

GeDiPNet Cluster 188 -- 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: 2026-10-09 03:00:13. · Rows: 12 · Generated: 2026-10-09 09:24:04

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Polycystic kidney disease	Polycystic liver disease	0.18033	11	1.033e-22	2.206e-21
autosomal dominant polycystic kidney disease	Polycystic kidney disease	0.14286	7	1.811e-18	3.185e-17
autosomal dominant polycystic kidney disease	Polycystic liver disease	0.06897	2	4.462e-5	2.028e-4
autosomal dominant polycystic kidney disease	renal-hepatic-pancreatic dysplasia 2	0.12500	1	4.546e-4	9.721e-4
ALG9-associated autosomal dominant polycystic kidney disease	Polycystic liver disease	0.04167	1	1.494e-3	2.313e-3
polycystic kidney disease 3 with or without polycystic liver disease	Polycystic liver disease	0.04167	1	1.494e-3	2.313e-3
Polycystic liver disease	polycystic liver disease 1	0.04167	1	1.494e-3	2.313e-3
Polycystic liver disease	SEC61B-related polycystic liver disease	0.04167	1	1.494e-3	2.313e-3
ALG9-associated autosomal dominant polycystic kidney disease	Polycystic kidney disease	0.02041	1	3.117e-3	4.095e-3
Polycystic kidney disease	polycystic kidney disease 3 with or without polycystic liver disease	0.02041	1	3.117e-3	4.095e-3
Polycystic kidney disease	polycystic liver disease 1	0.02041	1	3.117e-3	4.095e-3
Polycystic kidney disease	renal-hepatic-pancreatic dysplasia 2	0.02041	1	3.117e-3	4.095e-3