

GeDiPNet Cluster 354 -- 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: 7 · Generated: 2026-10-08 13:43:10

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Dyssegmental dysplasia	Schwartz-jampel syndrome	0.50000	2	2.531e-8	1.932e-7
Dyssegmental dysplasia	Silverman-Handmaker type dyssegmental dysplasia	0.33333	1	1.299e-4	3.908e-4
Schwartz-jampel syndrome	Silverman-Handmaker type dyssegmental dysplasia	0.25000	1	1.948e-4	5.325e-4
1p36 deletion syndrome	Silverman-Handmaker type dyssegmental dysplasia	0.07143	1	8.443e-4	1.497e-3
1p36 deletion syndrome	radio-tartaglia syndrome	0.07143	1	8.443e-4	1.497e-3
1p36 deletion syndrome	Dyssegmental dysplasia	0.06667	1	1.688e-3	2.529e-3
1p36 deletion syndrome	Schwartz-jampel syndrome	0.06250	1	2.531e-3	3.446e-3