



| Term | PValue | Sample |
|--|----------|--------|
| cytoplasmic translation | 0.000000 | HCT116 |
| translation | 0.000000 | HCT116 |
| regulation of translation | 0.006151 | HCT116 |
| negative regulation of translation | 0.006525 | HCT116 |
| mitochondrial electron transport, NADH to ubiquinone | 0.054501 | HCT116 |
| positive regulation of protein-containing complex assembly | 0.061396 | HCT116 |
| intracellular iron ion homeostasis | 0.078424 | HCT116 |
| proton motive force-driven mitochondrial ATP synthesis | 0.078424 | HCT116 |
| mitochondrial respiratory chain complex I assembly | 0.078424 | HCT116 |
| aerobic respiration | 0.079549 | HCT116 |
| cytoplasmic translation | 0.014674 | A549 |
| translation | 0.030397 | A549 |
| translation | 0.000217 | HepG2 |
| intracellular iron ion homeostasis | 0.000733 | HepG2 |
| cytoplasmic translation | 0.001528 | HepG2 |
| intracellular sequestering of iron ion | 0.004250 | HepG2 |
| translational elongation | 0.011496 | HepG2 |
| iron ion transport | 0.016300 | HepG2 |
| ribosomal small subunit biogenesis | 0.045261 | HepG2 |