

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: 2026-10-09 03:00:13. Filters applied: disease name contains "Complex partial epilepsy" (5 matching pairs). · Rows: 5 · Generated: 2026-10-09 08:24:10

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Complex partial epilepsy	Familial focal epilepsy with variable foci	1	1	SCN3A (4)	0.11111	1.00000	5.195e-4	1.060e-3	
Complex partial epilepsy	Perisylvian syndrome	1	1	SCN3A (2)	0.09091	1.00000	6.494e-4	1.241e-3	
Complex partial epilepsy	Polymicrogyria	1	1	SCN3A (2)	0.03846	1.00000	1.624e-3	2.460e-3	
Complex partial epilepsy	genetic developmental and epileptic encephalopathy	1	1	SCN3A (3)	0.02703	1.00000	2.338e-3	3.242e-3	
Complex partial epilepsy	Developmental delay	1	1	SCN3A (2)	0.02564	1.00000	2.468e-3	3.370e-3	