

GeDiPNet Cluster 212 -- 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 17:17:46. · Rows: 15 · Generated: 2026-10-07 21:04:47

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Benign congenital myopathy	Dystrophinopathy	0.50000	1	6.494e-5	2.317e-4
Benign congenital myopathy	progressive muscular dystrophy	0.50000	1	6.494e-5	2.317e-4
Dystrophinopathy	progressive muscular dystrophy	0.50000	1	6.494e-5	2.317e-4
Becker muscular dystrophy	Benign congenital myopathy	0.25000	1	1.948e-4	5.325e-4
Becker muscular dystrophy	Dystrophinopathy	0.25000	1	1.948e-4	5.325e-4
Becker muscular dystrophy	progressive muscular dystrophy	0.25000	1	1.948e-4	5.325e-4
Benign congenital myopathy	Leiomyosarcoma	0.12500	1	4.546e-4	9.697e-4
Dystrophinopathy	Leiomyosarcoma	0.12500	1	4.546e-4	9.697e-4
hereditary leiomyomatosis and renal cell cancer	Leiomyosarcoma	0.12500	1	4.546e-4	9.697e-4
Leiomyosarcoma	progressive muscular dystrophy	0.12500	1	4.546e-4	9.697e-4
Becker muscular dystrophy	Leiomyosarcoma	0.10000	1	1.363e-3	2.146e-3
Leiomyosarcoma	Pseudo-torch syndrome	0.10000	1	1.363e-3	2.146e-3
Benign congenital myopathy	Duchenne muscular dystrophy	0.03333	1	1.883e-3	2.760e-3
Duchenne muscular dystrophy	Dystrophinopathy	0.03333	1	1.883e-3	2.760e-3
Duchenne muscular dystrophy	progressive muscular dystrophy	0.03333	1	1.883e-3	2.760e-3