

A

NCBI Resources How To truchanghabin My NCBI Sign Out

ORFinder

Open Reading Frame Viewer Help

Sequence

ORFs found: 9 Genetic code: 1 Start codons: 'ATG' only

Find:

Tools:  Tracks:

ORF Finder 11.2.201046137

ORF4 ORF5 ORF6 ORF7 ORF8 ORF9

1: 1..1.2K (1.2Kbp) Tracks shown: 2/18

ORF4 (50 aa)

```

>|1|ORF4
PTIQKGLRASSSPVPTAQHGDSGGAAHFRARISLTLEAREATVLYAGV
ALFACLTVYTRVGLSLELNSGQVYRKHFRHNRDQKCL
  
```

Mark subset... Marked 0  as

Label	Strand	Frame	Start	Stop	Length (nt   aa)
ORF4	+	3	180	452	273   90
ORF2	+	2	605	874	270   89
ORF9	-	2	565	317	249   82
ORF7	-	2	1117	920	198   66
ORF5	+	3	465	608	144   47
ORF1	+	1	1054	>1176	123   40
ORF6	-	1	263	147	117   38
ORF3	+	2	1082	>1177	96   31
ORF8	-	2	830	773	78   26

Six frame translation...

ORF4

BLAST Database:

B

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BLAST<sup>+</sup>

Job title:

Results for:

Hit:

Query ID:

Query length:

Database Name:

Description:

Program:

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Support center

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NIH