

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 "long chain 3-hydroxyacyl-coa dehydrogenase deficiency" (5 matching pairs). · Rows: 5 · Generated: 2026-10-11 04:12:47

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
long chain 3-hydroxyacyl-coa dehydrogenase deficiency	Osteoarthritis	1	1	HADHA (2)	0.00137	1.00000	4.741e-2	4.920e-2	
long chain 3-hydroxyacyl-coa dehydrogenase deficiency	Myocardial ischemia	1	1	HADHA (3)	0.00255	1.00000	2.539e-2	2.713e-2	
Fatty liver	long chain 3-hydroxyacyl-coa dehydrogenase deficiency	1	1	HADHA (2)	0.01064	1.00000	6.040e-3	7.267e-3	
Fatty liver, alcoholic	long chain 3-hydroxyacyl-coa dehydrogenase deficiency	1	1	HADHA (2)	0.01149	1.00000	5.585e-3	6.785e-3	
long chain 3-hydroxyacyl-coa dehydrogenase deficiency	Trifunctional protein deficiency	1	1	HADHA (3)	0.25000	1.00000	1.948e-4	5.354e-4	