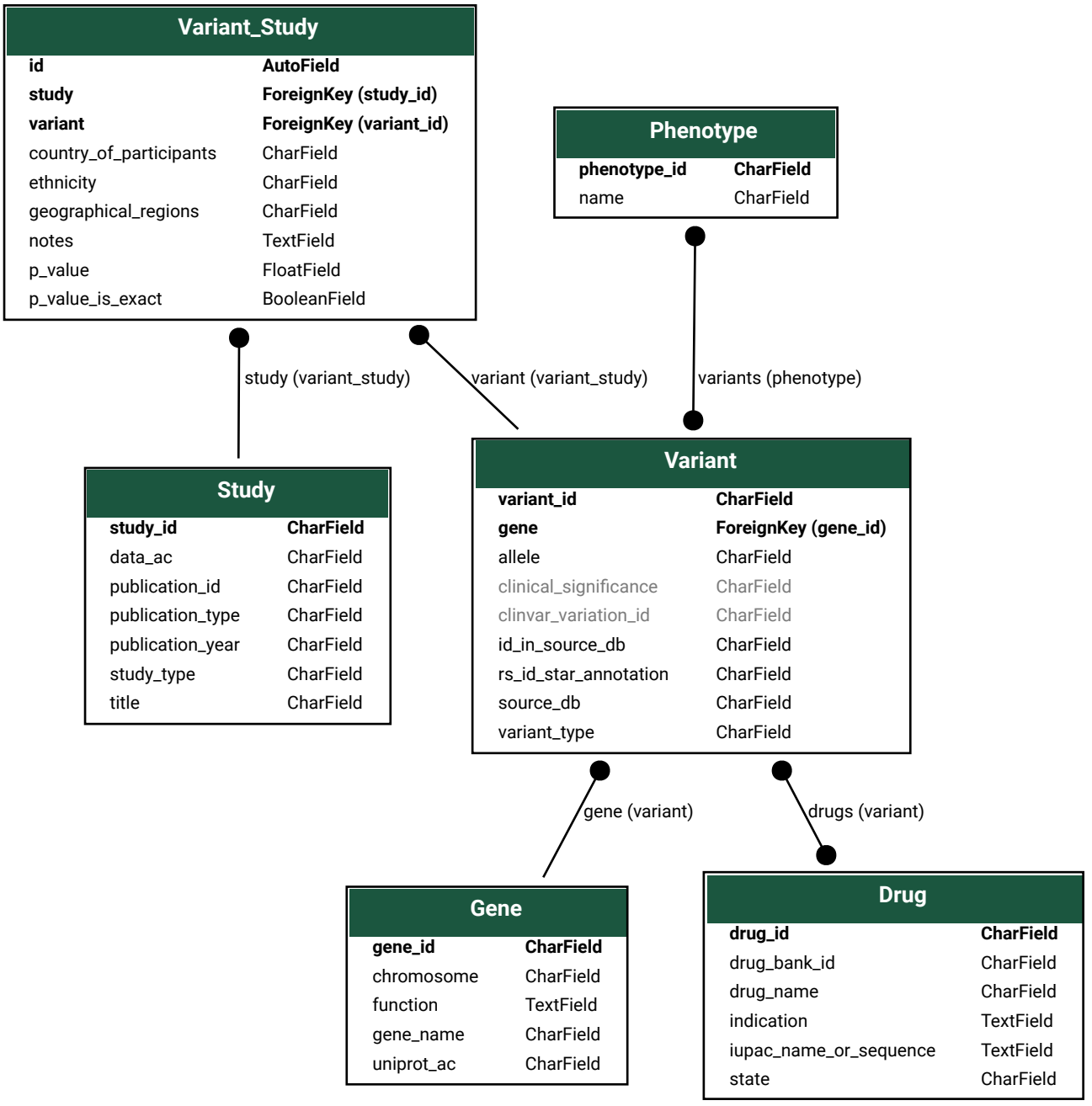
Variant_Study
id AutoField
study ForeignKey (study_id)
variant ForeignKey (variant_id)
country_of_participants CharField
ethnicity CharField
geographical_regions CharField
notes TextField
p_value FloatField
p_value_is_exact BooleanField
Phenotype
phenotype_id CharField
name CharField
study (variant_study)
variant (variant_study)
variants (phenotype)
Study
study_id CharField
data_ac CharField
publication_id CharField
publication_type CharField
publication_year CharField
study_type CharField
title CharField
Variant
variant_id CharField
gene ForeignKey (gene_id)
allele CharField
clinical_significance CharField
clinvar_variation_id CharField
id_in_source_db CharField
rs_id_star_annotation CharField
source_db CharField
variant_type CharField
gene (variant)
drugs (variant)
Gene
gene_id CharField
chromosome CharField
function TextField
gene_name CharField
uniprot_ac CharField
Drug
drug_id CharField
drug_bank_id CharField
drug_name CharField
indication TextField
iupac_name_or_sequence TextField
state CharField