

GeDiPNet Cluster 301 -- 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: 6 · Generated: 2026-10-08 14:22:35

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hyperoxaluria	Pfaundler-hurler syndrome	0.14286	2	1.670e-6	9.465e-6
mucopolysaccharidosis type 1	Pfaundler-hurler syndrome	0.25000	1	1.948e-4	5.325e-4
Brachydactyly-elbow wrist dysplasia syndrome	Pfaundler-hurler syndrome	0.20000	1	3.896e-4	8.637e-4
Alanine-glyoxylate aminotransferase deficiency	Hyperoxaluria	0.07692	1	7.793e-4	1.404e-3
Hyperoxaluria	mucopolysaccharidosis type 1	0.07692	1	7.793e-4	1.404e-3
Hyperoxaluria	primary hyperoxaluria type 3	0.07692	1	7.793e-4	1.404e-3