

GeDiPNet Cluster 258 -- 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: 13 · Generated: 2026-10-07 13:27:44

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Digitotalar dysmorphism	Distal arthrogryposis	0.35294	6	4.330e-19	7.842e-18
Digitotalar dysmorphism	Sheldon-hall syndrome	0.71429	5	8.323e-19	1.489e-17
Distal arthrogryposis	Sheldon-hall syndrome	0.29412	5	6.059e-16	9.211e-15
Arthrogryposis	Distal arthrogryposis	0.12000	3	1.511e-7	1.033e-6
Arthrogryposis	Sheldon-hall syndrome	0.13333	2	4.634e-6	2.425e-5
Arthrogryposis	Digitotalar dysmorphism	0.12500	2	6.949e-6	3.530e-5
Congenital finger flexion contractures	Sheldon-hall syndrome	0.16667	1	3.247e-4	7.680e-4
Sheldon-hall syndrome	TPM2-related myopathy	0.16667	1	3.247e-4	7.680e-4
Congenital finger flexion contractures	Digitotalar dysmorphism	0.14286	1	3.897e-4	8.637e-4
Digitotalar dysmorphism	TPM2-related myopathy	0.14286	1	3.897e-4	8.637e-4
Arthrogryposis	childhood-onset nemaline myopathy	0.08333	1	7.144e-4	1.333e-3
Arthrogryposis	TPM2-related myopathy	0.08333	1	7.144e-4	1.333e-3
Congenital finger flexion contractures	Distal arthrogryposis	0.05882	1	1.039e-3	1.737e-3