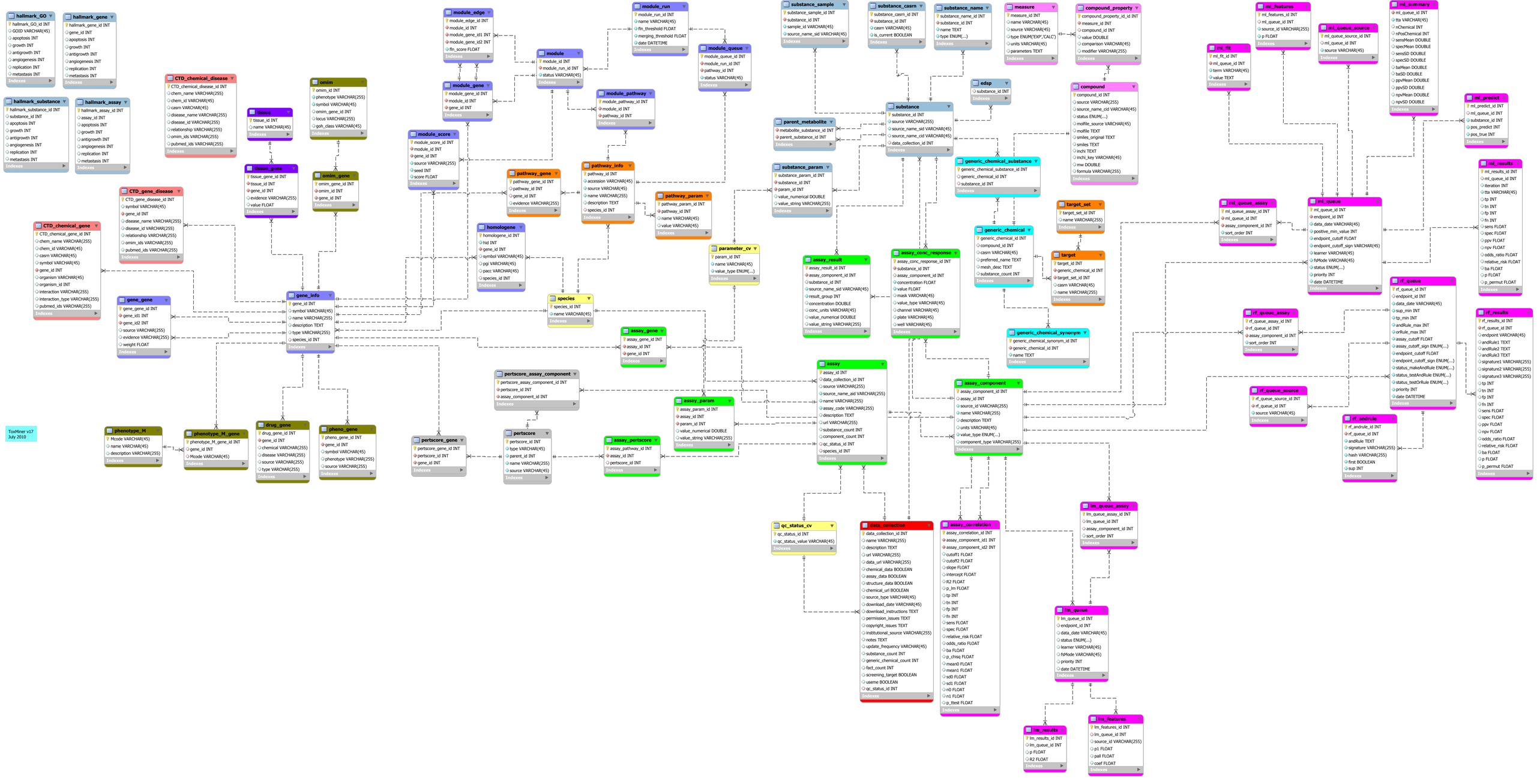
hallmark_GO
GOID VARCHAR(45)
apoptosis INT
growth INT
antigrowth INT
angiogenesis INT
replication INT
metastasis INT
hallmark_gene
hallmark_gene_id INT
gene_id INT
apoptosis INT
growth INT
antigrowth INT
angiogenesis INT
replication INT
metastasis INT
hallmark_substance
hallmark_substance_id INT
substance_id INT
apoptosis INT
growth INT
antigrowth INT
angiogenesis INT
replication INT
metastasis INT
hallmark_assay
hallmark_assay_id INT
assay_id INT
apoptosis INT
growth INT
antigrowth INT
angiogenesis INT
replication INT
metastasis INT
CTD_chemical_disease
CTD_chemical_disease_id INT
chem_name VARCHAR(255)
chem_id VARCHAR(45)
casrn VARCHAR(45)
disease_name VARCHAR(255)
disease_id VARCHAR(255)
relationship VARCHAR(255)
omim_ids VARCHAR(255)
pubmed_ids VARCHAR(255)
omim
omim_id INT
phenotype VARCHAR(255)
symbol VARCHAR(45)
omim_gene_id INT
locus VARCHAR(255)
goh_class VARCHAR(45)
tissue
tissue_id INT
name VARCHAR(45)
tissue_gene
tissue_gene_id INT
tissue_id INT
gene_id INT
evidence VARCHAR(255)
value FLOAT
omim_gene
omim_gene_id INT
omim_id INT
gene_id INT
CTD_gene_disease
CTD_gene_disease_id INT
symbol VARCHAR(45)
gene_id INT
disease_name VARCHAR(255)
disease_id VARCHAR(255)
relationship VARCHAR(255)
omim_ids VARCHAR(255)
pubmed_ids VARCHAR(255)
CTD_chemical_gene
CTD_chemical_gene_id INT
chem_name VARCHAR(255)
chem_id VARCHAR(45)
casrn VARCHAR(45)
symbol VARCHAR(45)
gene_id INT
organism VARCHAR(45)
organism_id INT
interaction VARCHAR(255)
interaction_type VARCHAR(255)
pubmed_ids VARCHAR(255)
gene_gene
gene_gene_id INT
gene_id1 INT
gene_id2 INT
source VARCHAR(255)
evidence VARCHAR(255)
weight FLOAT
gene_info
gene_id INT
symbol VARCHAR(45)
name VARCHAR(255)
description TEXT
type VARCHAR(255)
species_id INT
phenotype_M
Mcode VARCHAR(45)
name VARCHAR(45)
description VARCHAR(255)
phenotype_M_gene
phenotype_M_gene_id INT
gene_id INT
Mcode VARCHAR(45)
drug_gene
drug_gene_id INT
gene_id INT
chemical VARCHAR(255)
disease VARCHAR(255)
source VARCHAR(255)
type VARCHAR(255)
pheno_gene
pheno_gene_id INT
gene_id INT
symbol VARCHAR(45)
phenotype VARCHAR(255)
source VARCHAR(255)
module_edge
module_edge_id INT
module_id INT
module_gene_id1 INT
module_gene_id2 INT
fln_score FLOAT
module_gene
module_gene_id INT
module_id INT
gene_id INT
module_score
module_score_id INT
module_id INT
gene_id INT
source VARCHAR(255)
seed INT
score FLOAT
homologene
homologene_id INT
hid INT
gene_id INT
symbol VARCHAR(45)
pgi VARCHAR(45)
pacc VARCHAR(45)
species_id INT
module
module_id INT
module_run_id INT
status VARCHAR(45)
module_run
module_run_id INT
name VARCHAR(45)
fln_threshold FLOAT
merging_threshold FLOAT
date DATETIME
module_pathway
module_pathway_id INT
module_id INT
pathway_id INT
module_queue
module_queue_id INT
module_run_id INT
pathway_id INT
status VARCHAR(45)
pathway_gene
pathway_gene_id INT
pathway_id INT
gene_id INT
evidence VARCHAR(255)
pathway_info
pathway_id INT
accession VARCHAR(45)
source VARCHAR(45)
name VARCHAR(255)
description TEXT
species_id INT
pathway_param
pathway_param_id INT
pathway_id INT
name VARCHAR(45)
value VARCHAR(45)
parameter_cv
param_id INT
name VARCHAR(45)
value_type ENUM(...)
species
species_id INT
name VARCHAR(45)
assay_gene
assay_gene_id INT
assay_id INT
gene_id INT
pertscore_assay_component
pertscore_assay_component_id INT
pertscore_id INT
assay_component_id INT
pertscore_gene
pertscore_gene_id INT
pertscore_id INT
gene_id INT
pertscore
pertscore_id INT
type VARCHAR(45)
parent_id INT
name VARCHAR(255)
source VARCHAR(45)
assay_param
assay_param_id INT
assay_id INT
param_id INT
value_numerical DOUBLE
value_string VARCHAR(255)
assay_pertscore
assay_pathway_id INT
assay_id INT
pertscore_id INT
substance_sample
substance_sample_id INT
substance_id INT
sample_id VARCHAR(45)
source_name_sid VARCHAR(45)
substance_casrn
substance_casrn_id INT
substance_id INT
casrn VARCHAR(45)
is_current BOOLEAN
substance_name
substance_name_id INT
substance_id INT
name TEXT
type ENUM(...)
measure
measure_id INT
name VARCHAR(45)
source VARCHAR(45)
type ENUM('EXP','CALC')
units VARCHAR(45)
parameters TEXT
compound_property
compound_property_id_id INT
measure_id INT
compound_id INT
value DOUBLE
comparison VARCHAR(45)
modifier VARCHAR(255)
edsp
substance_id INT
compound
compound_id INT
source VARCHAR(255)
source_name_cid VARCHAR(45)
status ENUM(...)
molfile_source VARCHAR(45)
molfile TEXT
smiles_original TEXT
smiles TEXT
inchi TEXT
inchi_key VARCHAR(45)
mw DOUBLE
formula VARCHAR(255)
substance
substance_id INT
source VARCHAR(255)
source_name_sid VARCHAR(45)
source_name_cid VARCHAR(45)
data_collection_id INT
parent_metabolite
metabolite_substance_id INT
parent_substance_id INT
substance_param
substance_param_id INT
substance_id INT
param_id INT
value_numerical DOUBLE
value_string VARCHAR(255)
generic_chemical_substance
generic_chemical_substance_id INT
generic_chemical_id INT
substance_id INT
generic_chemical
generic_chemical_id INT
compound_id INT
casrn VARCHAR(45)
preferred_name TEXT
mesh_desc TEXT
substance_count INT
target_set
target_set_id INT
name VARCHAR(255)
target
target_id INT
generic_chemical_id INT
target_set_id INT
casrn VARCHAR(45)
name VARCHAR(255)
generic_chemical_synonym
generic_chemical_synonym_id INT
generic_chemical_id INT
name TEXT
assay_result
assay_result_id INT
assay_component_id INT
substance_id INT
source_name_sid VARCHAR(45)
result_group INT
concentration DOUBLE
conc_units VARCHAR(45)
value_numerical DOUBLE
value_string VARCHAR(255)
assay_conc_response
assay_conc_response_id INT
substance_id INT
assay_component_id INT
concentration FLOAT
value FLOAT
mask VARCHAR(45)
value_type VARCHAR(45)
channel VARCHAR(45)
plate VARCHAR(45)
well VARCHAR(45)
assay
assay_id INT
data_collection_id INT
source VARCHAR(255)
source_name_sid VARCHAR(255)
name VARCHAR(255)
assay_code VARCHAR(255)
description TEXT
url VARCHAR(255)
substance_count INT
component_count INT
qc_status_id INT
species_id INT
assay_component
assay_component_id INT
assay_id INT
source_id VARCHAR(255)
name VARCHAR(255)
description TEXT
units VARCHAR(45)
value_type ENUM(...)
component_type VARCHAR(255)
ml_features
ml_features_id INT
ml_queue_id INT
source_id VARCHAR(255)
p FLOAT
ml_queue_source
ml_queue_source_id INT
ml_queue_id INT
source VARCHAR(45)
ml_summary
ml_queue_id INT
tta VARCHAR(45)
nChemical INT
nProsChemical INT
sensMean DOUBLE
specMean DOUBLE
sensSD DOUBLE
specSD DOUBLE
baMean DOUBLE
baSD DOUBLE
ppvMean DOUBLE
ppvSD DOUBLE
npvMean DOUBLE
npvSD DOUBLE
ml_fit
ml_fit_id INT
ml_queue_id INT
term VARCHAR(45)
value TEXT
ml_predict
ml_predict_id INT
ml_queue_id INT
substance_id INT
pos_predict INT
pos_true INT
ml_results
ml_results_id INT
ml_queue_id INT
iteration INT
tta VARCHAR(45)
tp INT
tn INT
fp INT
fn INT
sens FLOAT
spec FLOAT
ppv FLOAT
npv FLOAT
odds_ratio FLOAT
relative_risk FLOAT
ba FLOAT
p FLOAT
p_permut FLOAT
ml_queue_assay
ml_queue_assay_id INT
ml_queue_id INT
assay_component_id INT
sort_order INT
ml_queue
ml_queue_id INT
endpoint_id INT
data_date VARCHAR(45)
positive_min_value INT
endpoint_cutoff FLOAT
endpoint_cutoff_sign VARCHAR(45)
learner VARCHAR(45)
fsMode VARCHAR(45)
status ENUM(...)
priority INT
date DATETIME
rf_queue
rf_queue_id INT
endpoint_id INT
data_date VARCHAR(45)
sup_min INT
tp_min INT
andRule_max INT
orRule_max INT
assay_cutoff FLOAT
assay_cutoff_sign ENUM(...)
endpoint_cutoff FLOAT
endpoint_cutoff_sign ENUM(...)
status_makeAndRule ENUM(...)
status_testAndRule ENUM(...)
status_testOrRule ENUM(...)
priority INT
date DATETIME
rf_queue_assay
rf_queue_assay_id INT
rf_queue_id INT
assay_component_id INT
sort_order INT
rf_results
rf_results_id INT
rf_queue_id INT
endpoint VARCHAR(45)
andRule1 TEXT
andRule2 TEXT
andRule3 TEXT
signature1 VARCHAR(255)
signature2 VARCHAR(255)
signature3 VARCHAR(255)
tp INT
tn INT
fp INT
fn INT
sens FLOAT
spec FLOAT
ppv FLOAT
npv FLOAT
odds_ratio FLOAT
relative_risk FLOAT
ba FLOAT
p FLOAT
p_permut FLOAT
rf_queue_source
rf_queue_source_id INT
rf_queue_id INT
source VARCHAR(45)
rf_andrule
rf_andrule_id INT
rf_queue_id INT
andRule TEXT
signature VARCHAR(255)
hash VARCHAR(255)
first BOOLEAN
sup INT
qc_status_cv
qc_status_id INT
qc_status_value VARCHAR(45)
data_collection
data_collection_id INT
name VARCHAR(255)
description TEXT
url VARCHAR(255)
data_url VARCHAR(255)
chemical_data BOOLEAN
assay_data BOOLEAN
structure_data BOOLEAN
chemical_url BOOLEAN
source_type VARCHAR(45)
download_date VARCHAR(45)
download_instructions TEXT
permission_issues TEXT
copyright_issues TEXT
institutional_source VARCHAR(255)
notes TEXT
update_frequency VARCHAR(45)
substance_count INT
generic_chemical_count INT
fact_count INT
screening_target BOOLEAN
useme BOOLEAN
qc_status_id INT
assay_correlation
assay_correlation_id INT
assay_component_id1 INT
assay_component_id2 INT
cutoff1 FLOAT
cutoff2 FLOAT
slope FLOAT
intercept FLOAT
R2 FLOAT
p_lm FLOAT
tp INT
tn INT
fp INT
fn INT
sens FLOAT
spec FLOAT
relative_risk FLOAT
odds_ratio FLOAT
ba FLOAT
p_chisq FLOAT
mean0 FLOAT
mean1 FLOAT
sd0 FLOAT
sd1 FLOAT
n0 FLOAT
n1 FLOAT
p_ttest FLOAT
lm_queue_assay
lm_queue_assay_id INT
lm_queue_id INT
assay_component_id INT
sort_order INT
lm_queue
lm_queue_id INT
endpoint_id INT
data_date VARCHAR(45)
status ENUM(...)
learner VARCHAR(45)
fsMode VARCHAR(45)
priority INT
date DATETIME
lm_results
lm_results_id INT
lm_queue_id INT
p FLOAT
R2 FLOAT
lm_features
lm_features_id INT
lm_queue_id INT
source_id VARCHAR(255)
p1 FLOAT
pall FLOAT
coef FLOAT
ToxMiner v17
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