

GeDiPNet Cluster 646 -- 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: 2026-10-21 12:15:07. · Rows: 7 · Generated: 2026-10-21 08:34:55

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hyper-immunoglobulin e syndrome 2, autosomal recessive, with recurrent infections	Hyper-immunoglobulin e syndrome with recurrent infections	0.50000	1	6.457e-5	1.985e-4
Hyper-immunoglobulin e syndrome 2, autosomal recessive, with recurrent infections	Intellectual developmental disorder, autosomal dominant 2	0.50000	1	6.457e-5	1.985e-4
Hyper-immunoglobulin e syndrome with recurrent infections	Intellectual developmental disorder, autosomal dominant 2	0.50000	1	6.457e-5	1.985e-4
Hyper-immunoglobulin e syndrome 2, autosomal recessive, with recurrent infections	Insulin dependent diabetes mellitus secretory diarrhea syndrome	0.25000	1	1.937e-4	4.415e-4
Hyper-immunoglobulin e syndrome with recurrent infections	Insulin dependent diabetes mellitus secretory diarrhea syndrome	0.25000	1	1.937e-4	4.415e-4
Insulin dependent diabetes mellitus secretory diarrhea syndrome	Intellectual developmental disorder, autosomal dominant 2	0.25000	1	1.937e-4	4.415e-4
Insulin dependent diabetes mellitus secretory diarrhea syndrome	Ritscher-schinzel syndrome 2	0.25000	1	1.937e-4	4.415e-4