

GeDiPNet Cluster 225 -- 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: 2026-10-10 03:00:13. · Rows: 12 · Generated: 2026-10-10 01:46:00

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hodgkin lymphoma	Lymphocytic leukemia	0.15228	30	1.883e-39	7.729e-38
Hodgkin lymphoma	Multiple myeloma	0.13839	31	5.272e-38	2.086e-36
Lymphocytic leukemia	Multiple myeloma	0.13100	30	8.999e-36	3.351e-34
Lymphocytic leukemia	Lymphoid leukemia	0.10769	14	2.728e-23	6.059e-22
Lymphoid leukemia	Multiple myeloma	0.07500	12	6.897e-18	1.179e-16
Lymphocytic b-cell leukemia	Lymphocytic leukemia	0.07143	10	4.089e-14	5.448e-13
Hodgkin lymphoma	Lymphoid leukemia	0.06870	9	2.328e-13	2.962e-12
Lymphocytic b-cell leukemia	Lymphoid leukemia	0.08621	5	3.659e-9	3.121e-8
Hepatic veno occlusive disease with immunodeficiency	Lymphocytic leukemia	0.01724	2	5.530e-5	2.331e-4
Hepatic veno occlusive disease with immunodeficiency	Lymphoid leukemia	0.03333	1	3.634e-3	4.667e-3
Hepatic veno occlusive disease with immunodeficiency	Lymphocytic b-cell leukemia	0.02778	1	4.411e-3	5.528e-3
Dentici novelli neurodevelopmental syndrome	Hodgkin lymphoma	0.00893	1	7.209e-3	8.515e-3