

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: 2026-10-20 03:00:20. Filters applied: disease name contains "Epileptic encephalopathy, early infantile, 32" (5 matching pairs). · Rows: 5 · Generated: 2026-10-20 03:55:09

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Developmental and epileptic encephalopathy	Epileptic encephalopathy, early infantile, 32	1	1	KCNA2 (7)	0.00431	1.00000	1.491e-2	1.630e-2	
Epileptic encephalopathy, early infantile, 32	Seizures	1	1	KCNA2 (2)	0.00433	1.00000	1.485e-2	1.628e-2	
Epileptic encephalopathy, early infantile, 32	Genetic developmental and epileptic encephalopathy	1	1	KCNA2 (2)	0.02703	1.00000	2.324e-3	3.054e-3	
Epileptic encephalopathy, early infantile, 32	Language development disorders	1	1	KCNA2 (3)	0.03704	1.00000	1.679e-3	2.343e-3	
Epileptic encephalopathy, early infantile, 32	Nizon-isidor syndrome	1	0	KCNA2 (1)	0.33333	1.00000	1.291e-4	3.269e-4	