

GeDiPNet Cluster 390 -- 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 17:17:47. · Rows: 9 · Generated: 2026-10-07 14:24:58

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Dentin dysplasia	Dentinogenesis imperfecta	0.70000	14	1.698e-43	7.723e-42
Amelogenesis imperfecta	Dentinogenesis imperfecta	0.44828	13	6.648e-36	2.497e-34
Amelogenesis imperfecta	Dentin dysplasia	0.43333	13	2.824e-35	1.029e-33
Amelogenesis imperfecta	Hypomaturation amelogenesis imperfecta	0.26923	7	1.182e-20	2.337e-19
Dentinogenesis imperfecta	Hypomaturation amelogenesis imperfecta	0.33333	6	3.029e-18	5.289e-17
Dentin dysplasia	Hypomaturation amelogenesis imperfecta	0.31579	6	4.681e-18	8.120e-17
Dentinogenesis imperfecta	Enamel-renal syndrome	0.05882	1	1.039e-3	1.737e-3
Dentin dysplasia	Enamel-renal syndrome	0.05556	1	1.104e-3	1.824e-3
Amelogenesis imperfecta	Enamel-renal syndrome	0.03846	1	1.624e-3	2.459e-3