

GeDiPNet Cluster 338 -- 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: 9 · Generated: 2026-10-08 19:29:12

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Cardiac anomalies-short stature-joint hypermobility-facial dysmorphism syndrome due to tab2 mutation	Chromosome 6q24-q25 deletion syndrome	0.50000	1	6.494e-5	2.317e-4
Cardiac anomalies-short stature-joint hypermobility-facial dysmorphism syndrome due to tab2 mutation	congenital heart defects, multiple types, 2	0.50000	1	6.494e-5	2.317e-4
Chromosome 6q24-q25 deletion syndrome	congenital heart defects, multiple types, 2	0.50000	1	6.494e-5	2.317e-4
Cardiac anomalies-short stature-joint hypermobility-facial dysmorphism syndrome due to tab2 mutation	Stress urinary incontinence	0.33333	1	1.299e-4	3.908e-4
Chromosome 6q24-q25 deletion syndrome	Stress urinary incontinence	0.33333	1	1.299e-4	3.908e-4
congenital heart defects, multiple types, 2	Stress urinary incontinence	0.33333	1	1.299e-4	3.908e-4
hereditary spastic paraplegia 8	Stress urinary incontinence	0.33333	1	1.299e-4	3.908e-4
hereditary spastic paraplegia 8	Ritscher-schinzler syndrome	0.20000	1	2.598e-4	6.484e-4
Ritscher-schinzler syndrome	Stress urinary incontinence	0.16667	1	5.195e-4	1.058e-3