

GeDiPNet Cluster 146 -- 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: 12 · Generated: 2026-10-08 22:42:38

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Burn-mckeown syndrome	Choanal atresia syndrome	0.33333	2	8.436e-8	5.939e-7
Branchial arch abnormalities syndrome	Branchial cleft anomalies	0.50000	1	6.494e-5	2.329e-4
Branchial arch abnormalities syndrome	Vissers-bodmer syndrome	0.25000	1	1.948e-4	5.350e-4
Branchial cleft anomalies	Vissers-bodmer syndrome	0.25000	1	1.948e-4	5.350e-4
holoprosencephaly 12 with or without pancreatic agenesis	Vissers-bodmer syndrome	0.25000	1	1.948e-4	5.350e-4
Branchial arch abnormalities syndrome	Choanal atresia syndrome	0.16667	1	3.247e-4	7.705e-4
Branchial cleft anomalies	Choanal atresia syndrome	0.16667	1	3.247e-4	7.705e-4
Duane-radial ray syndrome	Vissers-bodmer syndrome	0.20000	1	3.896e-4	8.660e-4
Branchial arch abnormalities syndrome	Vein of galen aneurysm	0.14286	1	3.897e-4	8.660e-4
Branchial cleft anomalies	Vein of galen aneurysm	0.14286	1	3.897e-4	8.660e-4
EPHB4-associated vascular malformation spectrum	Vein of galen aneurysm	0.14286	1	3.897e-4	8.660e-4
Choanal atresia syndrome	Vissers-bodmer syndrome	0.12500	1	9.739e-4	1.665e-3