

GeDiPNet Cluster 608 -- 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 23:20:00

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Lissencephaly 2 (norman-roberts type)	Sacroiliac arthritis	0.50000	1	6.457e-5	1.985e-4
Lissencephaly 2 (norman-roberts type)	Sacroiliac joint synovitis	0.50000	1	6.457e-5	1.985e-4
Sacroiliac arthritis	Sacroiliac joint synovitis	0.50000	1	6.457e-5	1.985e-4
Lissencephaly 2 (norman-roberts type)	Lissencephaly with cerebellar hypoplasia	0.33333	1	1.291e-4	3.269e-4
Lissencephaly 2 (norman-roberts type)	Roberts syndrome	0.33333	1	1.291e-4	3.269e-4
Lissencephaly with cerebellar hypoplasia	Sacroiliac arthritis	0.33333	1	1.291e-4	3.269e-4
Lissencephaly with cerebellar hypoplasia	Sacroiliac joint synovitis	0.33333	1	1.291e-4	3.269e-4
Roberts syndrome	Sacroiliac arthritis	0.33333	1	1.291e-4	3.269e-4
Roberts syndrome	Sacroiliac joint synovitis	0.33333	1	1.291e-4	3.269e-4
Lissencephaly with cerebellar hypoplasia	Roberts syndrome	0.25000	1	2.583e-4	5.384e-4