

```

gi | 194664329 | ref | XP_001788158.1 | MGRAIVARLGLGLLLLALLLPQTYSNQTTVVTPSNSSQTTSP---APHPANATTKASDGTLQSTASLFVISVSLHLYC 77
gi | 194664954 | ref | XP_001787821.1 | MGRAIVARLGLGLLLLALLLPQTYSNQTTVVTPSNSSQTTSP---APHPANATTKASDGTLQSTASLFVISVSLHLYC 77
gi | 119901305 | ref | XP_001254295.1 | MGRAIVARLGLGLLLLALLLPQTYSNQTTVVTPSNSSQTTSP---APHPANATTKASDGTLQSTASLFVISVSLHLYC 77
gi | 7019343 | ref | NP_037362.1 | MGRAMVARLGLGLLLLALLLPQTQIYSSETTTGTSNSNSQSTSN SGLAPNPTNATTKAAGGALQSTASLFVVSLSLHLYS 80
1.....10.....20.....30.....40.....50.....60.....70.....80

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