

GeDiPNet Cluster 319 -- 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 03:00:12. · Rows: 8 · Generated: 2026-10-07 09:18:07

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Bone marrow failure syndromes	Congenital bone marrow failure syndrome	0.30000	3	1.381e-10	1.432e-9
Aplasia and myelodysplasia	Congenital bone marrow failure syndrome	0.25000	1	1.948e-4	5.278e-4
bone marrow failure syndrome 3	Congenital bone marrow failure syndrome	0.25000	1	1.948e-4	5.278e-4
bone marrow failure syndrome 3	Shwachman-diamond syndrome	0.12500	1	4.546e-4	9.545e-4
Shwachman-diamond syndrome	shwachman-diamond syndrome 2	0.12500	1	4.546e-4	9.545e-4
Aplasia and myelodysplasia	Bone marrow failure syndromes	0.10000	1	5.845e-4	1.135e-3
bone marrow failure syndrome 3	Bone marrow failure syndromes	0.10000	1	5.845e-4	1.135e-3
Congenital bone marrow failure syndrome	Shwachman-diamond syndrome	0.10000	1	1.363e-3	2.129e-3