

GeDiPNet Cluster 198 -- 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: 10 · Generated: 2026-10-07 16:05:01

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Intracellular cobalamin metabolism disorder	Methylmalonic acidemia	0.23333	7	7.844e-19	1.408e-17
Cobalamin c disease	Methylmalonic acidemia	0.20690	6	1.119e-16	1.798e-15
Cobalamin c disease	Intracellular cobalamin metabolism disorder	0.41667	5	3.669e-16	5.689e-15
Intracellular cobalamin metabolism disorder	methylmalonic aciduria and homocystinuria	0.20000	2	3.037e-7	1.958e-6
Methylmalonic acidemia	methylmalonic aciduria and homocystinuria	0.07143	2	2.961e-6	1.605e-5
Cobalamin c disease	methylmalonic acidemia with homocystinuria, type cblJ	0.12500	1	4.546e-4	9.697e-4
Intracellular cobalamin metabolism disorder	methylmalonic acidemia with homocystinuria, type cblJ	0.10000	1	5.845e-4	1.154e-3
Alpha-mannosidosis	Methylmalonic acidemia	0.03571	1	1.753e-3	2.607e-3
Methylmalonic acidemia	methylmalonic acidemia due to methylmalonyl-CoA epimerase deficiency	0.03571	1	1.753e-3	2.607e-3
Methylmalonic acidemia	methylmalonic acidemia with homocystinuria, type cblJ	0.03571	1	1.753e-3	2.607e-3