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gi | 21614544 | ref | NP_002955.2 | MLTELEKALNSIIDVYHKYSLIKGNFHAVYRDDLKKLLETECPQYIRKKGADVWFKELDINTDGAVNFQEFLLILVIKMGVAAHKKSHESHKE 93
gi | 114559700 | ref | XP_513814.2 | MLTELEKALNSIIDVYHKYSLIKGNFHAVYRDDLKKLLETECPQYIRKKGADVWFKELDINTDGAVNFQEFLLILVIKMGVAAHKKSHESHKE 93
gi | 73961680 | ref | XP_547583.2 | MLTELESAINSLIEVYHKYSLVKGNYHALYRDDLKKLLETECPQYMKKKDADTWQELDVNSDGAINFEEFLILVIKVGVAASHKDIHKE---- 89
gi | 194665098 | ref | XP_001790368.1 | MLTDLECAINSLIDVYHKYSLKKGNYHAVYRDDLKQLLETECPKFMKKKDADTWFKELDINQDGGINFEEFLVLVIKVGLEAHEEIHKE---- 89
gi | 7305453 | ref | NP_038678.1 | MPSELEKALSNLIDVYHNYSGIKGNHHALYRDDFRKMVTTECPQFVQNKNTESLFKELDVNSDNAINFEEFLALVIRVGVAAHKDSHKE---- 89
gi | 16758672 | ref | NP_446274.1 | MATELEKALSNVIEVYHNYSGIKGNHHALYRDDFRKMVTTECPQFVQNKNTESLFKELDVNSDNAINFEEFLALVIRVGVAAHKDSHKE---- 89
1.....10.....20.....30.....40.....50.....60.....70.....80.....90...

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