Gene ontology in white module mRNA processing (GO:0006397) RNA processing (GO:0006396) % Genes in set mRNA splicing, via spliceosome (GO:0000398) 0.10 negative regulation of cellular macromolecule biosynthetic process (GO:2000113) 0.20 RNA splicing (GO:0008380) 0.30 nuclear export (GO:0051168) RNA splicing, via transesterification reactions with bulged adenosine as nucleophile (GO:0000377) negative regulation of nucleic acid-templated transcription (GO:1903507) positive regulation of nuclear-transcribed mRNA poly(A) tail shortening (GO:0060213) regulation of nuclear-transcribed mRNA poly(A) tail shortening (GO:0060211) 100 200 **Combined Score**