

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 "Primary adrenal insufficiency" (5 matching pairs). · Rows: 5 · Generated: 2026-10-10 21:38:46

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Carcinogenesis	Primary adrenal insufficiency	1	1	CUX1 (2)	0.05000	0.50000	2.337e-3	3.243e-3	
Duane retraction syndrome	Primary adrenal insufficiency	1	1	CUX1 (2)	0.07143	0.50000	1.558e-3	2.376e-3	
Keratinocyte carcinoma	Primary adrenal insufficiency	2	0	CUX1 (1), MYL10 (1)	0.02564	1.00000	2.468e-5	1.158e-4	
Angioedema	Primary adrenal insufficiency	2	0	CUX1 (1), MYL10 (1)	0.03448	1.00000	1.346e-5	6.544e-5	
Glucocorticoid deficiency	Primary adrenal insufficiency	2	0	CUX1 (1), MYL10 (1)	0.20000	1.00000	3.037e-7	1.951e-6	