

GeDiPNet Cluster 511 -- 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 03:00:19. · Rows: 10 · Generated: 2026-10-19 23:18:36

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Deafness autosomal dominant type 13	Weissenbacher-zweymueller syndrome	0.50000	1	6.457e-5	1.985e-4
Deafness autosomal dominant type 13	Deafness, autosomal recessive 53	0.50000	1	6.457e-5	1.985e-4
Deafness autosomal dominant type 13	Fibrochondrogenesis 2	0.50000	1	6.457e-5	1.985e-4
Deafness, autosomal recessive 53	Weissenbacher-zweymueller syndrome	0.50000	1	6.457e-5	1.985e-4
Deafness, autosomal recessive 53	Fibrochondrogenesis 2	0.50000	1	6.457e-5	1.985e-4
Fibrochondrogenesis 2	Weissenbacher-zweymueller syndrome	0.50000	1	6.457e-5	1.985e-4
Conductive hearing loss	Deafness autosomal dominant type 13	0.16667	1	3.228e-4	6.412e-4
Conductive hearing loss	Deafness, autosomal recessive 53	0.16667	1	3.228e-4	6.412e-4
Conductive hearing loss	Fibrochondrogenesis 2	0.16667	1	3.228e-4	6.412e-4
Conductive hearing loss	Weissenbacher-zweymueller syndrome	0.16667	1	3.228e-4	6.412e-4