

GeDiPNet Cluster 326 -- 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 03:00:12. · Rows: 12 · Generated: 2026-10-07 04:42:45

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Coronary artery vasospasm	Coronary vasospasm	0.60000	3	6.575e-12	7.736e-11
Capillary leak syndrome	Coronary artery vasospasm	0.20000	1	3.896e-4	8.521e-4
Capillary leak syndrome	Coronary vasospasm	0.16667	1	5.195e-4	1.041e-3
Coronary artery vasospasm	Resistant hypertension	0.16667	1	5.844e-4	1.135e-3
Coronary artery vasospasm	Intestinal perforation	0.16667	1	5.844e-4	1.135e-3
Intestinal perforation	Resistant hypertension	0.16667	1	5.844e-4	1.135e-3
Coronary vasospasm	Resistant hypertension	0.14286	1	7.792e-4	1.386e-3
Coronary vasospasm	Intestinal perforation	0.14286	1	7.792e-4	1.386e-3
Resistant hypertension	Ventricular ectopy	0.14286	1	7.792e-4	1.386e-3
Coronary artery vasospasm	Ventricular ectopy	0.14286	1	7.792e-4	1.386e-3
Intestinal perforation	Ventricular ectopy	0.14286	1	7.792e-4	1.386e-3
Coronary vasospasm	Ventricular ectopy	0.12500	1	1.039e-3	1.720e-3