

GeDiPNet Cluster 145 -- 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: 18 · Generated: 2026-10-08 09:05:21

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Cavernous malformations of cns	Cerebral cavernous malformation	0.36364	4	4.483e-13	5.626e-12
Congenital cerebral aneurysm	Congenital malformation of cerebral vessels	0.66667	2	8.436e-9	6.837e-8
Congenital cerebral aneurysm	Developmental venous anomaly	0.40000	2	5.062e-8	3.718e-7
Congenital malformation of cerebral vessels	Developmental venous anomaly	0.40000	2	5.062e-8	3.718e-7
Cerebral cavernous malformation	Congenital cerebral aneurysm	0.28571	2	1.265e-7	8.736e-7
Cerebral cavernous malformation	Congenital malformation of cerebral vessels	0.28571	2	1.265e-7	8.736e-7
Developmental venous anomaly	Parkes weber syndrome	0.33333	2	1.518e-7	1.033e-6
Cavernous malformations of cns	Congenital cerebral aneurysm	0.22222	2	2.362e-7	1.556e-6
Cavernous malformations of cns	Congenital malformation of cerebral vessels	0.22222	2	2.362e-7	1.556e-6
Hereditary hemorrhagic telangiectasia	Parkes weber syndrome	0.22222	2	5.313e-7	3.282e-6
Angiokeratoma	Cavernous malformations of cns	0.20000	2	7.084e-7	4.282e-6
Cerebral cavernous malformation	Developmental venous anomaly	0.22222	2	7.590e-7	4.550e-6
Developmental venous anomaly	Hereditary hemorrhagic telangiectasia	0.20000	2	1.062e-6	6.229e-6
Cavernous malformations of cns	Developmental venous anomaly	0.18182	2	1.416e-6	8.118e-6
Hydrops fetalis	Parkes weber syndrome	0.11111	2	3.035e-6	1.636e-5
Developmental venous anomaly	Hydrops fetalis	0.10526	2	6.066e-6	3.107e-5
Hereditary hemorrhagic telangiectasia	Hydrops fetalis	0.09091	2	2.119e-5	1.005e-4
Angiokeratoma	Cerebral cavernous malformation	0.11111	1	1.169e-3	1.894e-3