```
                                 ***:****:**: *****:****::: * : **::*:*::***.***:** *:*:**. ** **.***:  ** .:* **  **:***:**:    ****.**::* .*::**: *      :: :*:  :  *   * **:** :**
gi|7305267|ref|NP_038626.1|      MKKTQTWIITCIYLQLLLFNPLVKTKEICGNPVTDNVKDITKLVANLPNDYMITLNYVAGMDVLPSHCWLRDMVIQLSLSLTTLLDKFSNISEG---LSNYSIIDKLGKIVDDLVLCMEENAPKN-IKESPKRPETRSFTPEEFFSIFNR   146
gi|33667068|ref|NP_068615.1|      MKKTQTWIITCIYLQLLLFNPLVKTQEICRNPVTDNVKDITKLVANLPNDYMITLNYVAGMDVLPSHCWLRDMVTHLSVSLTTLLDKFSNISEG---LSNYSIIDKLGKIVDDLVACMEENAPKN-VKESLKKPETRNFTPEEFFSIFNR   146
gi|4505175|ref|NP_000890.1|      MKKTQTWILTCIYLQLLLFNPLVKTEGICRNRVTNNVKDVTKLVANLPKDYMITLKYVPGMDVLPSHCWISEMVVQLSDSLTDLLDKFSNISEG---LSNYSIIDKLVNIVDDLVECVKENSSKD-LKKSFKSPEPRLFTPEEFFRIFNR   146
gi|114646140|ref|XP_509255.2|    MKKTQTWILTCIYLQLLLFNPLVKTEGICRNRVTNNVKDVTKLVANLPKDYMITLKYVPGMDVLPSHCWISEMVVQLSDSLTDLLDKFSNISEG---LSNYSIIDKLVNIVDDLVECVKENSSKD-LKKSFKSPEPRLFTPEEFFRIFNR   146
gi|60678250|ref|NP_001012753.1|  MKKTQTWIITCIYLQLLLFNPLVKTKGICGKRVTDDVKDVTKLVANLPKDYKIALKYVPGMDVLPSHCWISVMVEQLSVSLTDLLDKFSNISEG---LSNYSIIDKLVKIVDDLVECTEGYSFEN-VKKAPKSPELRLFTPEEFFRIFNR   146
gi|27805973|ref|NP_776800.1|     MKKTQTWIITCIYLQLLLFNPLVHTQGICSNRVTDDVKDVTKLVANLPKDYMITLKYVPGMDVLPSHCWISEMVEQLSVSLTDLLDKFSNISEG---LSNYCIIDKLVKIVDDLVECMEEHSSEN-VKKSSKSPEPRQFTPEKFFGIFNK   146
gi|45384094|ref|NP_990461.1|     MKKAQTWIITCFCLQLLLLNPLVKAQSSCGNPVTDDVNDIAKLVGNLPNDYLITLKYVPKMDSLPNHCWLHLMVPEFSRSLHNLLQKFSDISDMSDVLSNYSIINNLTRIINDLMACLAFDKNKDFIKENGHLYEEDRFIPENFFRLFNS   150
                                 1.......10........20........30........40........50........60........70........80........90.......100.......110.......120.......130.......140.......150

                                 :*:.:*::   : . .:*::.**   *::****:***.: :***********      **.**::*     ...*:  ::**.:::**:*** :**:**** :. . ..: *. * :  :*:********:**:*. :*
gi|7305267|ref|NP_038626.1|      SIDAFKDF-MVASDTSDCVLSSTLG-PEKDSRVSVTKPFMLPPVAASSLRND------SSSSNRKAAKAPEDSGLQWTAMALPALISLVIGFAFGALYWKKKQS-SLTRAVENIQIN--EEDNEISMLQQKEREFQEV   273
gi|33667068|ref|NP_068615.1|      SIDAFKDF-MVASDTSDCVLSSTLG-PEKDSRVSVTKPFMLPPVAASSLRND------SSSSNRKAAKSPEDPGLQWTAMALPALISLVIGFAFGALYWKKKQS-SLTRAVENIQIN--EEDNEISMLQQKEREFQEV   273
gi|4505175|ref|NP_000890.1|      SIDAFKDF-VVASETSDCVVSSTLS-PEKDSRVSVTKPFMLPPVAASSLRND------SSSSNRKAKNPPGDSSLHWAAMALPALFSLIIGFAFGALYWKKRQP-SLTRAVENIQIN--EEDNEISMLQEKEREFQEV   273
gi|114646140|ref|XP_509255.2|    SIDAFKDF-VVASETSDCVVSSTLS-PEKDSRVSVTKPFMLPPVAASSLRND------SSSSNRKAKNPPGDSSLHWAAMALPALFSLIIGFAFGALYWKKRQP-SLTRAVENIQIN--EEDNEISMLQEKEREFQEV   273
gi|60678250|ref|NP_001012753.1|  SIDAFKDLETVASKSSECVVSSTLS-PDKDSRVSVTKPFMLPPVAASSLRND------SSSSNRKASNSIGDSNLQWAAMALPAFFSLVIGFAFGALYWKKKQP-NLTRTVENIQIN--EEDNEISMLQEKEREFQEV   274
gi|27805973|ref|NP_776800.1|     SIDAFKDLEIVASKMSECVISSTSS-PEKDSRVSVTKPFMLPPVAASSLRND------SSSSNRKASNSIEDSSLQWAAVALPAFFSLVIGFAFGAFYWKKKQP-NLTRTVENRQIN--EEDNEISMLQEKEREFQEV   274
gi|45384094|ref|NP_990461.1|     TIEVYKEF-ADSLDKNDCIMPSTVETPENDSRVAVTKTISFPPVAASSLRNDSIGSNTSSNSNKEALGFISSSSLQGISIALTSLLSLLIGFILGAIYWKKTHPKSRPESNETIQCHGCQEENEISMLQQKEKEHLQV   287
                                 .......160.......170.......180.......190.......200.......210.......220.......230.......240.......250.......260.......270.......280.......
```