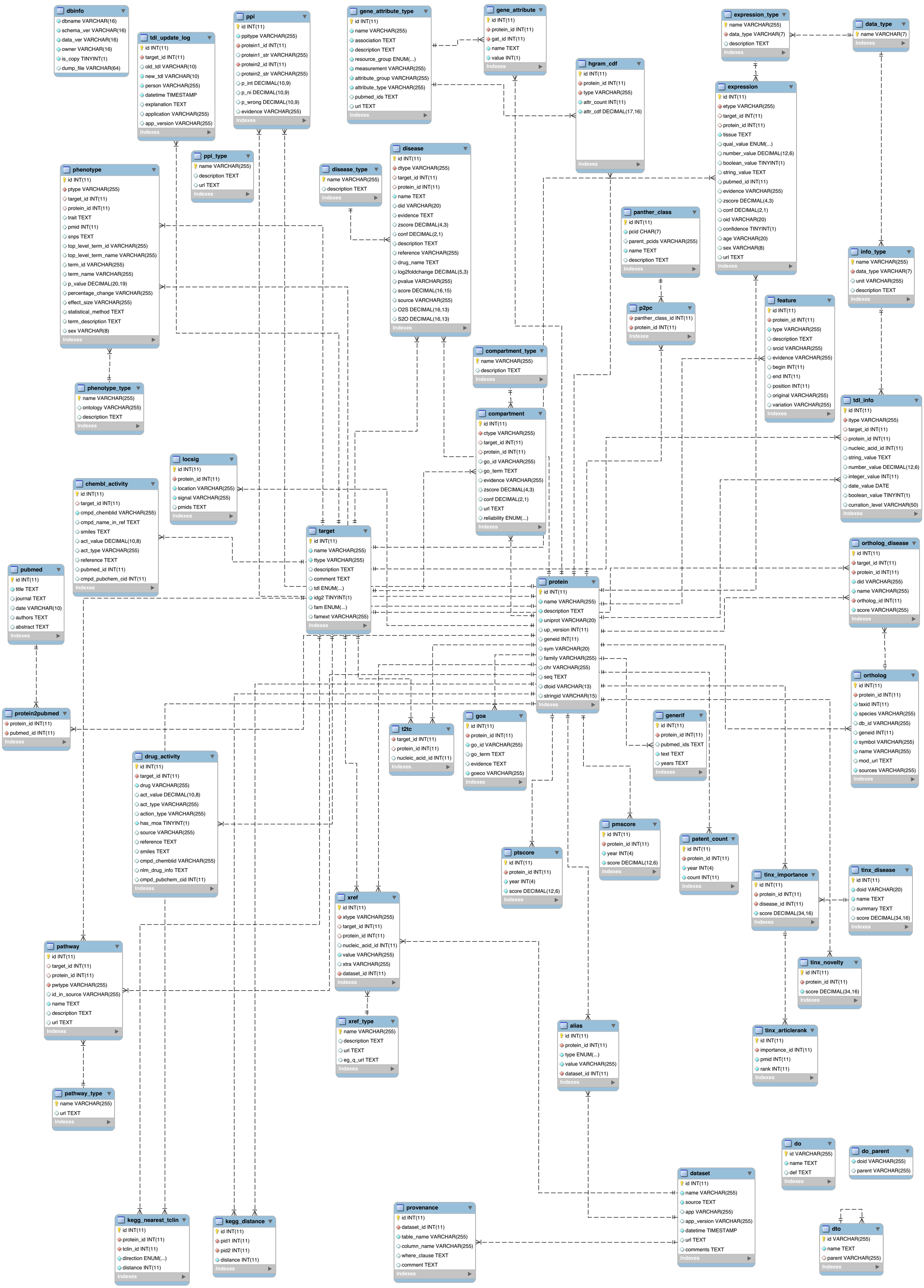
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data_ver VARCHAR(16)
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explanation TEXT
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app_version VARCHAR(255)
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protein2_str VARCHAR(255)
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p_ni DECIMAL(10,9)
p_wrong DECIMAL(10,9)
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description TEXT
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measurement VARCHAR(255)
attribute_group VARCHAR(255)
attribute_type VARCHAR(255)
pubmed_ids TEXT
url TEXT
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gene_attribute
id INT(11)
protein_id INT(11)
gat_id INT(11)
name TEXT
value INT(1)
Indexes
hgram_cdf
id INT(11)
protein_id INT(11)
type VARCHAR(255)
attr_count INT(11)
attr_cdf DECIMAL(17,16)
Indexes
expression_type
name VARCHAR(255)
data_type VARCHAR(7)
description TEXT
Indexes
data_type
name VARCHAR(7)
Indexes
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target_id INT(11)
protein_id INT(11)
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boolean_value TINYINT(1)
string_value TEXT
pubmed_id INT(11)
evidence VARCHAR(255)
zscore DECIMAL(4,3)
conf DECIMAL(2,1)
oid VARCHAR(20)
confidence TINYINT(1)
age VARCHAR(20)
sex VARCHAR(8)
url TEXT
Indexes
ppi_type
name VARCHAR(255)
description TEXT
url TEXT
Indexes
disease_type
name VARCHAR(255)
description TEXT
Indexes
disease
id INT(11)
dtype VARCHAR(255)
target_id INT(11)
protein_id INT(11)
name TEXT
did VARCHAR(20)
evidence TEXT
zscore DECIMAL(4,3)
conf DECIMAL(2,1)
description TEXT
reference VARCHAR(255)
drug_name TEXT
log2foldchange DECIMAL(5,3)
pvalue VARCHAR(255)
score DECIMAL(16,15)
source VARCHAR(255)
O2S DECIMAL(16,13)
S2O DECIMAL(16,13)
Indexes
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id INT(11)
pcid CHAR(7)
parent_pcids VARCHAR(255)
name TEXT
description TEXT
Indexes
p2pc
panther_class_id INT(11)
protein_id INT(11)
Indexes
feature
id INT(11)
protein_id INT(11)
type VARCHAR(255)
description TEXT
srcid VARCHAR(255)
evidence VARCHAR(255)
begin INT(11)
end INT(11)
position INT(11)
original VARCHAR(255)
variation VARCHAR(255)
Indexes
info_type
name VARCHAR(255)
data_type VARCHAR(7)
unit VARCHAR(255)
description TEXT
Indexes
phenotype
id INT(11)
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target_id INT(11)
protein_id INT(11)
trait TEXT
pmid INT(11)
snps TEXT
top_level_term_id VARCHAR(255)
top_level_term_name VARCHAR(255)
term_id VARCHAR(255)
term_name VARCHAR(255)
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percentage_change VARCHAR(255)
effect_size VARCHAR(255)
statistical_method TEXT
term_description TEXT
sex VARCHAR(8)
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phenotype_type
name VARCHAR(255)
ontology VARCHAR(255)
description TEXT
Indexes
compartment_type
name VARCHAR(255)
description TEXT
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id INT(11)
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protein_id INT(11)
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conf DECIMAL(2,1)
url TEXT
reliability ENUM(...)
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tdl_info
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target_id INT(11)
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nucleic_acid_id INT(11)
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number_value DECIMAL(12,6)
integer_value INT(11)
date_value DATE
boolean_value TINYINT(1)
curration_level VARCHAR(50)
Indexes
locsig
id INT(11)
protein_id INT(11)
location VARCHAR(255)
signal VARCHAR(255)
pmids TEXT
Indexes
chembl_activity
id INT(11)
target_id INT(11)
cmpd_chemblid VARCHAR(255)
cmpd_name_in_ref TEXT
smiles TEXT
act_value DECIMAL(10,8)
act_type VARCHAR(255)
reference TEXT
pubmed_id INT(11)
cmpd_pubchem_cid INT(11)
Indexes
target
id INT(11)
name VARCHAR(255)
ttype VARCHAR(255)
description TEXT
comment TEXT
tdl ENUM(...)
idg2 TINYINT(1)
fam ENUM(...)
famext VARCHAR(255)
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ortholog_disease
id INT(11)
target_id INT(11)
protein_id INT(11)
did VARCHAR(255)
name VARCHAR(255)
ortholog_id INT(11)
score VARCHAR(255)
Indexes
pubmed
id INT(11)
title TEXT
journal TEXT
date VARCHAR(10)
authors TEXT
abstract TEXT
Indexes
protein
id INT(11)
name VARCHAR(255)
description TEXT
uniprot VARCHAR(20)
up_version INT(11)
geneid INT(11)
sym VARCHAR(20)
family VARCHAR(255)
chr VARCHAR(255)
seq TEXT
dtoid VARCHAR(13)
stringid VARCHAR(15)
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id INT(11)
protein_id INT(11)
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species VARCHAR(255)
db_id VARCHAR(255)
geneid INT(11)
symbol VARCHAR(255)
name VARCHAR(255)
mod_url TEXT
sources VARCHAR(255)
Indexes
protein2pubmed
protein_id INT(11)
pubmed_id INT(11)
Indexes
t2tc
target_id INT(11)
protein_id INT(11)
nucleic_acid_id INT(11)
Indexes
goa
id INT(11)
protein_id INT(11)
go_id VARCHAR(255)
go_term TEXT
evidence TEXT
goeco VARCHAR(255)
Indexes
generif
id INT(11)
protein_id INT(11)
pubmed_ids TEXT
text TEXT
years TEXT
Indexes
drug_activity
id INT(11)
target_id INT(11)
drug VARCHAR(255)
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act_type VARCHAR(255)
action_type VARCHAR(255)
has_moa TINYINT(1)
source VARCHAR(255)
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smiles TEXT
cmpd_chemblid VARCHAR(255)
nlm_drug_info TEXT
cmpd_pubchem_cid INT(11)
Indexes
pmscore
id INT(11)
protein_id INT(11)
year INT(4)
score DECIMAL(12,6)
Indexes
patent_count
id INT(11)
protein_id INT(11)
year INT(4)
count INT(11)
Indexes
ptscore
id INT(11)
protein_id INT(11)
year INT(4)
score DECIMAL(12,6)
Indexes
tinx_importance
id INT(11)
protein_id INT(11)
disease_id INT(11)
score DECIMAL(34,16)
Indexes
tinx_disease
id INT(11)
doid VARCHAR(20)
name TEXT
summary TEXT
score DECIMAL(34,16)
Indexes
xref
id INT(11)
xtype VARCHAR(255)
target_id INT(11)
protein_id INT(11)
nucleic_acid_id INT(11)
value VARCHAR(255)
xtra VARCHAR(255)
dataset_id INT(11)
Indexes
tinx_novelty
id INT(11)
protein_id INT(11)
score DECIMAL(34,16)
Indexes
pathway
id INT(11)
target_id INT(11)
protein_id INT(11)
pwtype VARCHAR(255)
id_in_source VARCHAR(255)
name TEXT
description TEXT
url TEXT
Indexes
xref_type
name VARCHAR(255)
description TEXT
url TEXT
eg_q_url TEXT
Indexes
alias
id INT(11)
protein_id INT(11)
type ENUM(...)
value VARCHAR(255)
dataset_id INT(11)
Indexes
tinx_articlerank
id INT(11)
importance_id INT(11)
pmid INT(11)
rank INT(11)
Indexes
pathway_type
name VARCHAR(255)
url TEXT
Indexes
do
id VARCHAR(255)
name TEXT
def TEXT
Indexes
do_parent
doid VARCHAR(255)
parent VARCHAR(255)
dataset
id INT(11)
name VARCHAR(255)
source TEXT
app VARCHAR(255)
app_version VARCHAR(255)
datetime TIMESTAMP
url TEXT
comments TEXT
Indexes
kegg_nearest_tclin
id INT(11)
protein_id INT(11)
tclin_id INT(11)
direction ENUM(...)
distance INT(11)
Indexes
kegg_distance
id INT(11)
pid1 INT(11)
pid2 INT(11)
distance INT(11)
Indexes
provenance
id INT(11)
dataset_id INT(11)
table_name VARCHAR(255)
column_name VARCHAR(255)
where_clause TEXT
comment TEXT
Indexes
dto
id VARCHAR(255)
name TEXT
parent VARCHAR(255)
Indexes