

GeDiPNet Cluster 378 -- 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: 2026-10-09 03:00:12. · Rows: 6 · Generated: 2026-10-09 04:55:28

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	1.353e-9
Aplasia and myelodysplasia	Congenital bone marrow failure syndrome	0.25000	1	1.948e-4	5.350e-4
Congenital bone marrow failure syndrome	Growth hormone insensitivity syndrome with immune dysregulation	0.20000	1	3.896e-4	8.660e-4
Aplasia and myelodysplasia	Bone marrow failure syndromes	0.10000	1	5.845e-4	1.156e-3
Bone marrow failure syndromes	Congenital progressive bone marrow failure-b-cell immunodeficiency-skeletal dysplasia syndrome	0.10000	1	5.845e-4	1.156e-3
Bone marrow failure syndromes	Growth hormone insensitivity syndrome with immune dysregulation	0.09091	1	1.169e-3	1.895e-3