

GeDiPNet Cluster 440 -- 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: 14 · Generated: 2026-10-19 23:20:08

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Ascending aortic dilation	Pectus excavatum	0.50000	1	6.457e-5	1.985e-4
Ascending aortic dilation	Failure to thrive in infancy	0.50000	1	6.457e-5	1.985e-4
Ascending aortic dilation	Generalized joint laxity	0.50000	1	6.457e-5	1.985e-4
Failure to thrive in infancy	Pectus excavatum	0.50000	1	6.457e-5	1.985e-4
Failure to thrive in infancy	Generalized joint laxity	0.50000	1	6.457e-5	1.985e-4
Generalized joint laxity	Pectus excavatum	0.50000	1	6.457e-5	1.985e-4
Ascending aortic dilation	Congenital finger flexion contractures	0.33333	1	1.291e-4	3.269e-4
Congenital finger flexion contractures	Pectus excavatum	0.33333	1	1.291e-4	3.269e-4
Congenital finger flexion contractures	Failure to thrive in infancy	0.33333	1	1.291e-4	3.269e-4
Congenital finger flexion contractures	Generalized joint laxity	0.33333	1	1.291e-4	3.269e-4
Ascending aortic dilation	Congenital musculoskeletal anomalies	0.14286	1	3.874e-4	7.329e-4
Congenital musculoskeletal anomalies	Pectus excavatum	0.14286	1	3.874e-4	7.329e-4
Congenital musculoskeletal anomalies	Failure to thrive in infancy	0.14286	1	3.874e-4	7.329e-4
Congenital musculoskeletal anomalies	Generalized joint laxity	0.14286	1	3.874e-4	7.329e-4