

GeDiPNet Cluster 179 -- 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: 21 · Generated: 2026-10-09 11:40:02

| Disease A                                          | Disease B                                          | Similarity Score | Shared Genes | P-value  | FDR Q-value |
|----------------------------------------------------|----------------------------------------------------|------------------|--------------|----------|-------------|
| Transthyretin amyloid cardiomyopathy               | Wild-type transthyretin-related amyloidosis        | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| Dystransthyretinemic euthyroidal hyperthyroxinemia | Senile systemic amyloidosis                        | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| Dystransthyretinemic euthyroidal hyperthyroxinemia | Transthyretin amyloid cardiomyopathy               | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| Dystransthyretinemic euthyroidal hyperthyroxinemia | Wild-type transthyretin-related amyloidosis        | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| Dystransthyretinemic euthyroidal hyperthyroxinemia | obsolete hereditary ATTR amyloidosis               | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| Senile systemic amyloidosis                        | Wild-type transthyretin-related amyloidosis        | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| Senile systemic amyloidosis                        | Transthyretin amyloid cardiomyopathy               | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| obsolete hereditary ATTR amyloidosis               | Wild-type transthyretin-related amyloidosis        | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| obsolete hereditary ATTR amyloidosis               | Transthyretin amyloid cardiomyopathy               | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| obsolete hereditary ATTR amyloidosis               | Senile systemic amyloidosis                        | 0.50000          | 1            | 6.494e-5 | 2.329e-4    |
| Amyloid polyneuropathy                             | Senile systemic amyloidosis                        | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Amyloid polyneuropathy                             | Wild-type transthyretin-related amyloidosis        | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Amyloid polyneuropathy                             | obsolete hereditary ATTR amyloidosis               | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Amyloid polyneuropathy                             | Transthyretin amyloid cardiomyopathy               | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Amyloid polyneuropathy                             | Dystransthyretinemic euthyroidal hyperthyroxinemia | 0.33333          | 1            | 1.299e-4 | 3.929e-4    |
| Hyperthyroxinemia                                  | obsolete hereditary ATTR amyloidosis               | 0.25000          | 1            | 1.948e-4 | 5.350e-4    |
| Hyperthyroxinemia                                  | Wild-type transthyretin-related amyloidosis        | 0.25000          | 1            | 1.948e-4 | 5.350e-4    |
| Hyperthyroxinemia                                  | Transthyretin amyloid cardiomyopathy               | 0.25000          | 1            | 1.948e-4 | 5.350e-4    |
| Hyperthyroxinemia                                  | Senile systemic amyloidosis                        | 0.25000          | 1            | 1.948e-4 | 5.350e-4    |
| Dystransthyretinemic euthyroidal hyperthyroxinemia | Hyperthyroxinemia                                  | 0.25000          | 1            | 1.948e-4 | 5.350e-4    |
| Hyperthyroxinemia                                  | Thyroid hormone metabolism disorder                | 0.20000          | 1            | 3.896e-4 | 8.660e-4    |