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** : ** : *.***** ** .:**** .* ** * * ** . * ** ** ** :****.*****.*****.*****.*****.*****.*****.*****
gi | 61175244 | ref | NP_001012996.1 | MENVPKENKVV EKAPVQN - EAPALGGGEYQEPGGNVKGVWAPPAPGFGEDVPNRLVDNIDMIDGDGDDMERFMEEMLRELRRKIRELQLRYSRLRILIGDPPHDDHDEFCLMP 111
gi | 55663803 | ref | XP_521186.1 | MENVPKENKVV EKAPVQN - EAPALGGGEYQEPGGNVKGVWAPPAPGFGEDVPNRLVDNIDMIDGDGDDMERFMEEMLRELRRKIRELQLRYSRLRILIGDPPHDDHDEFCLMP 111
gi | 115496308 | ref | NP_001069203.1 | MEKDPKERREEEQAPVQNEEACPMGGGEGPKPRENVRGDWPPAQDFREDMPNGLVNNIDIIDGDADDMERFMEEMLRELRRKIRELQLRYSRLRILIGDPPHDDHDEFCLMP 112
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110..

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