

GeDiPNet Cluster 331 -- 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 03:00:13. · Rows: 12 · Generated: 2026-10-07 07:08:21

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital central hypoventilation syndrome	Haddad syndrome	0.90000	9	7.475e-33	2.661e-31
Central hypoventilation syndrome	Haddad syndrome	0.30000	3	1.381e-10	1.432e-9
Central hypoventilation syndrome	Congenital central hypoventilation syndrome	0.30000	3	1.381e-10	1.432e-9
Haddad syndrome	Hirschsprung disease	0.05479	4	4.058e-8	3.206e-7
Congenital central hypoventilation syndrome	Hirschsprung disease	0.05479	4	4.058e-8	3.206e-7
Biliary atresia	Haddad syndrome	0.05128	2	1.400e-4	4.194e-4
Biliary atresia	Congenital central hypoventilation syndrome	0.05128	2	1.400e-4	4.194e-4
Central hypoventilation syndrome	central hypoventilation syndrome, congenital, 1, with or without Hirschsprung disease	0.25000	1	1.948e-4	5.278e-4
central hypoventilation syndrome, congenital, 1, with or without Hirschsprung disease	Haddad syndrome	0.10000	1	5.845e-4	1.135e-3
central hypoventilation syndrome, congenital, 1, with or without Hirschsprung disease	Congenital central hypoventilation syndrome	0.10000	1	5.845e-4	1.135e-3
central hypoventilation syndrome, congenital, 1, with or without Hirschsprung disease	Hirschsprung disease	0.01471	1	4.351e-3	5.453e-3
Central hypoventilation syndrome	Hirschsprung disease	0.01429	1	1.300e-2	1.453e-2