```
                                 ***:****.  .****:::*****.**** *  :.  * **.:***:* .        .*.***  .*.***  .*.***  .*.***  .*.***  .*.*** :.*. **   *.*        **                  :*
gi|6678117|ref|NP_033291.1|      MSSHQQKQPCTAPPQLHEQQVKQPCQPPPPEPCVSQVKTPCDTKVPEPCH--------PKAPEPCHPKAPEPCHPKAPEPCHPKAPEPCHPKAPEPCHPKAPEPCHPKAPEPCHPKAP--------EPCHPKVPEPCLPKAPEPCQPIVP   134
gi|109465129|ref|XP_001065728.1| MSSHQQKQPCTAPPQLHEQQVKQPCQPPPQEPCVSQIKSPCDTKVPEPCD--------TKTPEPCHPKVPEPCHPKAPEPCHPKAPEPCHPKAPEPCHPKAPEPCHSKVPEPCHPKAP--------EPCHP--------KAPEPCHPKVP   126
gi|77735555|ref|NP_001029473.1|  MSSHQQKQPCIPPPQLQQHQVKQPCQPPPQEPCIPQTKEPCHTKVPEPCH--------PKVPEPCHPKVPEPCHPKVPEPCHPKVPEPCHPKVPEPCHPKAPEPCHPKVPEPCHPKVP--------EPCHP----------------KVP   118
gi|4885607|ref|NP_005407.1|      MSSYQQKQTFTPPPQLQQQQVKQPSQPPPQEIFVPTTKEPCHSKVPQPGNTKIPEPGCTKVPEPGCTKVPEPGCTKVPEPGCTKVPEPGCTKVPEPGCTKVPEPGYTKVPEPGSIKVPDQGFIKFPEPGAIKVPEQGYTKVPVPGYTKLP   150
gi|114559645|ref|XP_001136610.1| MSSYQQKQTFTPPPQLQQQQVKQPSQPPPQEIFVPTTKEPCHSKVPQPGN--------TKVPEPGCTKVPEPGCTKVPEPGCTKVPEPDCTKVPEPGCTKVPEPGYTKVLEPGSIKVPDQGFIKFPEPGAIKVPEQGYTKVPVPGYTKVP   142
                                 1.......10........20........30........40........50........60........70........80........90.......100.......110.......120.......130.......140.......150
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                                 ** **.* *  ********
gi|6678117|ref|NP_033291.1|      EPCPSTVTPILAQQKTKQK   153
gi|109465129|ref|XP_001065728.1| EPCPSTVAPILAQQKTKQK   145
gi|77735555|ref|NP_001029473.1|  EPCPSPVIPGPAQQKTKQK   137
gi|4885607|ref|NP_005407.1|      EPCPSTVTPGPAQQKTKQK   169
gi|114559645|ref|XP_001136610.1| EPRPSTVTPGPAQQKTKQK   161
                                 .......160.........
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