

GeDiPNet Cluster 466 -- 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: 15 · Generated: 2026-10-19 23:20:01

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Dystonia 4, hypomyelinating leukodystrophy 6	Whispering dysphonia	0.50000	1	6.457e-5	1.985e-4
Dystonia 4, hypomyelinating leukodystrophy 6	Neurologic disorder	0.50000	1	6.457e-5	1.985e-4
Dystonia 4, hypomyelinating leukodystrophy 6	Hypomyelination with atrophy of the basal ganglia and cerebellum	0.50000	1	6.457e-5	1.985e-4
Dystonia 4, hypomyelinating leukodystrophy 6	Dystonia 4, torsion, autosomal dominant	0.50000	1	6.457e-5	1.985e-4
Dystonia 4, torsion, autosomal dominant	Whispering dysphonia	0.50000	1	6.457e-5	1.985e-4
Dystonia 4, torsion, autosomal dominant	Neurologic disorder	0.50000	1	6.457e-5	1.985e-4
Dystonia 4, torsion, autosomal dominant	Hypomyelination with atrophy of the basal ganglia and cerebellum	0.50000	1	6.457e-5	1.985e-4
Hypomyelination with atrophy of the basal ganglia and cerebellum	Whispering dysphonia	0.50000	1	6.457e-5	1.985e-4
Hypomyelination with atrophy of the basal ganglia and cerebellum	Neurologic disorder	0.50000	1	6.457e-5	1.985e-4
Neurologic disorder	Whispering dysphonia	0.50000	1	6.457e-5	1.985e-4
Brown tendon sheath syndrome	Whispering dysphonia	0.25000	1	1.937e-4	4.414e-4
Brown tendon sheath syndrome	Neurologic disorder	0.25000	1	1.937e-4	4.414e-4
Brown tendon sheath syndrome	Hypomyelination with atrophy of the basal ganglia and cerebellum	0.25000	1	1.937e-4	4.414e-4
Brown tendon sheath syndrome	Dystonia 4, torsion, autosomal dominant	0.25000	1	1.937e-4	4.414e-4
Brown tendon sheath syndrome	Dystonia 4, hypomyelinating leukodystrophy 6	0.25000	1	1.937e-4	4.414e-4