

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  $\geq 2$  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: 2026-10-11 03:00:14. Filters applied: disease name contains "long qt syndrome 5" (5 matching pairs). · Rows: 5 · Generated: 2026-10-11 17:31:17

| Disease A                                       | Disease B                 | Shared Genes | Corroborated Shared Genes | Shared Gene Names | Similarity Score | Overlap Coefficient | P-value  | FDR Q-value | Shared Cluster |
|-------------------------------------------------|---------------------------|--------------|---------------------------|-------------------|------------------|---------------------|----------|-------------|----------------|
| Arrhythmogenic right ventricular cardiomyopathy | long qt syndrome 5        | 1            | 1                         | KCNE1 (2)         | 0.01887          | 1.00000             | 3.377e-3 | 4.374e-3    |                |
| Hereditary atrial fibrillation                  | long qt syndrome 5        | 1            | 1                         | KCNE1 (2)         | 0.04000          | 1.00000             | 1.559e-3 | 2.376e-3    |                |
| Congenital hearing disorder                     | long qt syndrome 5        | 1            | 1                         | KCNE1 (2)         | 0.14286          | 1.00000             | 3.897e-4 | 8.665e-4    |                |
| long qt syndrome 5                              | Long qt syndrome, digenic | 1            | 1                         | KCNE1 (2)         | 0.16667          | 1.00000             | 3.247e-4 | 7.709e-4    |                |
| Jervell and lange-nielsen syndrome              | long qt syndrome 5        | 1            | 1                         | KCNE1 (7)         | 0.33333          | 1.00000             | 1.299e-4 | 3.931e-4    |                |