

GeDiPNet Cluster 89 -- 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: 15 · Generated: 2026-10-07 02:51:05

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hodgkin lymphoma	Lymphocytic leukemia	0.15228	30	1.883e-39	8.184e-38
Hodgkin lymphoma	Multiple myeloma	0.13839	31	5.272e-38	2.209e-36
Lymphocytic leukemia	Multiple myeloma	0.13100	30	8.999e-36	3.542e-34
Lymphocytic leukemia	Lymphoid leukemia	0.10769	14	2.728e-23	6.416e-22
Lymphoid leukemia	Multiple myeloma	0.07500	12	6.897e-18	1.246e-16
Lymphocytic b-cell leukemia	Lymphocytic leukemia	0.07143	10	4.089e-14	5.765e-13
Myeloid leukemia	Promyelocytic leukemia	0.06349	12	4.132e-14	5.819e-13
Hodgkin lymphoma	Lymphoid leukemia	0.06870	9	2.328e-13	3.137e-12
Leukemia	Myeloid leukemia	0.05435	10	8.281e-12	9.682e-11
Lymphocytic b-cell leukemia	Lymphoid leukemia	0.08621	5	3.659e-9	3.303e-8
Leukemia	Lymphoid leukemia	0.07463	5	1.252e-8	1.053e-7
Leukemia	SYNCRIP-related neurodevelopmental disorder	0.02273	1	2.793e-3	3.736e-3
immunodeficiency, common variable, 14	Promyelocytic leukemia	0.01961	1	3.247e-3	4.237e-3
Dentici novelli neurodevelopmental syndrome	Hodgkin lymphoma	0.00893	1	7.209e-3	8.517e-3
hypotaurinemic retinal degeneration and cardiomyopathy	Myeloid leukemia	0.00662	1	9.742e-3	1.106e-2