GeDiPNet Shared-Gene Disease Pairs

Similarity: Jaccard index on shared curated genes (disease_gdp). Significance: upper-tail hypergeometric test (P(X>=k)), Benjamini-Hochberg FDR-corrected across all tested pairs. Corroborated Shared Genes: count of shared genes backed by >=2 distinct disease_gdp.main_source values combined across both diseases (e.g. GWAS-backed for one, ClinVar-backed for the other still counts); Shared Gene Names lists each gene's own source count in parentheses. Overlap Coefficient: shared genes / MIN(each disease's own total gene count) -- complements the Jaccard-based Similarity Score by showing how much of the SMALLER disease's entire known gene-set is covered by this pair, which Jaccard (dividing by the union) can understate. Shared Cluster: cluster id if both diseases fall in the same label-propagation cluster (scripts/build_disease_clusters.php), blank otherwise. See scripts/build_related.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-20 03:00:20. Filters applied: disease name contains "Retinal dystrophy with or without macular staphyloma" (5 matching pairs). · Rows: 5 · Generated: 2026-10-20 06:33:11

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Cone-rod dystrophy	Retinal dystrophy with or without macular staphyloma	1	1	CFAP410 (4)	0.01299	1.00000	4.907e-3	5.888e-3	
Leber congenital amaurosis	Retinal dystrophy with or without macular staphyloma	1	1	CFAP410 (2)	0.01351	1.00000	4.713e-3	5.685e-3	
Cone dystrophy	Retinal dystrophy with or without macular staphyloma	1	1	CFAP410 (2)	0.02500	1.00000	2.518e-3	3.269e-3	
Retinal dystrophy with or without macular staphyloma	Spondylometaphyseal dysplasia	1	1	CFAP410 (3)	0.06667	1.00000	9.039e-4	1.411e-3	
Axial spondylometaphyseal dysplasia	Retinal dystrophy with or without macular staphyloma	1	1	CFAP410 (3)	0.50000	1.00000	6.457e-5	1.985e-4	