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gi|5454132|ref|NP_006281.1|  :: . . . : : * * : * : * . . : * * : * . * * * * . * : . * * * : . * * * : * : * * * * : * * * * *
-SSQRRDVPRTTQSTSRPKCARASCKNILACRSEELCMCOHPNORMGPGAHRGEPAPEDPPKQRCRAPACDHFGNAKNGYCNCECFQFKOMYG- 995
gi|114609563|ref|XP_527515.2| -SSQRRDVPRTTQSTSRPKCARASCKNILACRSEELCMCOHPNORMGPGAHRGEPAPEDPPKQRCRAPACDHFGNAKNGYCNCECFQFKOMYG- 995
gi|31543880|ref|NP_033423.2| -SSQHRDMPRTTQVSRPKCARASCKNILACRSEELCMCOHLSORVGSVAHRGEPPTPEEPPKQRCRAPACDHFGNAKNGYCNCEYQFKOMYG- 995
gi|57032092|ref|XP_541123.1| -SSQRRDAPRSTPSASRPKCARASCKNILACRSEELCMCOHLSORVGAGVHRGEPVPEEPPKQRCRAPACDHFGNAKCSGYCNCECFQFKOLYG- 995
gi|76626126|ref|XP_584832.2| -SSQRRDLPRASQASRSKCARATCRNVLACCSQELCMCOHLGQRAGISLRPEHTPEEPPQRCRAPACDHFGNAKNGYCNCECFQFKOMYG- 995
gi|189526589|ref|XP_692922.3| LVTIRASKARPPPPVLTQACRRSGCKNLSFG-CTDLCFDCLSRGREG---RRAQAPKPKSKQRCKIQGCDHYANQEKQGYCNCECDHFQIYRG 995
.....910.....920.....930.....940.....950.....960.....970.....980.....990.....

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