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