

GeDiPNet Cluster 475 -- 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: 2026-10-20 03:00:18. · Rows: 14 · Generated: 2026-10-19 23:19:43

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Dc type 4 and 5	Dyskeratosiscongenita, autosomal dominant 4	0.50000	1	6.457e-5	1.985e-4
Dc type 4 and 5	Dyskeratosis congenita, autosomal recessive 5	0.50000	1	6.457e-5	1.985e-4
Dc type 4 and 5	Dyskeratosis congenita, autosomal dominant 4	0.50000	1	6.457e-5	1.985e-4
Dyskeratosis congenita, autosomal dominant 4	Dyskeratosiscongenita, autosomal dominant 4	0.50000	1	6.457e-5	1.985e-4
Dyskeratosis congenita, autosomal dominant 4	Dyskeratosis congenita, autosomal recessive 5	0.50000	1	6.457e-5	1.985e-4
Dyskeratosis congenita, autosomal recessive 5	Dyskeratosiscongenita, autosomal dominant 4	0.50000	1	6.457e-5	1.985e-4
Dc type 4 and 5	X-linked dyskeratosis congenita	0.25000	1	1.937e-4	4.414e-4
Dyskeratosis congenita, autosomal dominant 4	X-linked dyskeratosis congenita	0.25000	1	1.937e-4	4.414e-4
Dyskeratosis congenita, autosomal recessive 5	X-linked dyskeratosis congenita	0.25000	1	1.937e-4	4.414e-4
Dyskeratosiscongenita, autosomal dominant 4	X-linked dyskeratosis congenita	0.25000	1	1.937e-4	4.414e-4
Dc type 4 and 5	Telomere syndrome	0.16667	1	3.228e-4	6.412e-4
Dyskeratosis congenita, autosomal dominant 4	Telomere syndrome	0.16667	1	3.228e-4	6.412e-4
Dyskeratosis congenita, autosomal recessive 5	Telomere syndrome	0.16667	1	3.228e-4	6.412e-4
Dyskeratosiscongenita, autosomal dominant 4	Telomere syndrome	0.16667	1	3.228e-4	6.412e-4