

GeDiPNet Cluster 84 -- 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 03:00:13. · Rows: 19 · Generated: 2026-10-09 09:23:20

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Xeroderma pigmentosum	Xeroderma pigmentosum-cockayne syndrome	0.38462	5	1.099e-16	1.759e-15
Cerebrooculofacioskeletal syndrome	Xeroderma pigmentosum	0.26667	4	3.169e-12	3.614e-11
Cerebrooculofacioskeletal syndrome	Xeroderma pigmentosum-cockayne syndrome	0.33333	3	3.287e-10	3.122e-9
Trichothiodystrophy	Xeroderma pigmentosum-cockayne syndrome	0.13333	2	4.634e-6	2.418e-5
Trichothiodystrophy	Xeroderma pigmentosum	0.09091	2	3.050e-5	1.411e-4
xeroderma pigmentosum group B	Xeroderma pigmentosum-cockayne syndrome	0.16667	1	3.247e-4	7.705e-4
xeroderma pigmentosum group D	Xeroderma pigmentosum-cockayne syndrome	0.16667	1	3.247e-4	7.705e-4
xeroderma pigmentosum group G	Xeroderma pigmentosum-cockayne syndrome	0.16667	1	3.247e-4	7.705e-4
Cerebrooculofacioskeletal syndrome	xeroderma pigmentosum group D	0.14286	1	3.897e-4	8.660e-4
Cerebrooculofacioskeletal syndrome	xeroderma pigmentosum group G	0.14286	1	3.897e-4	8.660e-4
Trichorrhexis nodosa syndrome	Trichothiodystrophy	0.08333	1	7.144e-4	1.335e-3
Trichothiodystrophy	xeroderma pigmentosum group B	0.08333	1	7.144e-4	1.335e-3
Amish brittle hair brain syndrome	Trichothiodystrophy	0.08333	1	7.144e-4	1.335e-3
Trichothiodystrophy	xeroderma pigmentosum group D	0.08333	1	7.144e-4	1.335e-3
Xeroderma pigmentosum	xeroderma pigmentosum variant type	0.07692	1	7.793e-4	1.405e-3
Xeroderma pigmentosum	xeroderma pigmentosum group G	0.07692	1	7.793e-4	1.405e-3
Xeroderma pigmentosum	xeroderma pigmentosum group E	0.07692	1	7.793e-4	1.405e-3
Xeroderma pigmentosum	xeroderma pigmentosum group C	0.07692	1	7.793e-4	1.405e-3
Xeroderma pigmentosum	xeroderma pigmentosum group B	0.07692	1	7.793e-4	1.405e-3