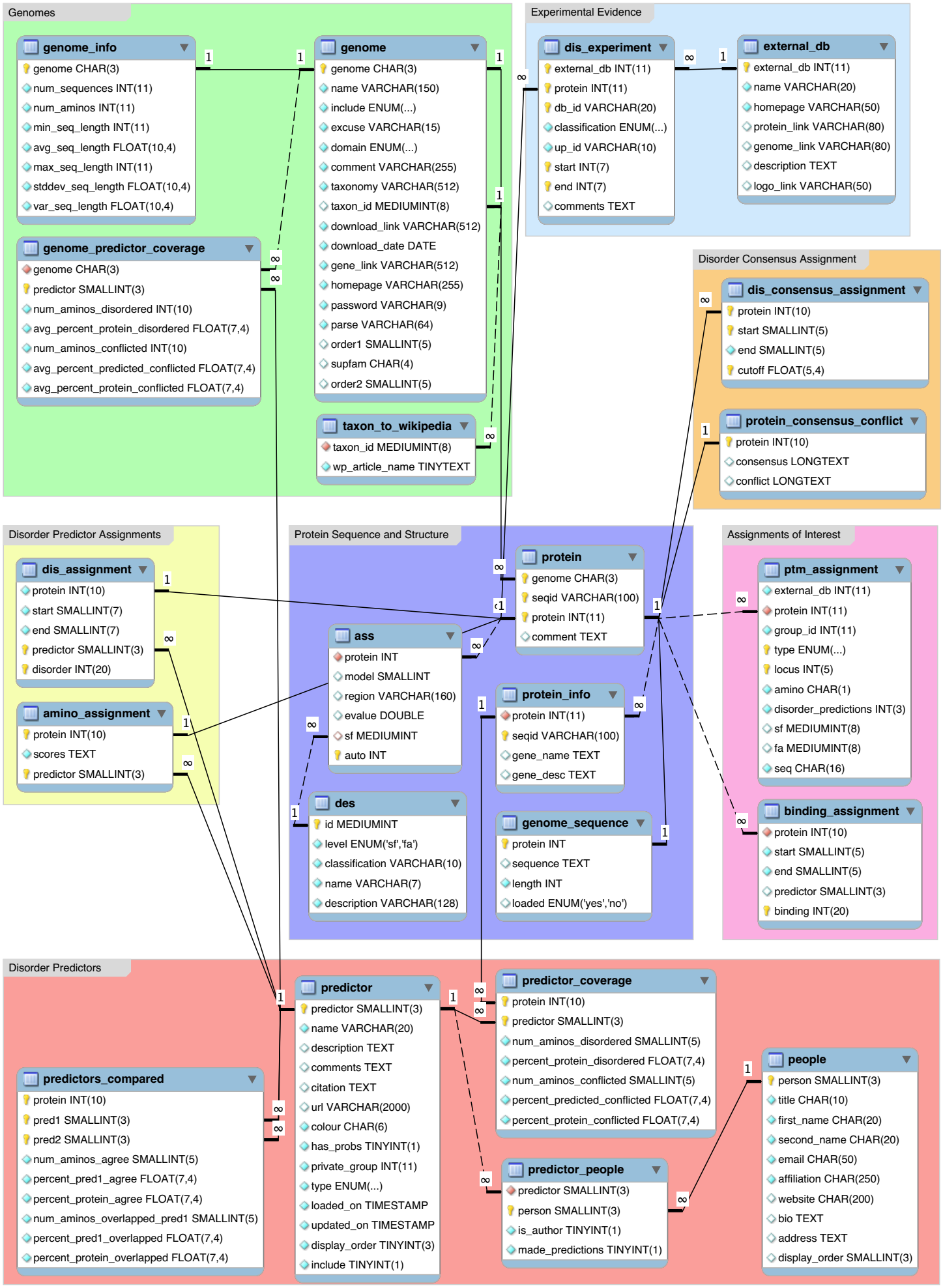
Genomes
genome_info
genome CHAR(3)
num_sequences INT(11)
num_aminos INT(11)
min_seq_length INT(11)
avg_seq_length FLOAT(10,4)
max_seq_length INT(11)
stddev_seq_length FLOAT(10,4)
var_seq_length FLOAT(10,4)
genome
genome CHAR(3)
name VARCHAR(150)
include ENUM(...)
excuse VARCHAR(15)
domain ENUM(...)
comment VARCHAR(255)
taxonomy VARCHAR(512)
taxon_id MEDIUMINT(8)
download_link VARCHAR(512)
download_date DATE
gene_link VARCHAR(512)
homepage VARCHAR(255)
password VARCHAR(9)
parse VARCHAR(64)
order1 SMALLINT(5)
supfam CHAR(4)
order2 SMALLINT(5)
genome_predictor_coverage
genome CHAR(3)
predictor SMALLINT(3)
num_aminos_disordered INT(10)
avg_percent_protein_disordered FLOAT(7,4)
num_aminos_conflicted INT(10)
avg_percent_predicted_conflicted FLOAT(7,4)
avg_percent_protein_conflicted FLOAT(7,4)
taxon_to_wikipedia
taxon_id MEDIUMINT(8)
wp_article_name TINYTEXT
Experimental Evidence
dis_experiment
external_db INT(11)
protein INT(11)
db_id VARCHAR(20)
classification ENUM(...)
up_id VARCHAR(10)
start INT(7)
end INT(7)
comments TEXT
external_db
external_db INT(11)
name VARCHAR(20)
homepage VARCHAR(50)
protein_link VARCHAR(80)
genome_link VARCHAR(80)
description TEXT
logo_link VARCHAR(50)
Disorder Consensus Assignment
dis_consensus_assignment
protein INT(10)
start SMALLINT(5)
end SMALLINT(5)
cutoff FLOAT(5,4)
protein_consensus_conflict
protein INT(10)
consensus LONGTEXT
conflict LONGTEXT
Disorder Predictor Assignments
dis_assignment
protein INT(10)
start SMALLINT(7)
end SMALLINT(7)
predictor SMALLINT(3)
disorder INT(20)
amino_assignment
protein INT(10)
scores TEXT
predictor SMALLINT(3)
Protein Sequence and Structure
protein
genome CHAR(3)
seqid VARCHAR(100)
protein INT(11)
comment TEXT
ass
protein INT
model SMALLINT
region VARCHAR(160)
evalue DOUBLE
sf MEDIUMINT
auto INT
protein_info
protein INT(11)
seqid VARCHAR(100)
gene_name TEXT
gene_desc TEXT
des
id MEDIUMINT
level ENUM('sf','fa')
classification VARCHAR(10)
name VARCHAR(7)
description VARCHAR(128)
genome_sequence
protein INT
sequence TEXT
length INT
loaded ENUM('yes','no')
Assignments of Interest
ptm_assignment
external_db INT(11)
protein INT(11)
group_id INT(11)
type ENUM(...)
locus INT(5)
amino CHAR(1)
disorder_predictions INT(3)
sf MEDIUMINT(8)
fa MEDIUMINT(8)
seq CHAR(16)
binding_assignment
protein INT(10)
start SMALLINT(5)
end SMALLINT(5)
predictor SMALLINT(3)
binding INT(20)
Disorder Predictors
predictors_compared
protein INT(10)
pred1 SMALLINT(3)
pred2 SMALLINT(3)
num_aminos_agree SMALLINT(5)
percent_pred1_agree FLOAT(7,4)
percent_protein_agree FLOAT(7,4)
num_aminos_overlapped_pred1 SMALLINT(5)
percent_pred1_overlapped FLOAT(7,4)
percent_protein_overlapped FLOAT(7,4)
predictor
predictor SMALLINT(3)
name VARCHAR(20)
description TEXT
comments TEXT
citation TEXT
url VARCHAR(2000)
colour CHAR(6)
has_probs TINYINT(1)
private_group INT(11)
type ENUM(...)
loaded_on TIMESTAMP
updated_on TIMESTAMP
display_order TINYINT(3)
include TINYINT(1)
predictor_coverage
protein INT(10)
predictor SMALLINT(3)
num_aminos_disordered SMALLINT(5)
percent_protein_disordered FLOAT(7,4)
num_aminos_conflicted SMALLINT(5)
percent_predicted_conflicted FLOAT(7,4)
percent_protein_conflicted FLOAT(7,4)
predictor_people
predictor SMALLINT(3)
person SMALLINT(3)
is_author TINYINT(1)
made_predictions TINYINT(1)
people
person SMALLINT(3)
title CHAR(10)
first_name CHAR(20)
second_name CHAR(20)
email CHAR(50)
affiliation CHAR(250)
website CHAR(200)
bio TEXT
address TEXT
display_order SMALLINT(3)