

GeDiPNet Cluster 305 -- 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 17:17:47. · Rows: 7 · Generated: 2026-10-07 20:07:48

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Chondrodysplasia	Rhizomelic chondrodysplasia punctata	0.21053	4	5.079e-12	5.752e-11
alkylglycerone-phosphate synthase deficiency	Rhizomelic chondrodysplasia punctata	0.16667	1	3.247e-4	7.680e-4
glyceronephosphate O-acyltransferase deficiency	Rhizomelic chondrodysplasia punctata	0.16667	1	3.247e-4	7.680e-4
alkylglycerone-phosphate synthase deficiency	Chondrodysplasia	0.05556	1	1.104e-3	1.824e-3
Chondrodysplasia	chondrodysplasia with joint dislocations, gpapp type	0.05556	1	1.104e-3	1.824e-3
Chondrodysplasia	glyceronephosphate O-acyltransferase deficiency	0.05556	1	1.104e-3	1.824e-3
Chondrodysplasia	X-linked dominant chondrodysplasia chassaing-lacombe type	0.05556	1	1.104e-3	1.824e-3