

GeDiPNet Cluster 387 -- Pairs Within Cluster

Similarity: Jaccard index on shared curated genes (disease_gdp). Significance: upper-tail hypergeometric test, Benjamini-Hochberg FDR-corrected across all tested pairs. See scripts/build_related.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-07 17:17:46. · Rows: 1 · Generated: 2026-10-07 21:03:04

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Alpha-methylacyl-coa racemase deficiency	Congenital bile acid synthesis defect	0.14286	1	3.897e-4	8.637e-4