

GeDiPNet Cluster 355 -- 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: 2026-10-09 03:00:12. · Rows: 10 · Generated: 2026-10-08 23:34:29

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
3mc syndrome	Malpuech facial clefting syndrome	0.75000	3	1.644e-12	1.924e-11
3mc syndrome	Craniofacial ulnar renal syndrome	0.60000	3	6.575e-12	7.307e-11
Craniofacial ulnar renal syndrome	Malpuech facial clefting syndrome	0.60000	3	6.575e-12	7.307e-11
3mc syndrome	Carnevale syndrome	0.40000	2	7.592e-8	5.414e-7
Carnevale syndrome	Malpuech facial clefting syndrome	0.40000	2	7.592e-8	5.414e-7
Carnevale syndrome	Craniofacial ulnar renal syndrome	0.33333	2	1.518e-7	1.030e-6
3mc syndrome	Oculopalatosekeletal syndrome	0.28571	2	2.530e-7	1.653e-6
Carnevale syndrome	Oculopalatosekeletal syndrome	0.28571	2	2.530e-7	1.653e-6
Malpuech facial clefting syndrome	Oculopalatosekeletal syndrome	0.28571	2	2.530e-7	1.653e-6
Craniofacial ulnar renal syndrome	Oculopalatosekeletal syndrome	0.25000	2	5.060e-7	3.142e-6