

GeDiPNet Cluster 360 -- 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: 6 · Generated: 2026-10-09 04:52:07

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hemolytic uremic syndrome	Mesangiocapillary glomerulonephritis	0.23529	8	1.775e-22	3.760e-21
C3 glomerulonephritis	Mesangiocapillary glomerulonephritis	0.54545	6	4.238e-20	8.112e-19
C3 glomerulonephritis	Hemolytic uremic syndrome	0.20000	7	8.392e-19	1.495e-17
atypical hemolytic-uremic syndrome with B factor anomaly	C3 glomerulonephritis	0.11111	1	5.195e-4	1.060e-3
atypical hemolytic-uremic syndrome with B factor anomaly	Mesangiocapillary glomerulonephritis	0.11111	1	5.195e-4	1.060e-3
Hemolytic uremic syndrome	hemolytic uremic syndrome, atypical, 8, with rhizomelic short stature	0.02941	1	2.143e-3	3.043e-3