

GeDiPNet Cluster 80 -- 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: 2026-10-08 03:00:13. · Rows: 19 · Generated: 2026-10-08 03:36:03

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	1.766e-15
Cerebrooculofacioskeletal syndrome	Xeroderma pigmentosum	0.26667	4	3.169e-12	3.632e-11
Cerebrooculofacioskeletal syndrome	Xeroderma pigmentosum-cockayne syndrome	0.33333	3	3.287e-10	3.138e-9
Trichothiodystrophy	Xeroderma pigmentosum-cockayne syndrome	0.13333	2	4.634e-6	2.425e-5
Trichothiodystrophy	Xeroderma pigmentosum	0.09091	2	3.050e-5	1.415e-4
xeroderma pigmentosum group B	Xeroderma pigmentosum-cockayne syndrome	0.16667	1	3.247e-4	7.680e-4
xeroderma pigmentosum group D	Xeroderma pigmentosum-cockayne syndrome	0.16667	1	3.247e-4	7.680e-4
xeroderma pigmentosum group G	Xeroderma pigmentosum-cockayne syndrome	0.16667	1	3.247e-4	7.680e-4
Cerebrooculofacioskeletal syndrome	xeroderma pigmentosum group D	0.14286	1	3.897e-4	8.637e-4
Cerebrooculofacioskeletal syndrome	xeroderma pigmentosum group G	0.14286	1	3.897e-4	8.637e-4
Trichorrhexis nodosa syndrome	Trichothiodystrophy	0.08333	1	7.144e-4	1.333e-3
Trichothiodystrophy	xeroderma pigmentosum group B	0.08333	1	7.144e-4	1.333e-3
Amish brittle hair brain syndrome	Trichothiodystrophy	0.08333	1	7.144e-4	1.333e-3
Trichothiodystrophy	xeroderma pigmentosum group D	0.08333	1	7.144e-4	1.333e-3
Xeroderma pigmentosum	xeroderma pigmentosum variant type	0.07692	1	7.793e-4	1.404e-3
Xeroderma pigmentosum	xeroderma pigmentosum group G	0.07692	1	7.793e-4	1.404e-3
Xeroderma pigmentosum	xeroderma pigmentosum group E	0.07692	1	7.793e-4	1.404e-3
Xeroderma pigmentosum	xeroderma pigmentosum group C	0.07692	1	7.793e-4	1.404e-3
Xeroderma pigmentosum	xeroderma pigmentosum group B	0.07692	1	7.793e-4	1.404e-3