

GeDiPNet Cluster 598 -- 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: 2026-10-20 03:00:19. · Rows: 8 · Generated: 2026-10-19 23:20:06

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	Sengers syndrome	0.60000	3	6.461e-12	6.555e-11
Cataract 38, autosomal recessive	Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	0.25000	1	1.937e-4	4.414e-4
Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	Triokinase and fmh cyclase deficiency	0.25000	1	1.937e-4	4.414e-4
Cataract 38, autosomal recessive	Sengers syndrome	0.20000	1	2.583e-4	5.384e-4
Cataract 38, autosomal recessive	Trichohepatoenteric syndrome	0.20000	1	2.583e-4	5.384e-4
Sengers syndrome	Triokinase and fmh cyclase deficiency	0.20000	1	2.583e-4	5.384e-4
Congenital cataract hypertrophic cardiomyopathy mitochondrial myopathy syndrome	Trichohepatoenteric syndrome	0.14286	1	7.746e-4	1.247e-3
Sengers syndrome	Trichohepatoenteric syndrome	0.12500	1	1.033e-3	1.574e-3