

GeDiPNet Cluster 268 -- 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: 2026-10-10 03:00:12. · Rows: 11 · Generated: 2026-10-10 09:42:20

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
heyne-sprout-jackson syndrome	Microcephalic dwarfism	0.50000	1	6.494e-5	2.331e-4
heyne-sprout-jackson syndrome	Tatton-Brown-Rahman overgrowth syndrome	0.50000	1	6.494e-5	2.331e-4
Microcephalic dwarfism	Tatton-Brown-Rahman overgrowth syndrome	0.50000	1	6.494e-5	2.331e-4
Clonal cytopenia of undetermined significance	Microcephalic dwarfism	0.25000	1	1.948e-4	5.354e-4
Clonal cytopenia of undetermined significance	heyne-sprout-jackson syndrome	0.25000	1	1.948e-4	5.354e-4
Clonal cytopenia of undetermined significance	Tatton-Brown-Rahman overgrowth syndrome	0.25000	1	1.948e-4	5.354e-4
coffin-lowry syndrome	Specific learning disability	0.07692	1	7.793e-4	1.405e-3
Developmental delay with behavioral abnormalities	Specific learning disability	0.07692	1	7.793e-4	1.405e-3
Microcephalic dwarfism	Specific learning disability	0.07692	1	7.793e-4	1.405e-3
heyne-sprout-jackson syndrome	Specific learning disability	0.07692	1	7.793e-4	1.405e-3
Specific learning disability	Tatton-Brown-Rahman overgrowth syndrome	0.07692	1	7.793e-4	1.405e-3