

Query ID

lcl|57111

Description

Pros28.1ADsim (5'UTR and 5'region intron of GD19372 = 228 bp)

Molecule type

nucleic acid

Query Length

228

Subject ID

57113

Description

Pros28.1Amel (246 bp 5' region intron and part of exon of CG42322 +26bp 5'UTR = 272bp of the construct)

Molecule type

nucleic acid

Subject Length

272

Program

BLASTN 2.2.22


```
Query 49 AATAATTTATTT 60
      ||| |||
Sbjct 130 AATAATTTCTTT 119
```

Score = 15.7 bits (16), Expect = 1.1
Identities = 11/13 (84%), Gaps = 0/13 (0%)
Strand=Plus/Minus

```
Query 48 AAATAATTTATTT 60
      ||| ||| |||
Sbjct 201 AAATCCTTTATTT 189
```

Score = 15.7 bits (16), Expect = 1.1
Identities = 10/11 (90%), Gaps = 0/11 (0%)
Strand=Plus/Minus

```
Query 52 AATTTATTTAC 62
      || ||| |||
Sbjct 255 AAGTTATTTAC 245
```

Score = 13.9 bits (14), Expect = 3.7
Identities = 7/7 (100%), Gaps = 0/7 (0%)
Strand=Plus/Minus

```
Query 199 TATTTTG 205
      ||| |||
Sbjct 84 TATTTTG 78
```

Score = 13.9 bits (14), Expect = 3.7
Identities = 7/7 (100%), Gaps = 0/7 (0%)
Strand=Plus/Plus

```
Query 171 CAAAATA 177
        |||
Sbjct 78 CAAAATA 84
```

Score = 13.9 bits (14), Expect = 3.7
Identities = 9/10 (90%), Gaps = 0/10 (0%)
Strand=Plus/Minus

```
Query 22 ATAATATTTT 31
        |||
Sbjct 160 ATAATATGTT 151
```