

GeDiPNet Cluster 435 -- 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: 2026-10-20 03:00:19. · Rows: 11 · Generated: 2026-10-19 23:19:56

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Polycystic kidney disease	Polycystic liver disease	0.17460	11	1.600e-22	2.532e-21
Autosomal dominant polycystic kidney disease	Polycystic kidney disease	0.15686	8	6.550e-21	9.880e-20
Autosomal dominant polycystic kidney disease	Polycystic liver disease	0.10345	3	1.594e-7	1.084e-6
Polycystic kidney disease 3 with or without polycystic liver disease	Polycystic liver disease	0.08333	2	2.110e-6	1.233e-5
Polycystic liver disease	Polycystic liver disease 1	0.08333	2	2.110e-6	1.233e-5
Polycystic kidney disease	Polycystic kidney disease 3 with or without polycystic liver disease	0.03922	2	1.021e-5	5.522e-5
Autosomal dominant polycystic kidney disease	Gillessen-kaesbach-nishimura syndrome	0.11111	1	5.165e-4	9.125e-4
Autosomal dominant polycystic kidney disease	Polycystic kidney disease 3 with or without polycystic liver disease	0.10000	1	1.033e-3	1.574e-3
Gillessen-kaesbach-nishimura syndrome	Polycystic liver disease	0.04167	1	1.485e-3	2.122e-3
Gillessen-kaesbach-nishimura syndrome	Polycystic kidney disease	0.01961	1	3.228e-3	4.073e-3
Polycystic kidney disease	Polycystic liver disease 1	0.01923	1	6.446e-3	7.555e-3