

GeDiPNet Cluster 297 -- 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: 2026-10-07 03:00:12. · Rows: 14 · Generated: 2026-10-07 03:20:20

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital hypothalamic hamartoma syndrome	Curry-jones syndrome	0.50000	1	6.494e-5	2.342e-4
Congenital hypothalamic hamartoma syndrome	Winter shortland temple syndrome	0.50000	1	6.494e-5	2.342e-4
Congenital hypothalamic hamartoma syndrome	mosaic SMO syndrome	0.50000	1	6.494e-5	2.342e-4
Curry-jones syndrome	Winter shortland temple syndrome	0.50000	1	6.494e-5	2.342e-4
Curry-jones syndrome	mosaic SMO syndrome	0.50000	1	6.494e-5	2.342e-4
mosaic SMO syndrome	Winter shortland temple syndrome	0.50000	1	6.494e-5	2.342e-4
Ameloblastoma	Congenital hypothalamic hamartoma syndrome	0.33333	1	1.299e-4	3.898e-4
Ameloblastoma	Curry-jones syndrome	0.33333	1	1.299e-4	3.898e-4
Ameloblastoma	Winter shortland temple syndrome	0.33333	1	1.299e-4	3.898e-4
Ameloblastoma	mosaic SMO syndrome	0.33333	1	1.299e-4	3.898e-4
Congenital hypothalamic hamartoma syndrome	Meningioma	0.04000	1	1.559e-3	2.360e-3
Curry-jones syndrome	Meningioma	0.04000	1	1.559e-3	2.360e-3
Meningioma	mosaic SMO syndrome	0.04000	1	1.559e-3	2.360e-3
Meningioma	Winter shortland temple syndrome	0.04000	1	1.559e-3	2.360e-3