

GeDiPNet Cluster 510 -- 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 23:20:31

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hereditary spastic paraplegia 8	Intellectual disability, congenital cardiac malformation and dandy-walker malformation	0.50000	1	6.457e-5	1.985e-4
Hereditary spastic paraplegia 8	Ritscher-schinzel syndrome 1	0.50000	1	6.457e-5	1.985e-4
Intellectual disability, congenital cardiac malformation and dandy-walker malformation	Ritscher-schinzel syndrome 1	0.50000	1	6.457e-5	1.985e-4
Hereditary spastic paraplegia 8	Ritscher-schinzler syndrome	0.20000	1	2.583e-4	5.384e-4
Intellectual disability, congenital cardiac malformation and dandy-walker malformation	Ritscher-schinzler syndrome	0.20000	1	2.583e-4	5.384e-4
Ritscher-schinzel syndrome 1	Ritscher-schinzler syndrome	0.20000	1	2.583e-4	5.384e-4
Ritscher-schinzel syndrome 3	Ritscher-schinzler syndrome	0.20000	1	2.583e-4	5.384e-4
Ritscher-schinzel syndrome 4	Ritscher-schinzler syndrome	0.20000	1	2.583e-4	5.384e-4