

GeDiPNet Cluster 157 -- 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: 19 · Generated: 2026-10-07 22:48:20

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Night blindness, congenital stationary	Oguchi disease	0.48148	13	1.665e-37	6.533e-36
Congenital stationary night blindness	Night blindness, congenital stationary	0.46154	12	1.408e-35	5.190e-34
Congenital stationary night blindness	Oguchi disease	0.58824	10	3.207e-31	1.023e-29
Congenital stationary night blindness	inherited retinal dystrophy	0.12000	3	1.637e-7	1.109e-6
inherited retinal dystrophy	Oguchi disease	0.11111	3	2.705e-7	1.767e-6
inherited retinal dystrophy	Night blindness, congenital stationary	0.07895	3	1.698e-6	9.624e-6
Congenital stationary night blindness	GRM6-related retinopathy	0.07692	1	7.793e-4	1.404e-3
Congenital stationary night blindness	GPR179-related retinopathy	0.07692	1	7.793e-4	1.404e-3
Congenital stationary night blindness	NYX-related retinopathy	0.07692	1	7.793e-4	1.404e-3
Congenital stationary night blindness	TRPM1-related retinopathy	0.07692	1	7.793e-4	1.404e-3
GRM6-related retinopathy	Oguchi disease	0.06667	1	9.092e-4	1.575e-3
NYX-related retinopathy	Oguchi disease	0.06667	1	9.092e-4	1.575e-3
Oguchi disease	retinitis pigmentosa 47	0.06667	1	9.092e-4	1.575e-3
Oguchi disease	TRPM1-related retinopathy	0.06667	1	9.092e-4	1.575e-3
GRM6-related retinopathy	Night blindness, congenital stationary	0.03846	1	1.624e-3	2.459e-3
GPR179-related retinopathy	Night blindness, congenital stationary	0.03846	1	1.624e-3	2.459e-3
Night blindness, congenital stationary	NYX-related retinopathy	0.03846	1	1.624e-3	2.459e-3
Night blindness, congenital stationary	retinitis pigmentosa 47	0.03846	1	1.624e-3	2.459e-3
Night blindness, congenital stationary	TRPM1-related retinopathy	0.03846	1	1.624e-3	2.459e-3