

GeDiPNet Cluster 614 -- 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: 8 · Generated: 2026-10-19 23:20:03

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Combined immunodeficiency, enteropathy spectrum	Gastrointestinal defects and immunodeficiency syndrome	0.75000	3	1.615e-12	1.707e-11
Intestinal atresia, inflammatory bowel disease and immunodeficiency	Multiple intestinal atresia	0.50000	1	6.457e-5	1.985e-4
Combined immunodeficiency, enteropathy spectrum	Multiple intestinal atresia	0.25000	1	1.937e-4	4.414e-4
Combined immunodeficiency, enteropathy spectrum	Intestinal atresia, inflammatory bowel disease and immunodeficiency	0.25000	1	1.937e-4	4.414e-4
Gastrointestinal defects and immunodeficiency syndrome	Multiple intestinal atresia	0.25000	1	1.937e-4	4.414e-4
Gastrointestinal defects and immunodeficiency syndrome	Intestinal atresia, inflammatory bowel disease and immunodeficiency	0.25000	1	1.937e-4	4.414e-4
Combined factor v and factor viii deficiency	Combined immunodeficiency, enteropathy spectrum	0.20000	1	3.874e-4	7.329e-4
Combined factor v and factor viii deficiency	Gastrointestinal defects and immunodeficiency syndrome	0.20000	1	3.874e-4	7.329e-4