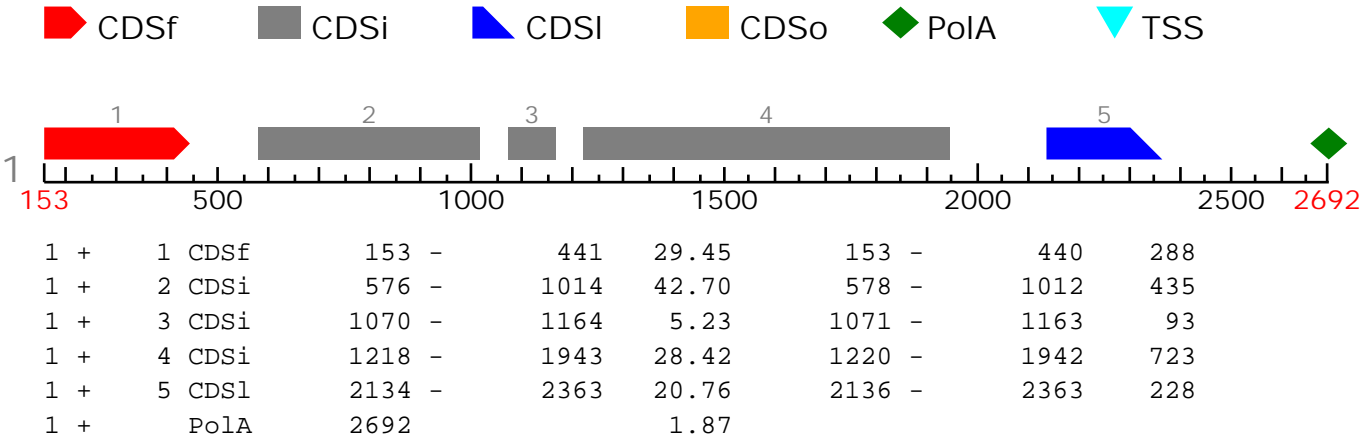


FGENESH 2.6 Prediction of potential genes in *Trichinella_spiralis* genomic DNA
Seq name: gb|CM027592.1| Dmelanogaster07
Length of sequence: 2902
Number of predicted genes 1: in +chain 1, in -chain 0.
Number of predicted exons 5: in +chain 5, in -chain 0.
Positions of predicted genes and exons: Variant 1 from 1, Score:117.105518



Predicted protein(s):

>FGENESH:[mRNA] 1 5 exon (s) 153 - 2363 1779 bp, chain +
ATGAAATATTGCAAATTTTGCTGCAAAGCTGTGACTGGAGTAAAATTAATTCACGTGCCG
AAGTGTGCTATTAAGAGAAAATTGTGGGAGCAGAGCCTTGGGTGCAGCCTTGGTGAAAAC
TCCCAAATTTGTGATACCCACTTTAATGATTTCGCAGTGAAGGCTGCACCTGCAAAAGGT
CAGACATTTAAAAGGAGGCGACTCAACGCAGATGCCGTACCTAGTAAAGTGATAGAGCCT
GAACCAGAAAAGAAAAGAAGGCTATAACCAGTGGGAGTACACAAACAGAAGCAGCAGCTG
AGGGAGTCTCAACAGTTGGAGGAGTCTCTACGCAAAATCTTCACGGACACGCAGATACGG
ATACTGAAGAATGGTGGACAAAGAGCTACGTTCAATTCCGACGACATTTCTACAGCTATT
TGTCTCCACACCGCAGGCCCTCGAGCGTATAACCATCTGTACAAAAGGATTTCCCTTTG
CCCAGTCGTACGACTTTGTACAGATGGTTATCAGATGTGGACATAAAAAGAGGATGTTTG
GATGTGGTCATAGACCTAATGGACAGTGATGGAGTTGATGACGCCGACAAGCTTTGCGTA
CTCGCTTTTCGACGAGATGAAGGTCGCTGCTGCCTTCGAGTATGACAGCTCTGCTGATATT
GTTTACGAGCCAAGCGACTATGTCCAATGGCTATTGTTCGTGGTCTAAAAAATCGTGGA
AGCAGCCAGAACTGCATAGGAAAGGATATTTAGTAGTTGCTATTGTATCCGATTTAGGT
ACCGGAAACCAAAAGCTATGGACAGAGCTCGGTATATCAGAATCAAAAACCTGGTTTAGC
CATCCTGCAGATGACCATTTAAAGATTTTCGTTTTTTTCGGATACGCCACATTTAATTAAG
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GATTTTCAGTTCTAAAGAAAACGAAAATAAAAGTAAAGATGATGAGCCTGTGCGATGATGAG
CCTGTGCGATGAGATGTTAAGCAATATAGATTTTACCGAAATGGATGAGTTGACGGAGGAT
GCGATGGAATATATCGCGGGCTATGTCATTAATAAATTGA

>FGENESH: 1 5 exon (s) 153 - 2363 592 aa, chain +

MKYCKFCCKAVTGVKLIHVPKCAIKRKLWEQSLGCSLGENSQICDTHFNDSQWKAAPAKG
QTFKRRRLNADAVPSKVIEPEPEKKAIPVGVHKQKQQLRESQQLLEESLRKIFTDTQIR
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DVVIDLMDSDGVDDADKLCVLAFFDEMKVAAAFEYDSSADIVYEPSDYVQLAIVRGLKNRG
SSQKLHRKGYLVVAIVSDLGTGNQKLWTELGISESKTWFSHPADDHLKIFVFSDTPHLIK
LVRNHYVDSGLTINGKKLTKKTIQEALHLCNKSDLSILFKINENHINVRSLAKQKVKLAT
QLFSNTTASSIRRCYSLGYDIENATETADFFKLMNDWFDIFNSKLSTSNIECSQPYGKQ
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