

Eukaryotic GeneMark.hmm version 3.54

Sequence name: seq.fna

FASTA defline: >NT_033777.3:c23141906-23128169 Drosophila melanogaster chromosome 3R

Sequence length: 13738 bp

G+C content: 45.57%

Matrices file:

/home/genemark/parameters/eukaryotic/Drosophila_melanogaster/drosophila_melanogaster.mod

Sun Mar 24 01:46:36 2019

Predicted genes/exons

Gene #	Exon #	Strand	Exon Type	Exon Range		Exon Length	Start/End Frame
1	1	+	Initial	386	865	480	1 3 - -
1	2	+	Internal	8644	8905	262	1 1 - -
1	3	+	Internal	9294	9412	119	2 3 - -
1	4	+	Terminal	10630	10716	87	1 3 - -
2	1	+	Initial	11792	11798	7	1 1 - -
2	2	+	Terminal	12496	13169	674	2 3 - -