

```

gi|6677833|ref|NP_033138.1|-----*:::* *::: .:::* :. *:::*: ** *:::*** **:****** *:*.
gi|13592079|ref|NP_112376.1|-----MPSQMEHAMEIMMLTFHRFAGDKD--HLTKEDLRVLMERFPGFLENQKDP LAVDKIMKDLDCRDGKVG 150
gi|73981599|ref|XP_533060.2|-----MPSQMEHAMEIMMLTFHRFAGEKN--VLTKEEDLRVLMERFPGFLENQKDP LAVDKIMKDLDCRDGKVG 150
gi|27807173|ref|NP_777075.1|-----MPPPP-----PSYRAAVSTDLTKMPSQMEHAMEIMMFTFHKFAGDKG--VLTKEEDLRVLMERFPGFLENQKDP LAVDKIMKDLDCRDGKVG 150
gi|4506761|ref|NP_002957.1|-----MPSQMEHAMEIMMFTFHKFAGDKG--VLTKEEDLRVLMERFPGFLENQKDP LAVDKIMKDLDCRDGKVG 150
gi|114559581|ref|XP_513807.2|-----MQEAHICPGIIEIAEWLREGGNARLKSLSKNFSFGSEORPFGHIGSOLAIFLIHPLNKTAGGGEGKTKWEKDASTDHTKMPSQMEHAMEIMMFTFHKFAGDKG--VLTKEEDLRVLMERFPGFLENQKDP LAVDKIMKDLDCRDGKVG 150
gi|47085671|ref|NP_998168.1|-----MPSDLERAMETLITVFHRYSGAEGNSSLSRRELKQLMEKELASFLKSKDPAAVDKIMKDL DANGDGEVN 150
gi|125844653|ref|XP_001341092.1|-----MPSLELAMESLIMVFRHYAGKEGRSGILTRRELRLIMENELSGFLKSKDPTFVDRIMKDL DANGDGEVD 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi|6677833|ref|NP_033138.1|FQSFLSLVAGLTIACNDYFVVNMKQKGGK-- 181
gi|13592079|ref|NP_112376.1|FQSFLSLVAGLTIACNDYFVVHMKQK--K-- 181
gi|73981599|ref|XP_533060.2|FQSFFSLIAGLTIACNDYFVIHMKQKGGK-- 181
gi|27807173|ref|NP_777075.1|FQSFFSLIAGLTIACNDYFVVHMKQKGGK-- 181
gi|4506761|ref|NP_002957.1|FQSFFSLIAGLTIACNDYFVVHMKQKGGK-- 181
gi|114559581|ref|XP_513807.2|FQSFFSLIAGLTIACNDYFVVHMKQKGGK-- 181
gi|47085671|ref|NP_998168.1|FEEFVSLVGLSIACEQLYQKQMKAAAKK-- 181
gi|125844653|ref|XP_001341092.1|FEEFVSLVGLSIACEACRYRVDLRKSQSWNL 181
.....160.....170.....180.

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