

GeDiPNet Cluster 460 -- 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: 12 · Generated: 2026-10-19 23:19:44

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Gm2 gangliosidosis	Tay-sachs disease	0.40000	2	7.504e-8	5.341e-7
Beta-hexosaminidase a deficiency	Hex a pseudodeficiency	0.50000	1	6.457e-5	1.985e-4
Beta-hexosaminidase a deficiency	Gm2-gangliosidosis, several forms, tay-sachs disease	0.50000	1	6.457e-5	1.985e-4
Gm2-gangliosidosis, several forms, tay-sachs disease	Hex a pseudodeficiency	0.50000	1	6.457e-5	1.985e-4
Beta-hexosaminidase a deficiency	Gm2 gangliosidosis	0.25000	1	1.937e-4	4.414e-4
Beta-hexosaminidase a deficiency	Tay-sachs disease	0.25000	1	1.937e-4	4.414e-4
Gm2 gangliosidosis	Hex a pseudodeficiency	0.25000	1	1.937e-4	4.414e-4
Gm2 gangliosidosis	Gm2-gangliosidosis, several forms, tay-sachs disease	0.25000	1	1.937e-4	4.414e-4
Gm2 gangliosidosis	Hexosaminidase activator deficiency	0.25000	1	1.937e-4	4.414e-4
Gm2-gangliosidosis, several forms, tay-sachs disease	Tay-sachs disease	0.25000	1	1.937e-4	4.414e-4
Hex a pseudodeficiency	Tay-sachs disease	0.25000	1	1.937e-4	4.414e-4
Hexosaminidase activator deficiency	Tay-sachs disease	0.25000	1	1.937e-4	4.414e-4