

GeDiPNet Cluster 128 -- 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: 2026-10-09 03:00:12. · Rows: 18 · Generated: 2026-10-09 09:24:13

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hyper-igm immunodeficiency syndrome	Hyper-immunoglobulin m syndrome	0.50000	4	1.495e-14	2.063e-13
Hyper-immunoglobulin m syndrome	Hyper-immunoglobulin syndrome	0.40000	2	5.062e-8	3.709e-7
Hyper-igm immunodeficiency syndrome	Hyper-immunoglobulin syndrome	0.25000	2	1.772e-7	1.182e-6
Hodgkin disease	Hyper-immunoglobulin m syndrome	0.10000	2	6.875e-6	3.488e-5
Hodgkin disease	Hyper-igm immunodeficiency syndrome	0.08696	2	2.401e-5	1.128e-4
Blast crisis	Hyper-immunoglobulin syndrome	0.25000	1	2.598e-4	6.505e-4
Hyper-immunoglobulin m syndrome	X-linked hyper-igm syndrome	0.20000	1	2.598e-4	6.505e-4
Blast crisis	Myelogenous leukemia	0.20000	1	3.896e-4	8.660e-4
Blast crisis	Chromosome 22q11.2 microdeletion syndrome	0.20000	1	3.896e-4	8.660e-4
Hyper-igm immunodeficiency syndrome	immunodeficiency, common variable, 2	0.12500	1	4.546e-4	9.721e-4
Hyper-igm immunodeficiency syndrome	X-linked hyper-igm syndrome	0.12500	1	4.546e-4	9.721e-4
Bloom syndrome	Hyper-immunoglobulin m syndrome	0.16667	1	5.195e-4	1.060e-3
Blast crisis	Hyper-immunoglobulin m syndrome	0.16667	1	5.195e-4	1.060e-3
Chromosome 22q11.2 microdeletion syndrome	Myelogenous leukemia	0.16667	1	5.844e-4	1.156e-3
Bloom syndrome	Hyper-igm immunodeficiency syndrome	0.11111	1	9.090e-4	1.575e-3
Blast crisis	Hyper-igm immunodeficiency syndrome	0.11111	1	9.090e-4	1.575e-3
Hodgkin disease	X-linked hyper-igm syndrome	0.05556	1	1.104e-3	1.826e-3
Hodgkin disease	Hyper-immunoglobulin syndrome	0.05263	1	2.207e-3	3.112e-3