

GeDiPNet Cluster 451 -- 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: 10 · Generated: 2026-10-19 23:22:20

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Autosomal recessive spinocerebellar ataxia 2	Cerebelloparenchymal disorder	0.50000	1	6.457e-5	1.985e-4
Autosomal recessive spinocerebellar ataxia 2	Spinocerebellar ataxia, autosomal recessive 2 ar	0.50000	1	6.457e-5	1.985e-4
Cerebelloparenchymal disorder	Spinocerebellar ataxia, autosomal recessive 2 ar	0.50000	1	6.457e-5	1.985e-4
Autosomal recessive spinocerebellar ataxia 2	Normal pressure hydrocephalus	0.33333	1	1.291e-4	3.269e-4
Cerebelloparenchymal disorder	Normal pressure hydrocephalus	0.33333	1	1.291e-4	3.269e-4
Normal pressure hydrocephalus	Spermatogenic failure 19	0.33333	1	1.291e-4	3.269e-4
Normal pressure hydrocephalus	Spinocerebellar ataxia, autosomal recessive 2 ar	0.33333	1	1.291e-4	3.269e-4
Autosomal recessive spinocerebellar ataxia 2	Ptosis	0.11111	1	5.165e-4	9.125e-4
Cerebelloparenchymal disorder	Ptosis	0.11111	1	5.165e-4	9.125e-4
Ptosis	Spinocerebellar ataxia, autosomal recessive 2 ar	0.11111	1	5.165e-4	9.125e-4