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**      *****.***** * * * :*:****      * * * . * . * * *****: * * . * * * *
gi |9506923|ref|NP_061933.1|-----MSIQOSAVSAKGFSSQGPAPCPAPATPAPASSSSSCCG-----GRGCCGDSGCCGSSSTSCCCFRRRRRQRSSGCCCGGG-SQR 84
gi |21311957|ref|NP_083074.1|-----MSQ-----GAKGFSSQSSAPCPAPTAPASSSSSCCGGCCRGCCGCCGDSGCCGSSSTGCCFRRRRRQRSSGCCCGGG-SQR 85
gi |73981608|ref|XP_864291.1|MAIFEGRVIMGNSKASGQYFIQLLLRCPANKASSNAEPLPSAPPSPAPRVHRELSSTMSFPOSAAAKGFSSQGPAPCPAPEPEPA--ASAACC-----CCGDSGCCGCC--GCCCFRRRRRQRSSGCCCGGG-SQR 137
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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* * * * :*.***. *
gi |9506923|ref|NP_061933.1|SQRS-NNRSSGCCSG-C 99
gi |21311957|ref|NP_083074.1|SQRS-CNNOSSGCCSGG-C 102
gi |73981608|ref|XP_864291.1|SQSSSNVDSQCCCGG-C 153
.....160.....

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