

GeDiPNet Shared-Gene Disease Pairs

Similarity: Jaccard index on shared curated genes (disease_gdp). Significance: upper-tail hypergeometric test ($P(X \geq k)$), Benjamini-Hochberg FDR-corrected across all tested pairs. Corroborated Shared Genes: count of shared genes backed by ≥ 2 distinct disease_gdp.main_source values combined across both diseases (e.g. GWAS-backed for one, ClinVar-backed for the other still counts); Shared Gene Names lists each gene's own source count in parentheses. Overlap Coefficient: shared genes / MIN(each disease's own total gene count) -- complements the Jaccard-based Similarity Score by showing how much of the SMALLER disease's entire known gene-set is covered by this pair, which Jaccard (dividing by the union) can understate. Shared Cluster: cluster id if both diseases fall in the same label-propagation cluster (scripts/build_disease_clusters.php), blank otherwise. See scripts/build_related.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-20 03:00:20. Filters applied: disease name contains "Primary myelofibrosis" (5 matching pairs). · Rows: 5 · Generated: 2026-10-20 02:12:25

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Budd-chiari syndrome	Primary myelofibrosis	2	2	JAK2 (5), CALR (2)	0.40000	1.00000	5.003e-8	3.625e-7	
Primary myelofibrosis	Thrombocythemia	2	2	JAK2 (5), CALR (4)	0.28571	1.00000	1.251e-7	8.632e-7	
Acute panmyelosis with myelofibrosis	Primary myelofibrosis	1	0	JAK2 (1)	0.33333	1.00000	1.291e-4	3.269e-4	
Myelofibrosis, somatic	Primary myelofibrosis	1	0	CALR (1)	0.25000	0.50000	2.583e-4	5.384e-4	
Hepatic vein thrombosis	Primary myelofibrosis	1	1	JAK2 (2)	0.20000	0.50000	3.874e-4	7.329e-4	