

GeDiPNet Cluster 556 -- 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: 9 · Generated: 2026-10-19 23:20:19

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hyperparathyroidism 2 with jaw tumors	Parathyroid neoplasm	0.50000	1	6.457e-5	1.985e-4
Hyperparathyroidism 2 with jaw tumors	Hyperparathyroidism, familial primary	0.50000	1	6.457e-5	1.985e-4
Hyperparathyroidism, familial primary	Parathyroid neoplasm	0.50000	1	6.457e-5	1.985e-4
Cervical adenosarcoma	Hyperparathyroidism 2 with jaw tumors	0.14286	1	3.874e-4	7.329e-4
Cervical adenosarcoma	Hyperparathyroidism, familial primary	0.14286	1	3.874e-4	7.329e-4
Cervical adenosarcoma	Parathyroid neoplasm	0.14286	1	3.874e-4	7.329e-4
Apocrine adenocarcinoma	Hyperparathyroidism 2 with jaw tumors	0.11111	1	5.165e-4	9.125e-4
Apocrine adenocarcinoma	Hyperparathyroidism, familial primary	0.11111	1	5.165e-4	9.125e-4
Apocrine adenocarcinoma	Parathyroid neoplasm	0.11111	1	5.165e-4	9.125e-4