

GeDiPNet Cluster 385 -- 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: 10 · Generated: 2026-10-07 14:17:09

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Cone-rod dystrophy, x-linked	X-linked cone-rod dystrophy	0.66667	2	8.436e-9	6.837e-8
Ciliary dyskinesia with retinitis pigmentosa	Retinitis pigmentosa, x-linked, and sinorespiratory infections with or without deafness	0.50000	1	6.494e-5	2.317e-4
Ciliary dyskinesia with retinitis pigmentosa	RPGR-related retinopathy	0.50000	1	6.494e-5	2.317e-4
Retinitis pigmentosa, x-linked, and sinorespiratory infections with or without deafness	RPGR-related retinopathy	0.50000	1	6.494e-5	2.317e-4
Ciliary dyskinesia with retinitis pigmentosa	Cone-rod dystrophy, x-linked	0.33333	1	1.299e-4	3.908e-4
Ciliary dyskinesia with retinitis pigmentosa	X-linked cone-rod dystrophy	0.33333	1	1.299e-4	3.908e-4
Cone-rod dystrophy, x-linked	Retinitis pigmentosa, x-linked, and sinorespiratory infections with or without deafness	0.33333	1	1.299e-4	3.908e-4
Retinitis pigmentosa, x-linked, and sinorespiratory infections with or without deafness	X-linked cone-rod dystrophy	0.33333	1	1.299e-4	3.908e-4
Cone-rod dystrophy, x-linked	RPGR-related retinopathy	0.33333	1	1.299e-4	3.908e-4
RPGR-related retinopathy	X-linked cone-rod dystrophy	0.33333	1	1.299e-4	3.908e-4