

GeDiPNet Cluster 348 -- 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 03:00:12. · Rows: 9 · Generated: 2026-10-08 03:37:04

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Chromosome 15q24 deletion syndrome	SIN3A-related intellectual disability syndrome	0.50000	1	6.494e-5	2.317e-4
15q24 microdeletion	Chromosome 15q24 deletion syndrome	0.33333	1	1.299e-4	3.908e-4
15q24 microdeletion	SIN3A-related intellectual disability syndrome	0.33333	1	1.299e-4	3.908e-4
Chromosome 15q24 deletion syndrome	Witteveen-kolk syndrome	0.33333	1	1.299e-4	3.908e-4
SIN3A-related intellectual disability syndrome	Witteveen-kolk syndrome	0.33333	1	1.299e-4	3.908e-4
15q24 microdeletion	Witteveen-kolk syndrome	0.25000	1	2.598e-4	6.484e-4
Chromosome 15q24 deletion syndrome	Congenital diaphragmatic hernia	0.04762	1	1.299e-3	2.060e-3
Congenital diaphragmatic hernia	SIN3A-related intellectual disability syndrome	0.04762	1	1.299e-3	2.060e-3
Congenital diaphragmatic hernia	Witteveen-kolk syndrome	0.04545	1	2.596e-3	3.512e-3