```
   Eukaryotic GeneMark.hmm version 3.54
   Sequence name: seq.fna
   FASTA defline: >NT_037436.4:c18846267-18828632 Drosophila melanogaster chromosome 3L
   Sequence length: 17636 bp
   G+C content: 37.42%
   Matrices file:
   /home/genemark/parameters/eukaryotic/Drosophila_melanogaster/drosophila_melanogaster.mod
   Sun Mar 24 01:47:47 2019


   Predicted genes/exons
```

Gene #	Exon #	Strand	Exon Type	Exon Range		Exon Length	Start/End Frame	
1	1	+	Initial	4183	4292	110	1 2	- -
1	2	+	Internal	4348	4459	112	3 3	- -
1	3	+	Internal	14284	14321	38	1 2	- -
1	4	+	Internal	14647	14773	127	3 3	- -
1	5	+	Internal	14841	14957	117	1 3	- -
1	6	+	Internal	15027	15175	149	1 2	- -
1	7	+	Internal	15240	15428	189	3 2	- -
1	8	+	Internal	15534	15992	459	3 2	- -
1	9	+	Internal	16311	16689	379	3 3	- -
1	10	+	Terminal	16755	17045	291	1 3	- -