

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: 2026-10-10 03:00:13. Filters applied: disease name contains "Dopamine beta-hydroxylase deficiency" (5 matching pairs). · Rows: 5 · Generated: 2026-10-10 20:16:18

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Dopamine beta-hydroxylase deficiency	Hematologic neoplasms	1	1	HSPA5 (2)	0.14286	0.50000	6.494e-4	1.241e-3	
Dopamine beta-hydroxylase deficiency	Paranoia	1	1	DBH (3)	0.07143	0.50000	1.558e-3	2.376e-3	
Dopamine beta-hydroxylase deficiency	Pheochromocytoma	1	1	DBH (3)	0.05556	0.50000	2.077e-3	2.969e-3	
Asbestosis	Dopamine beta-hydroxylase deficiency	1	1	HSPA5 (2)	0.05263	0.50000	2.207e-3	3.113e-3	
Dopamine beta-hydroxylase deficiency	Spinocerebellar ataxia	1	1	HSPA5 (2)	0.00943	0.50000	1.346e-2	1.499e-2	