

GeDiPNet Cluster 309 -- 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: 12 · Generated: 2026-10-08 03:40:54

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Distal myopathy	Vitelliform macular dystrophy	0.25000	4	2.136e-12	2.477e-11
Benign concentric annular macular dystrophy	IMPG1-related dominant retinopathy	0.50000	1	6.494e-5	2.317e-4
Benign concentric annular macular dystrophy	IMPG1-related recessive retinopathy	0.50000	1	6.494e-5	2.317e-4
IMPG1-related dominant retinopathy	IMPG1-related recessive retinopathy	0.50000	1	6.494e-5	2.317e-4
Benign concentric annular macular dystrophy	Vitelliform macular dystrophy	0.16667	1	3.247e-4	7.680e-4
IMPG1-related dominant retinopathy	Vitelliform macular dystrophy	0.16667	1	3.247e-4	7.680e-4
IMPG1-related recessive retinopathy	Vitelliform macular dystrophy	0.16667	1	3.247e-4	7.680e-4
IMPG2-related recessive retinopathy	Vitelliform macular dystrophy	0.16667	1	3.247e-4	7.680e-4
Benign concentric annular macular dystrophy	Distal myopathy	0.06667	1	9.092e-4	1.575e-3
Distal myopathy	IMPG1-related dominant retinopathy	0.06667	1	9.092e-4	1.575e-3
Distal myopathy	IMPG1-related recessive retinopathy	0.06667	1	9.092e-4	1.575e-3
Distal myopathy	IMPG2-related recessive retinopathy	0.06667	1	9.092e-4	1.575e-3