Gene ontology in dimgrey module protein export from nucleus (GO:0006611)

nucleotide-excision repair (GO:0006289) termination of RNA polymerase I transcription (GO:0006363)

regulation of proteasomal protein catabolic process (GO:0061136)

regulation of gene silencing by miRNA (GO:0060964) rRNA base methylation (GO:0070475)

aerobic electron transport chain (GO:0019646)

transcription initiation from RNA polymerase I promoter (GO:0006361) positive regulation of telomere maintenance (GO:0032206)

200 400 **Combined Score**

% Genes

0.08

in set

0.0 mitochondrial ATP synthesis coupled electron transport (GO:0042775) 0