

GeDiPNet Cluster 339 -- Pairs Within Cluster

Similarity: Jaccard index on shared curated genes (disease_gdp). Significance: upper-tail hypergeometric test, Benjamini-Hochberg FDR-corrected across all tested pairs. See scripts/build_related.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-07 03:00:13. · Rows: 9 · Generated: 2026-10-07 07:14:45

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Progressive myoclonic epilepsy	progressive myoclonus epilepsy	0.36000	9	9.773e-27	2.783e-25
Progressive myoclonic epilepsy	Unverricht-lundborg syndrome	0.12000	3	3.327e-9	3.012e-8
progressive myoclonus epilepsy	Unverricht-lundborg syndrome	0.18182	2	9.108e-7	5.670e-6
Myoclonus-renal failure syndrome	Unverricht-lundborg syndrome	0.25000	1	1.948e-4	5.278e-4
Myoclonus-renal failure syndrome	progressive myoclonus epilepsy	0.10000	1	5.845e-4	1.135e-3
Progressive myoclonic epilepsy with renal failure	progressive myoclonus epilepsy	0.10000	1	5.845e-4	1.135e-3
Myoclonus-renal failure syndrome	Progressive myoclonic epilepsy	0.04000	1	1.559e-3	2.360e-3
Progressive myoclonic epilepsy	Progressive myoclonic epilepsy with renal failure	0.04000	1	1.559e-3	2.360e-3
Lafora disease	Progressive myoclonic epilepsy	0.03846	1	3.115e-3	4.088e-3