	pgs_pubs		1,577
-	pg_publicationid	TEXT	NN
	release_date	TEXT	
	authors	TEXT	
	num_pgs_developed	INT	
	num_pgs_evaluated	INT	
	first_author	TEXT	
	title	TEXT	
	journal_name	TEXT	
	publication_date	TEXT	
	pmd_doi	TEXT	
ne 🖪	pmd_id	TEXT	

	pgs_samplesets		
m 🍇	pg_sampleset_id	TEXT	NN
m 2	pg_score_id	TEXT	
	num_individuals	INT	
	num_cases	TEXT	
	num_controls	TEXT	
	percent_male_participants	TEXT	
	sample_age	TEXT	
	broad_ancestry	TEXT	
	category_ancestry	TEXT	
	country_recruitment	TEXT	
	additional_ancestry	TEXT	
	description_phenotype	TEXT	
	followup_time	TEXT	
	gwas_catalogstudy_id	TEXT	
	source_pmdid	TEXT	
	source_doi	TEXT	
	cohort	TEXT	
	additional_information	TEXT	

	pgs_snp	ometa	
	pgs_score_id	TEXT	NN
	rsid	NUMERIC	
	gene	TEXT	
	func	TEXT	
	rank	REAL	
	snpcoord	TEXT	
	trait	TEXT	
	rank snpcoord	REAL TEXT	

		pgs_evalmetrics		
	pro de	pg_performancemetric_id	TEXT	NN
		evaluated_pgs_score_id	TEXT	
1	100 E	pg_sampleset_id	TEXT	
1	100 E	pg_publicationid	TEXT	
		reported_trait	TEXT	
		covariates	TEXT	
		pgsperformanceinfo	TEXT	
	per 🖬	pmd_id	TEXT	
		publication_doi	TEXT	
		hazard_ratio	TEXT	
		odds_ratio	TEXT	
		beta	TEXT	
		auc	TEXT	
		c_index	TEXT	
		other_metric	TEXT	

	pgs_meta		
100	pg_scoreid	TEXT	NN
	pgs_name	TEXT	
	reported_trait	TEXT	
	mapped_trait	TEXT	
m=1	trait_id	TEXT	
	pgs_development_method	TEXT	
	pgs_development_details	TEXT	
	original_genome_build	TEXT	
	num_variants	INT	
	number_of_interaction_terms	INT	
	type_of_variant_weight	TEXT	
In a	pg_publicationid	TEXT	
m=1	pmd_id	TEXT	
	ancestry_gwas	TEXT	
	ancestry_training	TEXT	
	ancestry_evaluation	TEXT	
	ftp_link	TEXT	
m-1	pgs_snpmeta_pgs_score_id	TEXT	NN

	pgs_traits	
ur 🍱	trait_id	TEXT NN
	mapped_ontology_trait	TEXT
	ontology_trait_description	TEXT
	ontology_url	TEXT