

GeDiPNet Cluster 628 -- 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 23:19:03

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Cutis laxa, autosomal dominant, 2	Cutis laxa, autosomal recessive, type ia	0.50000	1	6.457e-5	1.985e-4
Cutis laxa, autosomal dominant, 2	Cutis laxa, autosomal recessive, 1a	0.50000	1	6.457e-5	1.985e-4
Cutis laxa, autosomal recessive, 1a	Cutis laxa, autosomal recessive, type ia	0.50000	1	6.457e-5	1.985e-4
Cutis laxa, autosomal dominant, 2	Demyelinating hereditary motor and sensory neuropathy	0.33333	1	1.291e-4	3.269e-4
Cutis laxa, autosomal recessive, 1a	Demyelinating hereditary motor and sensory neuropathy	0.33333	1	1.291e-4	3.269e-4
Cutis laxa, autosomal recessive, type ia	Demyelinating hereditary motor and sensory neuropathy	0.33333	1	1.291e-4	3.269e-4
Cutis laxa, autosomal dominant, 2	Proximal spinal muscular atrophy	0.14286	1	3.874e-4	7.329e-4
Cutis laxa, autosomal recessive, 1a	Proximal spinal muscular atrophy	0.14286	1	3.874e-4	7.329e-4
Cutis laxa, autosomal recessive, type ia	Proximal spinal muscular atrophy	0.14286	1	3.874e-4	7.329e-4
Demyelinating hereditary motor and sensory neuropathy	Proximal spinal muscular atrophy	0.12500	1	7.747e-4	1.247e-3