

(1)srnaseq:atatacagggggagactctcat Length:22 Freq:21 chromosomal_pos:
 chr14.NT_026437.[82473123..82540116] (37169..37248)(+)
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 0.2555
 t - ga a a ttga
 ggtac taaa gagag t ccctttgtatgttc c t
 ||||| ||||| ||||| | ||||| ||||| ||||| |
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 c a ag - c gtaa
 MiPred Score: Real 85.3%

(2)srnaseq:atatacagggggagactcttat Length:22 Freq:181 chromosomal_pos:
 chr14.NT_026437.[82473123..82540116] (35947..36028)(+)
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 ca
 0.2533
 t g a ata a ttga
 tggtac tgaa ag gg ccctttgtatgttc c t
 ||||| ||||| || || || ||||| ||||| ||||| |
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 c a c aga c gtaa
 MiPred Score: Real 85.1%

(3)srnaseq:taaaaaccgtgactacttctg Length:21 Freq:2 chromosomal_pos:
 chr6.NT_025741.[12262885..12360950](4039..4122)(-)
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 agta
 0.3040
 g a a ag
 tattgggt ggtgca aagta ttgcggtttttgctatt t
 ||||| ||||| ||||| ||||| ||||| ||||| ||||| |
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 a c c ct
 MiPred Score: Real 83.7%

(4)srnaseq:cctcccactgcagagcctgggga Length:23 Freq:1 chromosomal_pos:
 chr10.NT_077569.[521740..607790] (35288..35409)(+)
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 caggctctgcagtgggaactgatgcctggaacagttcctgca
 0.2709
 gt acccactttt atgc cc cc c
 tg g gag ctgttccgggcatca tcccactgcagagcctggggag gga a
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 ac c ctt gacaaggctccgtagt agggtgacgtctcggacccttc cct g
 gt ----- ca -- c

(5)srnaseq:cggtttgaggctacagtgagat Length:22 Freq:7 chromosomal_pos:
chr11.NT_008984.[5728386..5729897] (75..191)(-)

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0.2815

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          |||      ||| ||||||||||||||| ||||||||||||||||||||| c
          cga      ctt gtgaactcgggtc ccaagctccgatgtcactctacac t
-----      ----- a              c              cg
```

MiPred Score: Real 76.4%