

GeDiPNet Cluster 500 -- 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: 2026-10-20 03:00:20. · Rows: 11 · Generated: 2026-10-19 23:20:18

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Dedifferentiated solitary fibrous tumor	Meninges hemangiopericytoma	0.66667	2	8.338e-9	6.596e-8
Dedifferentiated solitary fibrous tumor	Hemangiopericytoma	0.50000	2	2.501e-8	1.874e-7
Hemangiopericytoma	Meninges hemangiopericytoma	0.50000	2	2.501e-8	1.874e-7
Dedifferentiated solitary fibrous tumor	Peritoneal solitary fibrous tumor	0.33333	2	8.338e-8	5.852e-7
Meninges hemangiopericytoma	Peritoneal solitary fibrous tumor	0.33333	2	8.338e-8	5.852e-7
Hemangiopericytoma	Peritoneal solitary fibrous tumor	0.28571	2	2.501e-7	1.654e-6
Dedifferentiated solitary fibrous tumor	Hyper-immunoglobulin e syndrome 6, autosomal dominant, with recurrent infections	0.33333	1	1.291e-4	3.269e-4
Hyper-immunoglobulin e syndrome 6, autosomal dominant, with recurrent infections	Meninges hemangiopericytoma	0.33333	1	1.291e-4	3.269e-4
Hemangiopericytoma	Hyper-immunoglobulin e syndrome 6, autosomal dominant, with recurrent infections	0.25000	1	1.937e-4	4.414e-4
Hyper-immunoglobulin e syndrome 6, autosomal dominant, with recurrent infections	Peritoneal solitary fibrous tumor	0.16667	1	3.228e-4	6.412e-4
Peritoneal solitary fibrous tumor	Tarp syndrome	0.16667	1	3.228e-4	6.412e-4