

BLAST Results

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Job title: Y11471.1

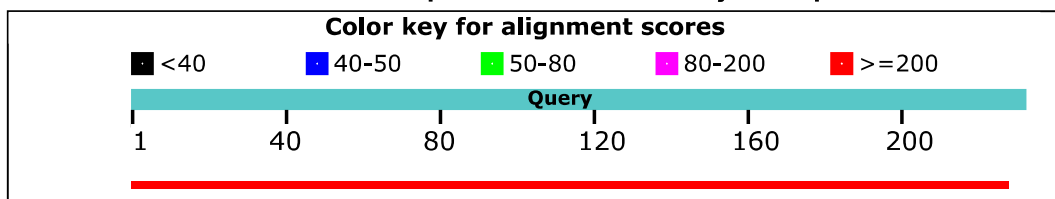
RID [FD5KCJW301R](#) (Expires on 08-14 02:05 am)

Query ID Icl|Query_169805
Description None
Molecule type dna
Query Length 227

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

Graphic Summary.

Distribution of the top 1 Blast Hits on 1 subject sequences



Descriptions

Sequences producing significant alignments:

Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
E.arvense ribosomal DNA, internal transcribed spacer 2	420	420	100%	5e-113	100.00%	Y11471.1

Alignments

E.arvense ribosomal DNA, internal transcribed spacer 2

Sequence ID: **Y11471.1** Length: 227 Number of Matches: 1

Range 1: 1 to 227

Score	Expect	Identities	Gaps	Strand	Frame
420 bits(227)	5e-113()	227/227(100%)	0/227(0%)	Plus/Plus	
Features:					
Query 1	TACAATCTATTGCGCCTCCCGTAGAGGGAGCGACAACGTGGTCGTCGCAACTCCTCCAGA	60			
Sbjct 1	TACAATCTATTGCGCCTCCCGTAGAGGGAGCGACAACGTGGTCGTCGCAACTCCTCCAGA	60			
Query 61	GGCACGGTTAGTTGAAATACATCGACGACACTGAGCGGCAAGACGAGTTGTAGGGGTGCT	120			
Sbjct 61	GGCACGGTTAGTTGAAATACATCGACGACACTGAGCGGCAAGACGAGTTGTAGGGGTGCT	120			
Query 121	CCTTCACCCGCTTGAAGTCTCGCGGGGGCTTACAGTGGTCACAGTCATCCGCCTCAGTGCG	180			
Sbjct 121	CCTTCACCCGCTTGAAGTCTCGCGGGGGCTTACAGTGGTCACAGTCATCCGCCTCAGTGCG	180			
Query 181	AAGCTAGAGAGGGCTTAGGGTCGACCTTGTGCGCTCCACCTACTC	227			
Sbjct 181	AAGCTAGAGAGGGCTTAGGGTCGACCTTGTGCGCTCCACCTACTC	227			

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