

GeDiPNet Cluster 635 -- 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: 10 · Generated: 2026-10-19 23:20:06

Disease A	Disease B	Similarity Score	Shared Genes	P-value	FDR Q-value
Pulmonary hypertension, familial primary, 1, with or without hht	Right bundle branch block	0.50000	1	6.457e-5	1.985e-4
Pulmonary hypertension, familial primary, 1, with or without hht	Right cardiac ventricular dilatation	0.50000	1	6.457e-5	1.985e-4
Pulmonary hypertension, familial primary, 1, with or without hht	Pulmonary venoocclusive disease 1	0.50000	1	6.457e-5	1.985e-4
Pulmonary hypertension, familial primary, 1, with or without hht	Pulmonary hypertension, primary, fenfluramine or dexfenfluramine-associated	0.50000	1	6.457e-5	1.985e-4
Pulmonary hypertension, primary, fenfluramine or dexfenfluramine-associated	Right bundle branch block	0.50000	1	6.457e-5	1.985e-4
Pulmonary hypertension, primary, fenfluramine or dexfenfluramine-associated	Right cardiac ventricular dilatation	0.50000	1	6.457e-5	1.985e-4
Pulmonary hypertension, primary, fenfluramine or dexfenfluramine-associated	Pulmonary venoocclusive disease 1	0.50000	1	6.457e-5	1.985e-4
Pulmonary venoocclusive disease 1	Right bundle branch block	0.50000	1	6.457e-5	1.985e-4
Pulmonary venoocclusive disease 1	Right cardiac ventricular dilatation	0.50000	1	6.457e-5	1.985e-4
Right bundle branch block	Right cardiac ventricular dilatation	0.50000	1	6.457e-5	1.985e-4