lipd domain model

Domain

clan_acc *string*
clan_id *string*
description *text*
name *string*
pfam_acc *string*
pfam_id *string*

Evidence

name *string*∗

Family

name *string*∗

Taxon

NCBIid *integer*
NCBIlineage *text*
taxon_name *string*∗

Protein

name *string*
protein_accession *string*∗
protein_sequence *text*

ProteinHasDomain

end_position *integer*
evalue *float*
start_position *integer*

ProteinsGoterm

Goterm

go_name *string*
go_term *string*∗