

GeDiPNet Cluster 266 -- 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 03:00:13. · Rows: 11 · Generated: 2026-10-07 11:13:23

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hyper-immunoglobulin m syndrome	Hyper-immunoglobulin syndrome	0.40000	2	5.062e-8	3.915e-7
Blast crisis	hyper-IgM syndrome type 2	0.33333	1	1.299e-4	3.898e-4
hyper-IgM syndrome type 2	Hyper-immunoglobulin syndrome	0.33333	1	1.299e-4	3.898e-4
hyper-IgM syndrome type 3	Hyper-immunoglobulin syndrome	0.33333	1	1.299e-4	3.898e-4
Blast crisis	Hyper-immunoglobulin syndrome	0.25000	1	2.598e-4	6.403e-4
hyper-IgM syndrome type 2	Hyper-immunoglobulin m syndrome	0.20000	1	2.598e-4	6.403e-4
hyper-IgM syndrome type 3	Hyper-immunoglobulin m syndrome	0.20000	1	2.598e-4	6.403e-4
Blast crisis	Myelogenous leukemia	0.20000	1	3.896e-4	8.521e-4
Blast crisis	Chromosome 22q11.2 microdeletion syndrome	0.20000	1	3.896e-4	8.521e-4
Blast crisis	Hyper-immunoglobulin m syndrome	0.16667	1	5.195e-4	1.041e-3
Chromosome 22q11.2 microdeletion syndrome	Myelogenous leukemia	0.16667	1	5.844e-4	1.135e-3