

GeDiPNet Cluster 275 -- 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 17:17:47. · Rows: 12 · Generated: 2026-10-07 13:49:30

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital secretory diarrhea	Congenital sodium diarrhea	0.40000	2	5.062e-8	3.718e-7
Congenital chronic diarrhea with protein-losing enteropathy	Congenital diarrhea	0.33333	2	8.436e-8	5.953e-7
Chronic infantile diarrhea due to guanylate cyclase 2c overactivity	Duodenal atresia	0.50000	1	6.494e-5	2.317e-4
Chronic infantile diarrhea due to guanylate cyclase 2c overactivity	Congenital sodium diarrhea	0.33333	1	1.299e-4	3.908e-4
Congenital sodium diarrhea	Duodenal atresia	0.33333	1	1.299e-4	3.908e-4
Chronic infantile diarrhea due to guanylate cyclase 2c overactivity	Congenital secretory diarrhea	0.20000	1	2.598e-4	6.484e-4
Congenital secretory diarrhea	Duodenal atresia	0.20000	1	2.598e-4	6.484e-4
Chronic infantile diarrhea due to guanylate cyclase 2c overactivity	Congenital diarrhea	0.16667	1	3.247e-4	7.680e-4
Congenital diarrhea	Duodenal atresia	0.16667	1	3.247e-4	7.680e-4
Congenital diarrhea	Congenital sodium diarrhea	0.14286	1	6.494e-4	1.239e-3
Congenital chloride diarrhea	Congenital secretory diarrhea	0.12500	1	1.039e-3	1.737e-3
Congenital chloride diarrhea	Congenital diarrhea	0.11111	1	1.298e-3	2.060e-3