

GeDiPNet Cluster 456 -- 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: 11 · Generated: 2026-10-19 23:20:18

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Mitochondrial complex v (atp synthase) deficiency, nuclear type 2	Mitochondrial encephalocardiomyopathy	0.50000	1	6.457e-5	1.985e-4
Mitochondrial complex v (atp synthase) deficiency, nuclear type 2	Oligohydramnios	0.50000	1	6.457e-5	1.985e-4
Mitochondrial encephalocardiomyopathy	Oligohydramnios	0.50000	1	6.457e-5	1.985e-4
Mitochondrial complex v (atp synthase) deficiency, nuclear type	Mitochondrial encephalocardiomyopathy	0.33333	1	1.291e-4	3.269e-4
Mitochondrial complex v (atp synthase) deficiency, nuclear type	Oligohydramnios	0.33333	1	1.291e-4	3.269e-4
Mitochondrial complex v (atp synthase) deficiency, nuclear type	Mitochondrial complex v (atp synthase) deficiency, nuclear type 2	0.33333	1	1.291e-4	3.269e-4
Mitochondrial complex v (atp synthase) deficiency, nuclear type	Mitochondrial complex v (atp synthase) deficiency, nuclear type 1	0.33333	1	1.291e-4	3.269e-4
Intrauterine growth restriction	Mitochondrial encephalocardiomyopathy	0.20000	1	2.583e-4	5.384e-4
Intrauterine growth restriction	Mitochondrial complex v (atp synthase) deficiency, nuclear type 2	0.20000	1	2.583e-4	5.384e-4
Intrauterine growth restriction	Oligohydramnios	0.20000	1	2.583e-4	5.384e-4
Intrauterine growth restriction	Mitochondrial complex v (atp synthase) deficiency, nuclear type	0.16667	1	5.165e-4	9.125e-4