

| | R-HSA-8982491 | R-HSA-1474290 | R-HSA-373755 | R-HSA-422475 | immune_only |
|---|---------------|---------------|--------------|--------------|-------------|
| Pairwise path length in reference tree | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Jaccard distance over extended gene set | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Dijkstra BMA PPI | 0.0007 | 0.0 | 0.0 | 0.0 | 0.0 |
| WM distance over summary W2V | 0.0052 | 0.0 | 0.0 | 0.0 | NaN |
| WM distance over gene symbols W2V | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Cosine distance over over summary W2V | 0.0027 | 0.0 | 0.0005 | 0.0 | NaN |
| Cosine distance over over NCBI summary W2V | 0.0005 | 0.0 | 0.0 | 0.0 | 0.0 |
| Cosine distance over gene symbols W2V | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Cosine distance GO MF description W2V | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Cosine distance GO CC description W2V | 0.0 | 0.0 | 0.0 | 0.0 | 0.0476 |
| Cosine distance GO BP description W2V | 0.2682 | 0.0 | 0.0 | 0.0 | 0.0 |
| GO-distance (go_type=MF, measure=Wang, combine=BMA) | 0.0 | 0.0 | 0.0 | 0.0 | 0.0004 |
| GO-distance (go_type=CC, measure=Wang, combine=BMA) | 0.0004 | 0.0 | 0.0 | 0.0 | 0.0029 |
| GO-distance (go_type=BP, measure=Wang, combine=BMA) | 0.0223 | 0.0 | 0.0 | 0.0 | 0.0 |
| Overlap distance over genes | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Overlap distance over gene traits | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Minkowski distance (p=2) over genes | 0.0061 | 0.2457 | 0.2867 | 0.0 | 0.0 |
| Minkowski distance (p=2) over gene traits | 0.9516 | 0.0009 | 0.6076 | 0.0 | 0.0 |
| Minkowski distance (p=2) over gene trait frequency | 0.1526 | 0.5101 | 0.0015 | 0.0 | 0.0 |
| Minkowski distance (p=1) over genes | 0.0061 | 0.2457 | 0.2867 | 0.0 | 0.0 |
| Minkowski distance (p=1) over gene traits | 0.9516 | 0.0009 | 0.6076 | 0.0 | 0.0 |
| Minkowski distance (p=1) over gene trait frequency | 0.408 | 0.1588 | 0.024 | 0.0 | 0.0 |
| Kappa distance over genes | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Kappa distance over gene traits | 0.0 | 0.0 | 0.0 | 0.0 | 0.0133 |
| Jaccard distance over genes | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Jaccard distance over gene traits | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Cosine distance over gene trait frequency | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Random (uniform, (0,1)) | 0.7903 | 0.555 | 0.4201 | 0.3821 | 0.5835 |