

Alignment to

```
1B_Full_26-5F_215.txt-- Matches:230; Mismatches:5; Gaps:211; Unattempted:0
1B_Full_26-5R_217.txt-- Matches:230; Mismatches:5; Gaps:210; Unattempted:0
```

* * * *
 400>ATCACAAC~~TTTGGAAATAAACTCGCTGTTTCTTGTA~~-TAAC~~TTACG~~->444
 191>ATCACAAC~~TTTGGAAATGA~~ACTCGCTG~~TTTCTTGTA~~AA~~TTACG~~TAACGA->237
 15<-----TGCA-TA-----GCAGAT-----G-----AA-----<1