

GeDiPNet Cluster 156 -- 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 03:00:13. · Rows: 15 · Generated: 2026-10-08 02:43:46

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	2.317e-4
Retinitis pigmentosa with choroidal involvement	RPE65-related dominant retinopathy	0.50000	1	6.494e-5	2.317e-4
RPE65-related dominant retinopathy	RPE65-related recessive retinopathy	0.50000	1	6.494e-5	2.317e-4
Congenital blindness	Retinitis pigmentosa with choroidal involvement	0.25000	1	1.948e-4	5.325e-4
Congenital blindness	RPE65-related recessive retinopathy	0.25000	1	1.948e-4	5.325e-4
Congenital blindness	RPE65-related dominant retinopathy	0.25000	1	1.948e-4	5.325e-4
Congenital nystagmus	Retinitis pigmentosa with choroidal involvement	0.11111	1	5.195e-4	1.058e-3
Congenital nystagmus	RPE65-related dominant retinopathy	0.11111	1	5.195e-4	1.058e-3
Congenital nystagmus	RPE65-related recessive retinopathy	0.11111	1	5.195e-4	1.058e-3
Aica-ribosiduria	Blindness	0.06667	1	9.092e-4	1.575e-3
AIPL1-related retinopathy	Blindness	0.06667	1	9.092e-4	1.575e-3
Blindness	LCA5-related retinopathy	0.06667	1	9.092e-4	1.575e-3
Blindness	Retinitis pigmentosa with choroidal involvement	0.06667	1	9.092e-4	1.575e-3
Blindness	RPE65-related dominant retinopathy	0.06667	1	9.092e-4	1.575e-3
Blindness	RPE65-related recessive retinopathy	0.06667	1	9.092e-4	1.575e-3