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*** :;*** *****:*****:;*: *.:***** * **.* ** *.:*****:*.:***** ***: : *; *.:*: *****:*.:*.:*** *.:**.*:***.***:***
gi |6754288|ref|NP_034631.1| MSEVNRSELEAILPOLKCHFTWNLFRREGSMSSHMEDRVCNQVEHLNSEEKATMYDLLAYIKHLDGESKAALECLGQAE DLRRSEHKDQAEIRRLVTWGN YAWIYYHMRGLSEAGAVDKVRVVCOKFANPYSMECPLECEEGWTRLKCG 150
gi |94405886|ref|XP_001000862.1| MSEVNRSELEAILPOLKCHFTWNLFRREGSMSSHMEDRVCNQVEHLNSEEKATMYDLLAYIKHLDGESKAALECLGQAE DLRRSEHKDQAEIRRLVTWGN YAWIYYHMRGLSEAGAVDKVRVVCOKFANPYSMECPLECEEGWTRLKCG 150
gi |56090427|ref|NP_001007695.1| MSEVNKESLEKILPOLKCHFTWNLFKEGSISSHMEDRVCNQIENLNSEKATMYDLLAYIKHLDGESKAALECLGQAE DLRRSEHKDQAEIRRLVTWGN YAWIYYRIGQLSEAGAVADKVRVVCOKFANPYSMECPLEDC EEGWTRLKCG 150
gi |72534658|ref|NP_001026853.1| MSEVTKNSEKILPOLKCHFTWNLFKEDSVSRDLEDRVCNQIEFLNTEFKATMYNLLAYIKHLDGESKAALECLGQAE DLRRSEHKDQAEIRRLVTWGN YAWIYYHMRGLSDACIYVDRVVKQCKKFSNPYSIEYSELDC EEGWTRLKCG 150
gi |115495365|ref|NP_001068882.1| MSE-DKNSLEQILPOLRCHFTWNLFKGGSVSRDLEDRVCNQIEFLNTEFKATMYNLLAYIKHLDGESKAALECLGQAE DLRRSEHKDQAEIRRLVTWGN YAWIYYHMRGLSDACIYVDRVVKQCKKFSNPYSIEYSELDC EEGWTRLVKCG 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

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gi |6754288|ref|NP_034631.1| --RNERAKMCFEKALEEKPDPCESSGMAIAMFRLEEKPEKQFSDALKQAMELNPNQYKLVLLALKLLRMGEEAEGERLTKDALGKAPNQTDLVQKAAQFYKKGKGNLDRATELLGKALRSTVNNSPLYSLVMCRYREITILEQLONK GDA 300
gi |94405886|ref|XP_001000862.1| --RNERAKMCFEKALEEKPDPCESSGMAIAMFRLEEKPEKQFSDALKQAMELNPNQYKLVLLALKLLRMGEEAEGERLTKDALGKAPNQTDLVQKAAQFYKKGKGNLDRATELLGKALRSTVNNSPLYSLVMCRYREITILEQLONK GDA 300
gi |56090427|ref|NP_001007695.1| --RNERAKMCFEKALEEKPDPCECRAGMAIAMYRLQEKPEKQFSDALKQAMELNSINQYKLVLLALKLQKMGEEAERLVEDALEKGFNQTDLVQKAAQFYKKGKGNLDRATEFLGKALRSTANNSPLYSLVMCRYREKLLQONTGEA 300
gi |72534658|ref|NP_001026853.1| --RNERAKVCFEKALEEKPNNPEFSSGLAIAMYYLDDRPQSSLEILKQAVELSPDNQYKLVLLALTLQKTNESQEGEQLVVEALEKAPCQTDVLLSAAKFYRKGKGLDKAIGLLRALEIPNNPILYHQIACCYRAKVKQIQHIGES 300
gi |115495365|ref|NP_001068882.1| GKKTIERAKVYFQKALEEKPNNPEFSSGLAIAMYYLDDRPQSSLEILKQAVELSPDNQYKLVLLALTLQKTNESQEGEQLVVEALEKAPCQTDVLLSAAKFYRKGKGLDKAIGLLRALEIPNNPILYHQIACCYRAKVKQIQHIGES 300
.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....260.....270.....280.....290.....300

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gi |6754288|ref|NP_034631.1| DSSERRORMAELRRLTMEFMQKTLQRRRSPPLNSYSDLLIDFPEVERCYQMVISKESPDVEEEDLYERYCNLQYHRKSEDLAAL ECLLOFPFRNERSIEKEEVKQOT----- 450
gi |94405886|ref|XP_001000862.1| DSSERRORMAELRRLTMEFMQKTLQRRRSPPLNSYSDLLIDFPEVERCYQMVISKESPDVEEEDLYERYCNLQYHRKSEDLAAL ECLLOFPFRNERSIEKEEVKQOT----- 450
gi |56090427|ref|NP_001007695.1| DKSERRQTMBELRRLTMEYVNNKALQRRPSPPLNSYSDLLIDFPEVEKCYQMVFSSESPRAEDQELFERYCNLQYHRKSEGLKALKDLMOFPFRPEKSTKEKEEVKQOTONGDLNLO----- 450
gi |72534658|ref|NP_001026853.1| EASGNKEMTEALKQYAMDYSNKALEKGLNPLNAYSDLAEFLEI-ECYQTPFNKEVPDAEKQOSHORYCNLQKVNKSEDTAVQHGLEGLSISKSTDKKEIKDQPNVSENLLPONAPNYWYLOGLIHKONGDLLQAAKCYEKELGRLLR 450
gi |115495365|ref|NP_001068882.1| EAIKQKEKIEELREHARNVYSKAIKGLNPLHACSD--EFLNEECYQTAFSKELPSTKQELHGDHCPQEDREISEHTAVQCSDLGSLISTESTEREKMKFQIQLNIANNQLPONVYVSWLQGLIHKMNGDLLQAECEYKALDHLK 450
.....310.....320.....330.....340.....350.....360.....370.....380.....390.....400.....410.....420.....430.....440.....450

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gi |6754288|ref|NP_034631.1| ----- 493
gi |94405886|ref|XP_001000862.1| ----- 493
gi |56090427|ref|NP_001007695.1| ----- 493
gi |72534658|ref|NP_001026853.1| DAPSGIGSIFLSAEELEDGSEEMCGAVSSPRELLSNSEQLN 493
gi |115495365|ref|NP_001068882.1| NSPSGIGSIFLPAIEREEGSEEMEDADSSILRELDPD----- 493
.....460.....470.....480.....490...

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