

GeDiPNet Cluster 229 -- 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: 2026-10-09 03:00:13. · Rows: 11 · Generated: 2026-10-09 09:27:04

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital epicanthus	Esophageal atresia	0.06818	3	2.922e-6	1.581e-5
ATP1A3-associated neurological disorder	Capos syndrome	0.50000	1	6.494e-5	2.329e-4
ATP1A3-associated neurological disorder	Cerebellar ataxia, areflexia, pes cavus, optic atrophy, and sensorineural hearing loss	0.50000	1	6.494e-5	2.329e-4
Capos syndrome	Cerebellar ataxia, areflexia, pes cavus, optic atrophy, and sensorineural hearing loss	0.50000	1	6.494e-5	2.329e-4
Alternating hemiplegia of childhood	Capos syndrome	0.33333	1	1.299e-4	3.929e-4
Alternating hemiplegia of childhood	Cerebellar ataxia, areflexia, pes cavus, optic atrophy, and sensorineural hearing loss	0.33333	1	1.299e-4	3.929e-4
Alternating hemiplegia of childhood	ATP1A3-associated neurological disorder	0.33333	1	1.299e-4	3.929e-4
ATP1A3-associated neurological disorder	Congenital epicanthus	0.06667	1	9.092e-4	1.575e-3
Capos syndrome	Congenital epicanthus	0.06667	1	9.092e-4	1.575e-3
Cerebellar ataxia, areflexia, pes cavus, optic atrophy, and sensorineural hearing loss	Congenital epicanthus	0.06667	1	9.092e-4	1.575e-3
Congenital epicanthus	fibrodysplasia ossificans progressiva	0.06667	1	9.092e-4	1.575e-3