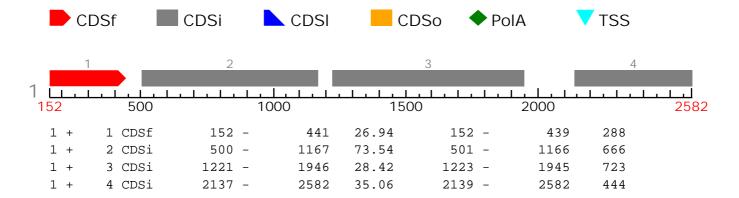
FGENESH 2.6 Prediction of potential genes in Trichinella\_spiralis genomic DNA

Seg name: gb|CM010516.1| Dmelanogaster33

Length of sequence: 2906

Number of predicted genes 1: in +chain 1, in -chain 0. Number of predicted exons 4: in +chain 4, in -chain 0.

Positions of predicted genes and exons: Variant 1 from 1, Score: 145.876233



## Predicted protein(s):

>FGENESH: [mRNA] 1 4 exon (s) 152 - 2582 2130 bp, chain + AAGTGTGCTATTAAGAGAAAATTGTGGGAGCAGAGCCTTGGGTGCAGCCTTGGTGAAAAC TCCCAAATTTGTGATACCCACTTTAATGATTCGCAGTGGAAGGCTGCACCTGCAAAAGGT CAGACATTTAAAAGGAGGCGACTCAACGCAGATGCCGTACCTAGTAAAGTGATAGAGCCT GAACCAGAAAAGATAAAAGAAGGCTATACCAGTGGGAGTACACAAACAGAGTCCTGTTCA TTGTTTAATGAAAATAAGAGCTTGAGGGAAAAAATTCGTACTTTGGAGTACGAAATGCGT CGTTTAGAGCAGCTGAGGGAGTCTCAACAGTTGGAGGAGTCTCTACGCAAAATCTTC ACGGACACGCAGATACGGATACTGAAGAATGGTGGACAAAGAGCTACGTTCAATTCCGAC GACATTTCTACAGCTATTTGTCTCCACACCGCAGGCCCTCGAGCGTATAACCATCTGTAC AAAAAAGGATTTCCTTTGCCCAGTCGTACGACTTTGTACAGATGGTTATCAGATGTGGAC ATAAAAAGAGGATGTTTGGATGTGGTCATAGACCTAATGGACAGTGATGGAGTTGATGAC GACAGCTCTGCTGATATTGTTTACGAGCCAAGCGACTATGTCCAACTGGCTATTGTTCGT GGTCTAAAAAAATCGTGGAAGCAGCCAGTTTTTTTCGATTTTAATACCCGAATGGACCCG GATACTCTTAACAATATTAAGGAAACTGCATAGGAAAGGATATTTAGTAGTTGCTATT GTATCCGATTTAGGTACCGGAAACCAAAAGCTATGGACAGAGCTCGGTATATCAGAATCA AAAACCTGGTTTAGCCATCCTGCAGATGACCATTTAAAGATTTTCGTTTTTTCGGATACG CCACATTTAATTAAGTTAGTCCGTAACCACTATGTGGATTCCGGATTAACAATAAATGGG AAAAAATTAACAAAAAAAACAATTCAGGAGGCACTTCATCTTTGCAACAAGTCCGATCTG GTTAAATTGGCTACCCAGCTGTTTTCGAATACCACCGCTAGCTCGATCAGACGCTGCTAT TCATTGGGGTATGACATTGAAAATGCCACCGAAACTGCGGACTTCTTCAAATTGATGAAT GATTGGTTCGACATTTTTAATTCTAAATTGTCCACATCCAATTGCATTGAGTGCTCGCAA CCTTATGGCAAGCAGTTGGATATACAGAATGATATTTTGAATCGAATGTCGGAAATTATG CGAACAGGAATTCTGGATAAACCCAAAAGGCTCCCATTTCAAAAAGGTATCATTGTGAAT AATGCTTCGCTTGATGGCTTGTATAAATATTTTGCAAGAAAACTTCAGTATGCAATACATA GGTGGACAATTCGACCATCCCACTCCACTGCAGTTTAAGTATAGGTTAAGAAAATATATA ATAGCCAGGAATACAGAAATGTTAAGAAATTCGGGAAATATCGAAGAGGACAACTCTGAA
AGCTGGCTTAATTTAGATTTCAGTTCTAAAGAAAACGAAAATAAAAGTAAAGATGATGAG
CCTGTCGATGATGAGCCTGTCGATGAGATGTTAAGCAATATAGATTTCACCGAAATGGAT
GAGTTGACGGAGGATGCGATGGAATATATCGCGGGGCTATGTCATTAAAAAATTGAGAATC
AGTGACAAAGTAAAAGAAAATTTGACATTTACATACGTCGACGAGGTGTCTCACGGCGGA
CTTATTAAGCCGTCCGAAAAATTTCAAGAGAAAGTTAAAAGAGCTAGAATGTATTTTTTG
CATTATACAAATAATAATTTTGAAATTACAAATAATGTAAAGGAAAAATTAATATA
GCAGCGCGAAACGTCGATGTTGATAAACAA

>FGENESH: 1 4 exon (s) 152 - 2582 710 aa, chain +
MKYCKFCCKAVTGVKLIHVPKCAIKRKLWEQSLGCSLGENSQICDTHFNDSQWKAAPAKG
QTFKRRRLNADAVPSKVIEPEPEKIKEGYTSGSTQTESCSLFNENKSLREKIRTLEYEMR
RLEQQLRESQQLEESLRKIFTDTQIRILKNGGQRATFNSDDISTAICLHTAGPRAYNHLY
KKGFPLPSRTTLYRWLSDVDIKRGCLDVVIDLMDSDGVDDADKLCVLAFDEMKVAAAFEY
DSSADIVYEPSDYVQLAIVRGLKKSWKQPVFFDFNTRMDPDTLNNILRKLHRKGYLVVAI
VSDLGTGNQKLWTELGISESKTWFSHPADDHLKIFVFSDTPHLIKLVRNHYVDSGLTING
KKLTKKTIQEALHLCNKSDLSILFKINENHINVRSLAKQKVKLATQLFSNTTASSIRRCY
SLGYDIENATETADFFKLMNDWFDIFNSKLSTSNCIECSQPYGKQLDIQNDILNRMSEIM
RTGILDKPKRLPFQKGIIVNNASLDGLYKYLQENFSMQYILTSRLNQDIVEHFFGSMRSR
GGQFDHPTPLQFKYRLRKYIIARNTEMLRNSGNIEEDNSESWLNLDFSSKENENKSKDDE
PVDDEPVDEMLSNIDFTEMDELTEDAMEYIAGYVIKKLRISDKVKENLTFTYVDEVSHGG
LIKPSEKFQEKLKELECIFLHYTNNNNFEITNNVKEKLILAARNVDVDKQ