

GeDiPNet Cluster 322 -- 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: 2026-10-09 03:00:12. · Rows: 8 · Generated: 2026-10-08 23:36:08

| Disease A                       | Disease B                         | Similarity Score | Shared Genes | P-value  | FDR Q-value |
|---------------------------------|-----------------------------------|------------------|--------------|----------|-------------|
| Hyper-immunoglobulin d syndrome | Mevalonate kinase deficiency      | 0.40000          | 2            | 5.062e-8 | 3.709e-7    |
| Arthralgia                      | Hyper-igd syndrome                | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Arthralgia                      | Bone marrow neoplasms             | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Hyper-igd syndrome              | Mevalonate kinase deficiency      | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Arthralgia                      | Mevalonate kinase deficiency      | 0.25000          | 1            | 2.598e-4 | 6.505e-4    |
| Hyper-igd syndrome              | Hyper-immunoglobulin d syndrome   | 0.20000          | 1            | 2.598e-4 | 6.505e-4    |
| Hyper-immunoglobulin d syndrome | methylmalonic aciduria, cblb type | 0.20000          | 1            | 2.598e-4 | 6.505e-4    |
| Arthralgia                      | Hyper-immunoglobulin d syndrome   | 0.16667          | 1            | 5.195e-4 | 1.060e-3    |