

GeDiPNet Cluster 340 -- 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 19:28:52

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Chromosome 3p25 monosomy	Rippling muscle disease	0.40000	2	7.592e-8	5.427e-7
Chromosome 3p25 monosomy	Tubular aggregate myopathy	0.22222	2	5.313e-7	3.282e-6
Rippling muscle disease	Tubular aggregate myopathy	0.22222	2	5.313e-7	3.282e-6
Caveolinopathy	Creatine phosphokinase elevation	0.50000	1	6.494e-5	2.317e-4
Caveolinopathy	Chromosome 3p25 monosomy	0.25000	1	1.948e-4	5.325e-4
Caveolinopathy	Rippling muscle disease	0.25000	1	1.948e-4	5.325e-4
Chromosome 3p25 monosomy	Creatine phosphokinase elevation	0.25000	1	1.948e-4	5.325e-4
Creatine phosphokinase elevation	Rippling muscle disease	0.25000	1	1.948e-4	5.325e-4
Caveolinopathy	Tubular aggregate myopathy	0.12500	1	4.546e-4	9.697e-4
Creatine phosphokinase elevation	Tubular aggregate myopathy	0.12500	1	4.546e-4	9.697e-4
DPAGT1-congenital disorder of glycosylation	Tubular aggregate myopathy	0.12500	1	4.546e-4	9.697e-4