

GeDiPNet Cluster 239 -- 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: 2026-10-08 03:00:13. · Rows: 11 · Generated: 2026-10-08 11:15:15

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hereditary spastic paraplegia	Spastic paraplegia	0.43226	67	3.878e-125	5.878e-123
complex hereditary spastic paraplegia	Hereditary spastic paraplegia	0.06863	7	4.230e-16	6.515e-15
complex hereditary spastic paraplegia	Spastic paraplegia	0.05785	7	1.463e-15	2.173e-14
Allan-herndon-dudley syndrome	Hereditary spastic paraplegia	0.00980	1	6.559e-3	7.817e-3
Hereditary spastic paraplegia	hereditary spastic paraplegia 11	0.00980	1	6.559e-3	7.817e-3
Hereditary spastic paraplegia	hereditary spastic paraplegia 18	0.00980	1	6.559e-3	7.817e-3
Hereditary spastic paraplegia	hereditary spastic paraplegia 62	0.00980	1	6.559e-3	7.817e-3
Allan-herndon-dudley syndrome	Spastic paraplegia	0.00826	1	7.793e-3	9.067e-3
hereditary spastic paraplegia 11	Spastic paraplegia	0.00826	1	7.793e-3	9.067e-3
hereditary spastic paraplegia 18	Spastic paraplegia	0.00826	1	7.793e-3	9.067e-3
hereditary spastic paraplegia 62	Spastic paraplegia	0.00826	1	7.793e-3	9.067e-3