

GeDiPNet Cluster 245 -- 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: 10 · Generated: 2026-10-08 00:35:47

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Congenital malformation syndromes predominantly involving limbs	Rubinstein-taybi syndrome	0.22222	2	2.362e-7	1.556e-6
Congenital malformation syndromes predominantly involving limbs	Vacterl association	0.11765	2	1.012e-6	5.965e-6
Rubinstein-taybi syndrome	Vacterl association	0.08696	2	2.824e-5	1.314e-4
Congenital cleft hand	Congenital malformation syndromes predominantly involving limbs	0.33333	1	1.299e-4	3.908e-4
Congenital malformation syndromes predominantly involving limbs	Waardenburg anophthalmia syndrome	0.33333	1	1.299e-4	3.908e-4
Chromosome 16p13.3 deletion syndrome	Rubinstein-taybi syndrome	0.11111	1	5.195e-4	1.058e-3
Congenital cleft hand	Rubinstein-taybi syndrome	0.11111	1	5.195e-4	1.058e-3
Rubinstein-taybi syndrome	Waardenburg anophthalmia syndrome	0.11111	1	5.195e-4	1.058e-3
fanconi anemia complementation group I	Vacterl association	0.05882	1	1.039e-3	1.737e-3
Vacterl association	Waardenburg anophthalmia syndrome	0.05882	1	1.039e-3	1.737e-3