

GeDiPNet Shared-Gene Disease Pairs

Similarity: Jaccard index on shared curated genes (disease_gdp). Significance: upper-tail hypergeometric test ($P(X \geq k)$), Benjamini-Hochberg FDR-corrected across all tested pairs. Corroborated Shared Genes: count of shared genes backed by ≥ 2 distinct disease_gdp.main_source values combined across both diseases (e.g. GWAS-backed for one, ClinVar-backed for the other still counts); Shared Gene Names lists each gene's own source count in parentheses. Overlap Coefficient: shared genes / MIN(each disease's own total gene count) -- complements the Jaccard-based Similarity Score by showing how much of the SMALLER disease's entire known gene-set is covered by this pair, which Jaccard (dividing by the union) can understate. Shared Cluster: cluster id if both diseases fall in the same label-propagation cluster (scripts/build_disease_clusters.php), blank otherwise. See scripts/build_related.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-11 03:00:14. Filters applied: disease name contains "Syndactyly of the toes" (5 matching pairs). · Rows: 5 · Generated: 2026-10-11 02:16:54

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Bullous pemphigoid	Syndactyly of the toes	1	1	DHCR7 (2)	0.04000	0.50000	2.985e-3	3.949e-3	
Metabolic bone disorder	Syndactyly of the toes	1	0	DHCR7 (1)	0.06250	0.50000	1.818e-3	2.677e-3	
Imperforate anus	Syndactyly of the toes	1	0	CUL9 (1)	0.09091	0.50000	1.169e-3	1.895e-3	
Smith-lemli-opitz syndrome	Syndactyly of the toes	1	1	DHCR7 (7)	0.16667	0.50000	5.195e-4	1.061e-3	
Peripheral pulmonary artery stenosis	Syndactyly of the toes	1	0	CUL9 (1)	0.25000	0.50000	2.598e-4	6.509e-4	