

GeDiPNet Cluster 126 -- 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: 2026-10-08 03:00:12. · Rows: 15 · Generated: 2026-10-08 04:38:38

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Erythrocytosis	Secondary polycythemia	0.57143	8	1.645e-26	4.423e-25
Erythrocytosis	Polycythemia	0.37500	9	4.926e-25	1.208e-23
Anemia	Polycythemia	0.13043	12	1.496e-23	3.379e-22
Anemia	Hemolytic anemia	0.11111	14	2.416e-20	4.692e-19
Polycythemia	Secondary polycythemia	0.27273	6	4.102e-17	6.756e-16
Hemolytic anemia	Hereditary hemolytic anemia	0.08065	5	2.194e-10	2.131e-9
Hereditary hemolytic anemia	Secondary polycythemia	0.11111	2	1.296e-5	6.337e-5
Erythrocytosis	Hereditary hemolytic anemia	0.08696	2	3.604e-5	1.662e-4
Bisphosphoglycerate mutase deficiency	Secondary polycythemia	0.11111	1	5.195e-4	1.058e-3
Bisphosphoglycerate mutase deficiency	Hereditary hemolytic anemia	0.08333	1	7.144e-4	1.333e-3
Coronary vessel anomalies	Hereditary hemolytic anemia	0.08333	1	7.144e-4	1.333e-3
Hereditary hemolytic anemia	inherited glutathione synthetase deficiency	0.08333	1	7.144e-4	1.333e-3
Bisphosphoglycerate mutase deficiency	Erythrocytosis	0.07143	1	8.443e-4	1.497e-3
Bisphosphoglycerate mutase deficiency	Hemolytic anemia	0.01786	1	3.572e-3	4.594e-3
Charcot-Marie-Tooth disease type 4B2	Hemolytic anemia	0.01786	1	3.572e-3	4.594e-3