

GeDiPNet Cluster 400 -- 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 03:00:12. · Rows: 9 · Generated: 2026-10-07 10:01:30

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Retinitis pigmentosa with choroidal involvement	RPE65-related recessive retinopathy	0.50000	1	6.494e-5	2.342e-4
Retinitis pigmentosa with choroidal involvement	RPE65-related dominant retinopathy	0.50000	1	6.494e-5	2.342e-4
RPE65-related dominant retinopathy	RPE65-related recessive retinopathy	0.50000	1	6.494e-5	2.342e-4
Congenital blindness	Retinitis pigmentosa with choroidal involvement	0.25000	1	1.948e-4	5.278e-4
Congenital blindness	RPE65-related recessive retinopathy	0.25000	1	1.948e-4	5.278e-4
Congenital blindness	RPE65-related dominant retinopathy	0.25000	1	1.948e-4	5.278e-4
Congenital nystagmus	Retinitis pigmentosa with choroidal involvement	0.11111	1	5.195e-4	1.041e-3
Congenital nystagmus	RPE65-related dominant retinopathy	0.11111	1	5.195e-4	1.041e-3
Congenital nystagmus	RPE65-related recessive retinopathy	0.11111	1	5.195e-4	1.041e-3