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*.: *** ** :. ***: * ** : * . *.: * ** :. ** *. * :*: .***.*** ** :*:*** *****.***.*** .***:***
gi|9966907|ref|NP_065160.1| MASR-WAVQLLVAAWSMGCCEALKCYTCKEPMISASCRTIIRCKPEDTACMTLLVTVAEYFFNQSPVVTSCSSSCVATDPSIGAAHLIFCCFRDLCNSE-L----- 150
gi|114622052|ref|XP_001149955.1| MASR-WAVQLLVAAWSMGCCEALKCYTCKEPMISASCRTIIRCKPEDTACMTLLVTVAEYFFNQSPVVTSCSSSCVATDPSIGAAHLIFCCFRDLCNSE-L----- 150
gi|10048458|ref|NP_065265.1| MTLR-WAMWLLLLAAWSMGYGEAFRCYTCEQPTAINCKNIQAQCKMEDTACKTIVLETVEAEFFNHSPMVTSCSSSCVATDPSIGAAHLIFCCFRDLCNSG-FPGFVAGL----- 150
gi|27662046|ref|XP_216966.1| MTLR-WAMWLLLLAAWSMGYGEAFRCYTCEQPTAINCKNIQAQCKMEDTACKTIVLETVEAEFFNHSPMVTSCSSSCVATDPSIGAAHLIFCCFRDLCNSG-FPGLVITGL----- 150
gi|73974816|ref|XP_539190.2| MARCGAALLLLLLAAASLRNGEAFRCFTCELPFVSLCKNIVVCKPEDTACKTIVLETVEAEFFNHSPMVTSCSSSCVATDPSIGAAHLIFCCFRDLCNSV-QASRLLGHLF----- 150
gi|119906252|ref|XP_876124.2| MAVPRALLALLLAASAVHGLGALSCTCEQPTALPLCKNITVCKPNEIACKTIVLETVEAEFFNHSPMVTSCSSSCVATDPSIGAAHLIFCCFRDLCNSVGLARLSAGALAPLGAVDGIVFMGGWVCGAQVCRGNKEGPGEEGLA----- 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi|9966907|ref|NP_065160.1| ----- 171
gi|114622052|ref|XP_001149955.1| ----- 171
gi|10048458|ref|NP_065265.1| ----- 171
gi|27662046|ref|XP_216966.1| ----- 171
gi|73974816|ref|XP_539190.2| ----- 171
gi|119906252|ref|XP_876124.2| GHFLRGLARLLKDGLOSVLGG 171
.....160.....170.

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