

GeDiPNet Cluster 541 -- 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 03:00:18. · Rows: 10 · Generated: 2026-10-19 23:19:42

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Amish nemaline myopathy	Nemaline myopathy 5	0.50000	1	6.457e-5	1.985e-4
Amish nemaline myopathy	Nemaline myopathy 5a, autosomal recessive, severe infantile	0.50000	1	6.457e-5	1.985e-4
Amish nemaline myopathy	Nemaline myopathy 5c, autosomal dominant	0.50000	1	6.457e-5	1.985e-4
Nemaline myopathy 5	Nemaline myopathy 5a, autosomal recessive, severe infantile	0.50000	1	6.457e-5	1.985e-4
Nemaline myopathy 5	Nemaline myopathy 5c, autosomal dominant	0.50000	1	6.457e-5	1.985e-4
Nemaline myopathy 5a, autosomal recessive, severe infantile	Nemaline myopathy 5c, autosomal dominant	0.50000	1	6.457e-5	1.985e-4
Amish nemaline myopathy	Childhood-onset nemaline myopathy	0.25000	1	1.937e-4	4.414e-4
Childhood-onset nemaline myopathy	Nemaline myopathy 5	0.25000	1	1.937e-4	4.414e-4
Childhood-onset nemaline myopathy	Nemaline myopathy 5a, autosomal recessive, severe infantile	0.25000	1	1.937e-4	4.414e-4
Childhood-onset nemaline myopathy	Nemaline myopathy 5c, autosomal dominant	0.25000	1	1.937e-4	4.414e-4