

GeDiPNet Cluster 398 -- 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 03:00:13. · Rows: 10 · Generated: 2026-10-07 03:52:08

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Arachnodactyly	Byzanthine arch palate	0.11765	2	6.571e-6	3.533e-5
Ck syndrome	Congenital hemidysplasia with ichthyosiform erythroderma and limb defects	0.50000	1	6.494e-5	2.342e-4
Ck syndrome	Perisylvian polymicrogyria	0.33333	1	1.299e-4	3.898e-4
Congenital hemidysplasia with ichthyosiform erythroderma and limb defects	Perisylvian polymicrogyria	0.33333	1	1.299e-4	3.898e-4
Arachnodactyly	Ck syndrome	0.16667	1	3.247e-4	7.577e-4
Arachnodactyly	Congenital hemidysplasia with ichthyosiform erythroderma and limb defects	0.16667	1	3.247e-4	7.577e-4
Arachnodactyly	Perisylvian polymicrogyria	0.14286	1	6.494e-4	1.219e-3
Byzanthine arch palate	Ck syndrome	0.07143	1	8.443e-4	1.478e-3
Byzanthine arch palate	Congenital hemidysplasia with ichthyosiform erythroderma and limb defects	0.07143	1	8.443e-4	1.478e-3
Byzanthine arch palate	Perisylvian polymicrogyria	0.06667	1	1.688e-3	2.514e-3