Gene ontology in darkgrey module mitochondrial translation (GO:0032543) mitochondrial translational elongation (GO:0070125) % Genes in set translational termination (GO:0006415) 0.40 translational elongation (GO:0006414) 0.50 mitochondrial translational termination (GO:0070126) 0.60 ubiquitin-dependent protein catabolic process (GO:0006511) proteasome-mediated ubiquitin-dependent protein catabolic process (GO:0043161) 12.5 mRNA splicing, via spliceosome (GO:0000398) 10.0 DNA metabolic process (GO:0006259) mRNA processing (GO:0006397) 100 200 300

**Combined Score**