

GeDiPNet Cluster 325 -- 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 17:17:47. · Rows: 9 · Generated: 2026-10-07 15:11:22

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Central retinal vein occlusion	hereditary thrombophilia due to congenital protein C deficiency	0.50000	1	6.494e-5	2.317e-4
Central retinal vein occlusion	Congenital thrombotic thrombocytopenic purpura	0.33333	1	1.299e-4	3.908e-4
Congenital thrombotic thrombocytopenic purpura	hereditary thrombophilia due to congenital protein C deficiency	0.33333	1	1.299e-4	3.908e-4
Congenital thrombotic thrombocytopenic purpura	Upshaw-schulman syndrome	0.33333	1	1.299e-4	3.908e-4
Three-vessel coronary artery disease	Upshaw-schulman syndrome	0.25000	1	1.948e-4	5.325e-4
Arteritis	Central retinal vein occlusion	0.20000	1	2.598e-4	6.484e-4
Arteritis	hereditary thrombophilia due to congenital protein C deficiency	0.20000	1	2.598e-4	6.484e-4
Congenital thrombotic thrombocytopenic purpura	Three-vessel coronary artery disease	0.20000	1	3.896e-4	8.637e-4
Arteritis	Congenital thrombotic thrombocytopenic purpura	0.16667	1	5.195e-4	1.058e-3