

GeDiPNet Cluster 473 -- 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: 2026-10-20 03:00:19. · Rows: 11 · Generated: 2026-10-19 23:20:56

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital heart defects, multiple types, 1, x-linked	VACTERL association, X-linked	0.50000	1	6.457e-5	1.985e-4
Congenital heart defects, multiple types, 1, x-linked	Heterotaxy, visceral, 1, x-linked	0.50000	1	6.457e-5	1.985e-4
Heterotaxy, visceral, 1, x-linked	VACTERL association, X-linked	0.50000	1	6.457e-5	1.985e-4
Congenital heart defects, multiple types, 1, x-linked	Congenital heart disease with heterotaxy syndrome	0.33333	1	1.291e-4	3.269e-4
Congenital heart disease with heterotaxy syndrome	VACTERL association, X-linked	0.33333	1	1.291e-4	3.269e-4
Congenital heart disease with heterotaxy syndrome	Heterotaxy, visceral, 1, x-linked	0.33333	1	1.291e-4	3.269e-4
Congenital heart defects, multiple types, 1, x-linked	Dextrocardia	0.12500	1	4.520e-4	8.260e-4
Dextrocardia	VACTERL association, X-linked	0.12500	1	4.520e-4	8.260e-4
Dextrocardia	Heterotaxy, visceral, 1, x-linked	0.12500	1	4.520e-4	8.260e-4
Dextrocardia	Gut malrotation	0.12500	1	4.520e-4	8.260e-4
Congenital heart disease with heterotaxy syndrome	Dextrocardia	0.11111	1	9.038e-4	1.411e-3