

GeDiPNet Cluster 618 -- 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: 10 · Generated: 2026-10-19 23:19:35

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Combined psap deficiency	Gaucher disease due to saposin C deficiency	0.50000	1	6.457e-5	1.985e-4
Combined psap deficiency	Krabbe disease due to saposin A deficiency	0.50000	1	6.457e-5	1.985e-4
Gaucher disease due to saposin C deficiency	Krabbe disease due to saposin A deficiency	0.50000	1	6.457e-5	1.985e-4
Combined psap deficiency	Combined saposin deficiency	0.33333	1	1.291e-4	3.269e-4
Combined psap deficiency	Krabbe disease	0.33333	1	1.291e-4	3.269e-4
Combined saposin deficiency	Gaucher disease due to saposin C deficiency	0.33333	1	1.291e-4	3.269e-4
Combined saposin deficiency	Krabbe disease due to saposin A deficiency	0.33333	1	1.291e-4	3.269e-4
Gaucher disease due to saposin C deficiency	Krabbe disease	0.33333	1	1.291e-4	3.269e-4
Krabbe disease	Krabbe disease due to saposin A deficiency	0.33333	1	1.291e-4	3.269e-4
Combined saposin deficiency	Krabbe disease	0.25000	1	2.583e-4	5.384e-4