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BLAST Results

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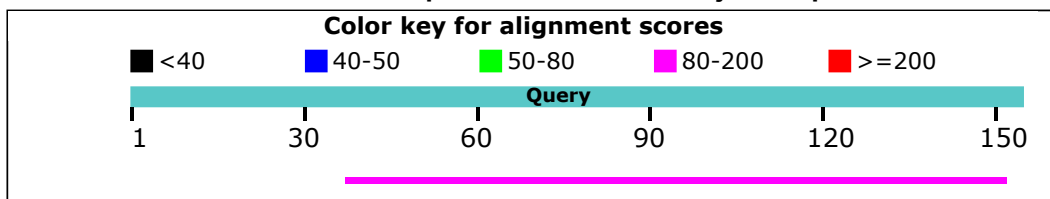
Blast 2 sequences

Job title: 64,976-1R

RID	7802F5US114 (Expires on 02-26 21:48 pm)	Subject ID	4799 subjects
Query ID	lcl Query_137753	Description	See details
Description	64,976-1R	Molecule type	nucleic acid
Molecule type	nucleic acid	Subject Length	4343931
Query Length	151	Program	BLASTN 2.8.1+

[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	Ident	Accession
MHIKPCM_02563 ATP-dependent 6-phosphofructokinase isozyme 1	147	147	75%	3e-36	89.74%	Query_140317

Alignments

MHIKPCM_02563 ATP-dependent 6-phosphofructokinase isozyme 1

Sequence ID: Query_140317 Length: 963 Number of Matches: 1

Range 1: 827 to 942

Score	Expect	Identities	Gaps	Strand	Frame
147 bits(79)	3e-36()	105/117(90%)	4/117(3%)	Plus/Minus	

Features:

```

Query 38  GTCCAGCCAGTCACCTTTGAACGGACGCTTTCA---TAAAGATAGCGTCMATGATGTCGT 94
Sbjct 942  GTCCAGCCAGTCACCTTTGAACGGACGCTT-CATGTTTTCGATAGCGTCGATGATGTCGT 884
Query 95  GWTGAACCCAGCTGCCCGTTTCRGGATACCTACACAACGACCGCCGTAACCTGCCAGCA 151
Sbjct 883  GGTGAACCCAGCTGTTCTGGATACCTACACAACGACCGCCGTAACCTGCCAGCA 827

```

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