

GeDiPNet Cluster 285 -- 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: 14 · Generated: 2026-10-08 04:32:15

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
greenberg dysplasia	Reynolds syndrome	0.50000	1	6.494e-5	2.317e-4
greenberg dysplasia	Rhizomelic skeletal dysplasia with or without pelger-huet anomaly	0.50000	1	6.494e-5	2.317e-4
greenberg dysplasia	regressive spondylometaphyseal dysplasia	0.50000	1	6.494e-5	2.317e-4
regressive spondylometaphyseal dysplasia	Reynolds syndrome	0.50000	1	6.494e-5	2.317e-4
regressive spondylometaphyseal dysplasia	Rhizomelic skeletal dysplasia with or without pelger-huet anomaly	0.50000	1	6.494e-5	2.317e-4
Reynolds syndrome	Rhizomelic skeletal dysplasia with or without pelger-huet anomaly	0.50000	1	6.494e-5	2.317e-4
greenberg dysplasia	Pelger-huet anomaly	0.33333	1	1.299e-4	3.908e-4
Pelger-huet anomaly	Reynolds syndrome	0.33333	1	1.299e-4	3.908e-4
Pelger-huet anomaly	Rhizomelic skeletal dysplasia with or without pelger-huet anomaly	0.33333	1	1.299e-4	3.908e-4
Pelger-huet anomaly	regressive spondylometaphyseal dysplasia	0.33333	1	1.299e-4	3.908e-4
Congenital retrognathism	greenberg dysplasia	0.14286	1	3.897e-4	8.637e-4
Congenital retrognathism	regressive spondylometaphyseal dysplasia	0.14286	1	3.897e-4	8.637e-4
Congenital retrognathism	Reynolds syndrome	0.14286	1	3.897e-4	8.637e-4
Congenital retrognathism	Rhizomelic skeletal dysplasia with or without pelger-huet anomaly	0.14286	1	3.897e-4	8.637e-4