

GeDiPNet Cluster 65 -- Member Diseases

Cluster grouping: label propagation over disease_related's top-5-per-disease shared-gene similarity graph. Cluster size: 14 diseases. Sorted by connections within the cluster (most central first). Connections/partners count deduplicated disease pairs; significant = FDR q 0.05. See scripts/build_disease_clusters.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-07 03:15:01. · Rows: 14 · Generated: 2026-10-07 13:28:28

Disease	Connections in Cluster	Significant Partners (q<0.05)	Curated Genes
Capillary malformation	6	6	3
Capillary malformation-arteriovenous malformation	6	6	8
Klippel-trenaunay syndrome	6	6	7
Anastomosing haemangioma	5	5	3
Congenital hemangioma	5	5	2
Parkes weber syndrome	5	5	3
Port-wine stain	5	5	2
Cutis marmorata telangiectatica congenita	4	4	2
Hereditary hemorrhagic telangiectasia	4	4	7
Arteriovenous malformations	3	3	2
Congenital lipomatous overgrowth, vascular malformations, and epidermal nevi	3	3	3
Hydrops fetalis	3	3	16
Cerebrofacial arteriovenous metameric syndrome	1	1	1
telangiectasia, hereditary hemorrhagic, type 5	0	0	0