

BLAST Results

[Questions/comments](#)

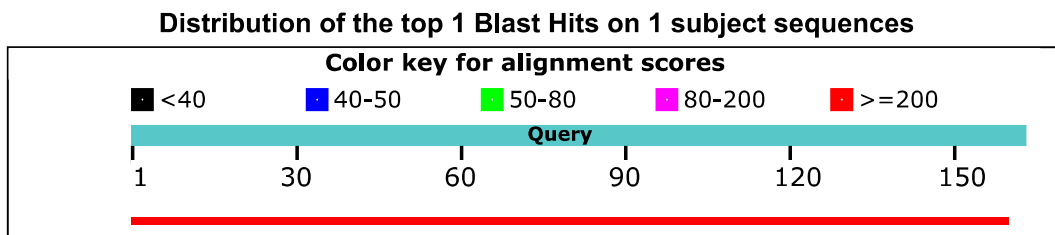
Job title: Y11470.1

RID [FD5HP52R01R](#) (Expires on 08-14 02:04 am)

Query ID Icl|Query_341209
Description None
Molecule type dna
Query Length 159

Database Name nt
Description Nucleotide collection (nt)
Program BLASTN 2.13.0+

Graphic Summary.



Descriptions

Sequences producing significant alignments:

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
E.arvense ribosomal DNA, internal transcribed spacer 1	294	294	100%	2e-75	100.00%	Y11470.1

Alignments

E.arvense ribosomal DNA, internal transcribed spacer 1

Sequence ID: **Y11470.1** Length: 159 Number of Matches: 1

Range 1: 1 to 159

Score	Expect	Identities	Gaps	Strand	Frame
294 bits(159)	2e-75()	159/159(100%)	0/159(0%)	Plus/Plus	

Features:

Query	1	CCACGCATCCGTGCAAACCCCTGCAAATTGTGGACATCCCCAAGGAGGCGACTTCAATAC	60
Sbjct	1	CCACGCATCCGTGCAAACCCCTGCAAATTGTGGACATCCCCAAGGAGGCGACTTCAATAC	60
Query	61	CCTCCTGCGCCTGTCCGTCCGACATCCAGGGGTCAACTTCTGCCCTCCGATGGAGCACTC	120
Sbjct	61	CCTCCTGCGCCTGTCCGTCCGACATCCAGGGGTCAACTTCTGCCCTCCGATGGAGCACTC	120
Query	121	TTCAATGCGCTGAGACTCCCACATAGGAATATGAAATCA	159
Sbjct	121	TTCAATGCGCTGAGACTCCCACATAGGAATATGAAATCA	159

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