```
   Eukaryotic GeneMark.hmm version 3.54
   Sequence name: seq.fna
   FASTA defline: >NT_033777.3:21378977-21381970 Drosophila melanogaster chromosome 3R
   Sequence length: 2994 bp
   G+C content: 48.26%
   Matrices file:
   /home/genemark/parameters/eukaryotic/Drosophila_melanogaster/drosophila_melanogaster.mod
   Sun Mar 24 01:52:05 2019


   Predicted genes/exons

   Gene Exon Strand Exon          Exon Range          Exon        Start/End
    #    #         Type                              Length       Frame


    1    1    +    Initial         271      455        185         1 2 - -
    1    2    +    Terminal        915     1134        220         3 3 - -

    2    1    +    Initial        1202     1305        104         1 2 - -
    2    2    +    Internal       1358     1968        611         3 1 - -
    2    3    +    Terminal       2397     2851        455         2 3 - -
```