

GeDiPNet Cluster 92 -- 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: 2026-10-07 03:00:12. · Rows: 16 · Generated: 2026-10-07 09:17:09

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Cerebellar ataxia	Spinocerebellar ataxia	0.20994	38	2.791e-56	1.704e-54
Cerebellar ataxia	Spastic ataxia	0.11446	19	3.210e-25	8.382e-24
Spastic ataxia	Spinocerebellar ataxia	0.07362	12	3.612e-14	5.115e-13
Spastic ataxia	Spastic ataxia optic atrophy dysarthria syndrome	0.01408	1	4.546e-3	5.665e-3
Spastic ataxia	spinocerebellar ataxia type 42	0.01408	1	4.546e-3	5.665e-3
autosomal recessive cerebellar ataxia	Spinocerebellar ataxia	0.00952	1	6.754e-3	8.028e-3
Spinocerebellar ataxia	spinocerebellar ataxia type 13	0.00952	1	6.754e-3	8.028e-3
Spinocerebellar ataxia	spinocerebellar ataxia type 42	0.00952	1	6.754e-3	8.028e-3
Spinocerebellar ataxia	spinocerebellar ataxia, autosomal recessive 23	0.00952	1	6.754e-3	8.028e-3
Spinocerebellar ataxia	spinocerebellar ataxia, autosomal recessive 27	0.00952	1	6.754e-3	8.028e-3
Spinocerebellar ataxia	spinocerebellar ataxia, autosomal recessive 28	0.00952	1	6.754e-3	8.028e-3
autosomal recessive cerebellar ataxia	Cerebellar ataxia	0.00870	1	7.404e-3	8.702e-3
Cayman type cerebellar ataxia	Cerebellar ataxia	0.00870	1	7.404e-3	8.702e-3
Cerebellar ataxia	spinocerebellar ataxia type 42	0.00870	1	7.404e-3	8.702e-3
Cerebellar ataxia	spinocerebellar ataxia, autosomal recessive 23	0.00870	1	7.404e-3	8.702e-3
3m syndrome	Spinocerebellar ataxia	0.00935	1	2.013e-2	2.191e-2