

GeDiPNet Cluster 287 -- Member Diseases

Cluster grouping: label propagation over disease_related's top-5-per-disease shared-gene similarity graph. Cluster size: 6 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: 2026-10-09 03:15:01. · Rows: 6 · Generated: 2026-10-09 13:16:02

Disease	Connections in Cluster	Significant Partners (q<0.05)	Curated Genes
Common arterial trunk with aortic dominance	5	5	2
Common arterial trunk with pulmonary dominance and interrupted aortic arch	5	5	2
Chromosome 22q11.2 microduplication syndrome	4	4	1
Conotruncal anomaly face syndrome	4	4	1
Velocardiofacial syndrome	4	4	5
GATA6-related congenital heart disease with or without pancreatic agenesis or neonatal diabetes	2	2	1