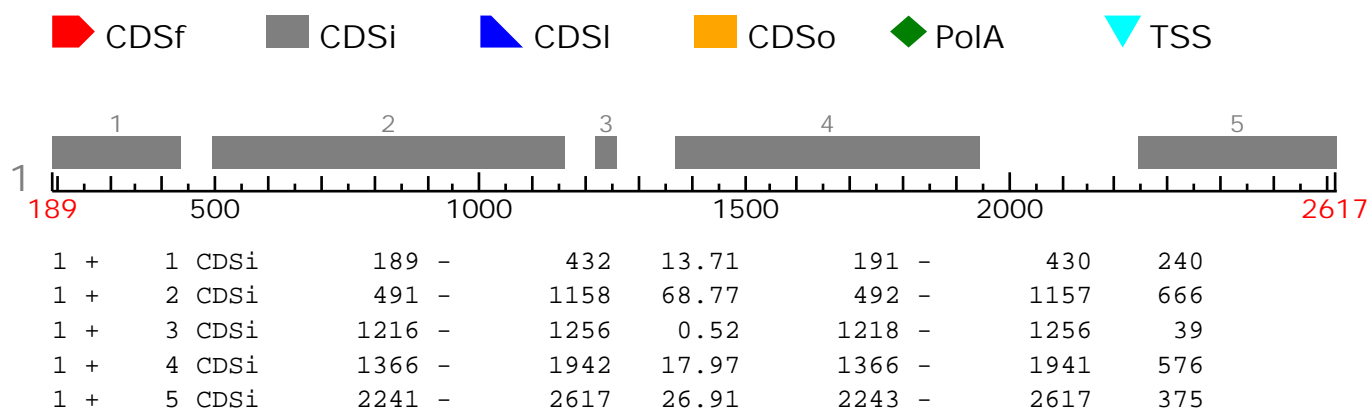


FGENESH 2.6 Prediction of potential genes in *Trichinella_spiralis* genomic DNA
 Seq name: gb|JAECWY010001388.1| Dprosaltans01
 Length of sequence: 2942
 Number of predicted genes 1: in +chain 1, in -chain 0.
 Number of predicted exons 5: in +chain 5, in -chain 0.
 Positions of predicted genes and exons: Variant 1 from 1, Score:107.760645



Predicted protein(s):

```
>FGENESH:[mRNA] 1 5 exon (s) 189 - 2617 1905 bp, chain +
GTGACTGGAGTGAAATTAATTCACGTGCCGAAGTGTGCTATTAAGAGAAAATTGTGGGAG
CAGAGCCTGGGGTGCAGCCTTGGTGAAGAACTCCCAAATTTGTGATACCCACTTTAATAAT
TCTGCAAAAGGTCAGACATTTAAAAGGAGGCGACTTAACGCAGATGCCATACCTAGTAAA
GTGGTGGAGCCTGAACCAGAAAAATAAAGAAGGCTATGCCAGTGGGAGTACACAAACA
GAGTCCTGTTTCATTGTTTAAATGAAAATAAGAGCTTGAGGGAAAAAATTCGTACCTTGGAG
TACGAAATGCGTCGCTTAGAGCAGCAGCTGAGGGAGTCTCAACAGTTGGAGGAGTCTCTA
CGCAAAATCTTCACGGAACGCAGATACGGATACTGAAGAATGGTGGACAAAGAGCGACG
TTCAATTCCGACGACATTTCTACGGCTATTTGTCTCCACACCGCAGGCTCTCGAGCGTAT
AACCATCTATACAAAAAAGGATTTCTTTTACCCAGTCGGAGCACCTTTGTACAGATGGTTA
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GCCTTCGAGTATGACAGCTCTGCTGATATTGTTTACGAGCCAAGCGACTATGTCCAACGTG
GCTATTGTTTCGTGGTCTAAAAAAATCGTGGAAGCAGCCAGTTTTTTTTTCGATTTTAATACC
CGAATGGACCCGGATACTCTTAACAACATATTAAGGAACTGCATAGGAAAGGATATTTA
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GCTAGCTCCATCAGACGCTGCTATTTCATTGGGGTATGACATTGAAAATGCCACCGAAACT
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AAAAAATTGAGACTCAGTGACAAAGTAAAAGAAAATTTGACATTTACATACGTCGACGAG
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GAAAAATTGATATTAGCAGCGCAAAACGTTGATGTCGATAAACTA

>FGENESH: 1 5 exon (s) 189 - 2617 635 aa, chain +

VTGVKLIHVPKCAIKRKLWEQSLGCSLGENSQICDTHFNNSAKGQTFKRRRLNADAIPSK
VVEPEPEKIKEGYASGSTQTESCSLFNENKSLREKIRTLEYEMRRLEQQLRRESQQLEESL
RKIFTETQIRILKNGGQRATFNSDDISTAICLHTAGSRAYNHLYKKGFPLPSRSTLYRWL
SDVDIKTGCLDVVIDLMDSDGVDDADKLCVLAFFDEMKVAAAFEYDSSADIVYEPSDYVQL
AIVRGLKKS WKQPVFDFNTRMDPDTLNNILRKLHRKGYLVVAIVSDLGTGNQKLWTELG
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FFGSMRSRGGQFDHPTPLQFKYRLRKYIIDYEPVDDEPVDDEPVADEPVADEL SLISNID
VTEMDELTEDAMEYIAGYVIKKLRLSDKVKENLTFTYVDEVSHGGLIKPSEKFQEKLKEL
ECIFSQYTNNNNFEITNNVKEKLILAAQNVDVDKL