

GeDiPNet Cluster 335 -- 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: 11 · Generated: 2026-10-08 14:22:53

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
dyskeratosis congenita, autosomal dominant 3	Revesz debuse syndrome	0.50000	1	6.494e-5	2.317e-4
dyskeratosis congenita, autosomal dominant 3	Revesz syndrome	0.50000	1	6.494e-5	2.317e-4
Revesz debuse syndrome	Revesz syndrome	0.50000	1	6.494e-5	2.317e-4
Bone marrow diseases	Revesz debuse syndrome	0.20000	1	2.598e-4	6.484e-4
Bone marrow diseases	Revesz syndrome	0.20000	1	2.598e-4	6.484e-4
Bone marrow diseases	dyskeratosis congenita, autosomal dominant 3	0.20000	1	2.598e-4	6.484e-4
Cerebellar diseases	Mowat-wilson syndrome	0.12500	1	4.546e-4	9.697e-4
Cerebellar diseases	Revesz debuse syndrome	0.12500	1	4.546e-4	9.697e-4
Cerebellar diseases	Revesz syndrome	0.12500	1	4.546e-4	9.697e-4
Cerebellar diseases	dyskeratosis congenita, autosomal dominant 3	0.12500	1	4.546e-4	9.697e-4
Bone marrow diseases	Cerebellar diseases	0.09091	1	1.817e-3	2.675e-3