

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: 2026-10-12 03:00:13. Filters applied: disease name contains "Chops syndrome" (5 matching pairs). · Rows: 5 · Generated: 2026-10-12 01:56:57

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Chops syndrome	Delirium, dementia, and cognitive disorders	1	1	AFF4 (3)	0.01754	1.00000	3.637e-3	4.668e-3	
Chops syndrome	Cognition disorder	1	1	AFF4 (3)	0.01818	1.00000	3.507e-3	4.517e-3	
Chops syndrome	Growth disorder	1	1	AFF4 (3)	0.02778	1.00000	2.273e-3	3.191e-3	
Chops syndrome	Cornelia de lange syndrome	1	1	AFF4 (3)	0.09091	1.00000	6.494e-4	1.241e-3	
Chops syndrome	Cognitive impairment-coarse facies-heart defects-obesity-pulmonary involvement-short stature-skeletal dysplasia syndrome	1	1	AFF4 (6)	0.50000	1.00000	6.494e-5	2.331e-4	