

GeDiPNet Cluster 388 -- 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: 2026-10-08 03:00:13. · Rows: 10 · Generated: 2026-10-08 14:22:06

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hereditary hyperekplexia	Hyperekplexia	0.83333	5	1.387e-19	2.579e-18
Hereditary hyperekplexia	Hyperexplexia hereditary	0.66667	4	2.135e-15	3.133e-14
Hyperekplexia	Hyperexplexia hereditary	0.66667	4	2.135e-15	3.133e-14
Retinitis pigmentosa, juvenile cataract, short stature, intellectual disability syndrome	sulfite oxidase deficiency due to molybdenum cofactor deficiency type C	0.33333	1	1.299e-4	3.908e-4
Hyperexplexia hereditary	sulfite oxidase deficiency due to molybdenum cofactor deficiency type C	0.20000	1	2.598e-4	6.484e-4
Hereditary hyperekplexia	sulfite oxidase deficiency due to molybdenum cofactor deficiency type C	0.16667	1	3.247e-4	7.680e-4
Hyperekplexia	sulfite oxidase deficiency due to molybdenum cofactor deficiency type C	0.16667	1	3.247e-4	7.680e-4
Hyperexplexia hereditary	Retinitis pigmentosa, juvenile cataract, short stature, intellectual disability syndrome	0.16667	1	5.195e-4	1.058e-3
Hereditary hyperekplexia	Retinitis pigmentosa, juvenile cataract, short stature, intellectual disability syndrome	0.14286	1	6.494e-4	1.239e-3
Hyperekplexia	Retinitis pigmentosa, juvenile cataract, short stature, intellectual disability syndrome	0.14286	1	6.494e-4	1.239e-3