

GeDiPNet Cluster 132 -- 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: 2026-10-10 03:00:13. · Rows: 16 · Generated: 2026-10-10 06:59:24

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Hereditary breast and ovarian cancer syndrome	Hereditary breast cancer	0.29730	11	6.085e-28	1.721e-26
Hereditary breast and ovarian cancer syndrome	Hereditary breast-ovarian cancer syndrome	0.31250	5	4.166e-16	6.403e-15
Fanconi anemia	Hereditary breast cancer	0.12500	8	7.923e-15	1.120e-13
Fanconi anemia	Hereditary breast and ovarian cancer syndrome	0.12245	6	8.685e-13	1.055e-11
Fanconi anemia	Hereditary breast-ovarian cancer syndrome	0.09756	4	1.753e-10	1.706e-9
Hereditary breast-ovarian cancer syndrome	RAD51D-related cancer predisposition	0.16667	1	3.247e-4	7.709e-4
BARD1-related cancer predisposition	Hereditary breast and ovarian cancer syndrome	0.06250	1	9.742e-4	1.665e-3
Hereditary breast and ovarian cancer syndrome	nijmegen breakage syndrome	0.06250	1	9.742e-4	1.665e-3
Hereditary breast and ovarian cancer syndrome	RAD51D-related cancer predisposition	0.06250	1	9.742e-4	1.665e-3
BARD1-related cancer predisposition	Hereditary breast cancer	0.03030	1	2.078e-3	2.969e-3
fanconi anemia complementation group p	Hereditary breast cancer	0.03030	1	2.078e-3	2.969e-3
Fanconi anemia complementation group U	Hereditary breast cancer	0.03030	1	2.078e-3	2.969e-3
Hereditary breast cancer	nijmegen breakage syndrome	0.03030	1	2.078e-3	2.969e-3
Fanconi anemia	fanconi anemia complementation group f	0.02500	1	2.533e-3	3.446e-3
Fanconi anemia	fanconi anemia complementation group p	0.02500	1	2.533e-3	3.446e-3
Fanconi anemia	Fanconi anemia complementation group U	0.02500	1	2.533e-3	3.446e-3