

GeDiPNet Cluster 570 -- 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 03:00:17. · Rows: 10 · Generated: 2026-10-19 23:20:20

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Salivary gland basal cell adenocarcinoma	Salivary gland pleomorphic adenoma	0.55000	11	3.141e-33	6.621e-32
Salivary gland basal cell adenocarcinoma	Salivary gland carcinoma ex pleomorphic adenoma	0.28205	11	5.161e-27	9.432e-26
Salivary gland carcinoma ex pleomorphic adenoma	Salivary gland pleomorphic adenoma	0.27778	10	8.422e-26	1.468e-24
Salivary gland mucoepidermoid carcinoma	Salivary gland pleomorphic adenoma	0.27027	10	1.208e-25	2.098e-24
Salivary gland carcinoma ex pleomorphic adenoma	Salivary gland mucoepidermoid carcinoma	0.22222	12	1.973e-25	3.416e-24
Parotid gland pleomorphic adenoma	Salivary gland pleomorphic adenoma	0.46667	7	3.240e-22	5.081e-21
Parotid gland pleomorphic adenoma	Salivary gland basal cell adenocarcinoma	0.36842	7	3.671e-21	5.589e-20
Parotid gland pleomorphic adenoma	Salivary gland carcinoma ex pleomorphic adenoma	0.20588	7	6.348e-19	8.970e-18
Salivary gland basal cell adenocarcinoma	Salivary gland mucoepidermoid carcinoma	0.18605	8	4.065e-18	5.572e-17
Parotid gland pleomorphic adenoma	Salivary gland mucoepidermoid carcinoma	0.16667	6	1.614e-15	1.998e-14