

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: 2026-10-09 03:00:13. Filters applied: disease name contains "Hypochromic anemia" (5 matching pairs). · Rows: 5 · Generated: 2026-10-09 20:49:58

Disease A	Disease B	Shared Genes	Corroborated Shared Genes	Shared Gene Names	Similarity Score	Overlap Coefficient	P-value	FDR Q-value	Shared Cluster
Hypochromic anemia	Microcytic anemia	1	1	SLC11A2 (5)	0.20000	1.00000	2.598e-4	6.505e-4	98
Hypochromic anemia	Hypochromic microcytic anemia	1	1	SLC11A2 (3)	0.16667	1.00000	3.247e-4	7.705e-4	98
Hypochromic anemia	Iron deficiency anemia	1	1	SLC11A2 (2)	0.09091	1.00000	6.494e-4	1.241e-3	98
Hemochromatosis	Hypochromic anemia	1	1	SLC11A2 (2)	0.06667	1.00000	9.092e-4	1.575e-3	
Anemia	Hypochromic anemia	1	0	SLC11A2 (1)	0.01176	1.00000	5.455e-3	6.629e-3	