

GeDiPNet Cluster 230 -- 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 03:00:12. · Rows: 13 · Generated: 2026-10-07 00:37:14

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
holoprosencephaly 3	Preaxial polydactyly with upper back hypertrichosis	0.50000	1	6.494e-5	2.342e-4
holoprosencephaly 3	Skeletal system disorder	0.50000	1	6.494e-5	2.342e-4
Preaxial polydactyly with upper back hypertrichosis	Skeletal system disorder	0.50000	1	6.494e-5	2.342e-4
holoprosencephaly 3	Triphalangeal thumb-polysyndactyly syndrome	0.33333	1	1.299e-4	3.898e-4
Preaxial polydactyly with upper back hypertrichosis	Triphalangeal thumb-polysyndactyly syndrome	0.33333	1	1.299e-4	3.898e-4
Skeletal system disorder	Triphalangeal thumb-polysyndactyly syndrome	0.33333	1	1.299e-4	3.898e-4
holoprosencephaly 3	Partial agenesis of corpus callosum	0.25000	1	1.948e-4	5.278e-4
Partial agenesis of corpus callosum	Preaxial polydactyly with upper back hypertrichosis	0.25000	1	1.948e-4	5.278e-4
Partial agenesis of corpus callosum	Skeletal system disorder	0.25000	1	1.948e-4	5.278e-4
Congenital hypogonadotropic hypogonadism	Schizencephaly	0.16667	1	3.247e-4	7.577e-4
Preaxial polydactyly with upper back hypertrichosis	Schizencephaly	0.16667	1	3.247e-4	7.577e-4
Schizencephaly	Skeletal system disorder	0.16667	1	3.247e-4	7.577e-4
holoprosencephaly 3	Schizencephaly	0.16667	1	3.247e-4	7.577e-4