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                                 *:* : :. :   .        : .   *:* ***..*:**..*.* .: : .     *:: * . * * *..  :* :*:*:  * **.:    :*. *:*::******** *::: ::  *:**::***  * *.*** **.*: ***
gi|4503697|ref|NP_000791.1|      MAEGEITTFTALTEKFNLPPGNYKKPKLLYCSNGGHFLRILPDGTVDGTRDRSDQHIQLQLSAESVGEVYIKSTETGQYLAMDTDGLLYGSQTPNEECLFLERLEENHYNTYISKKHAEKN-WFVGLKKNGSCKRGPRTHYGQKAILFLP   149
gi|114602469|ref|XP_001153487.1| MAEGEITTFTALTEKFNLPPGNYKKPKLLYCSNGGHFLRILPDGTVDGTRDRSDQHIQLQLSAESVGEVYIKSTETGQYLAMDTDGLLYGSQTPNEECLFLERLEENHYNTYISKKHAEKN-WFVGLKKNGSCKRGPRTHYGQKAILFLP   149
gi|73949448|ref|XP_849274.1|     MAEGEITTFTALTEKFNLPPGNYMKPKLLYCSNGGHFLRILPDGTVDGTRDRSDQHIQLQLSAESVGEVYIKSTETGQYLAMDTDGLLYGSQTPNEECLFLERLEENHYNTYTSKKHAEKN-WFVGLKKNGSCKRGPRTHYGQKAILFLP   149
gi|6753850|ref|NP_034327.1|      MAEGEITTFAALTERFNLPLGNYKKPKLLYCSNGGHFLRILPDGTVDGTRDRSDQHIQLQLSAESAGEVYIKGTETGQYLAMDTEGLLYGSQTPNEECLFLERLEENHYNTYTSKKHAEKN-WFVGLKKNGSCKRGPRTHYGQKAILFLP   149
gi|6978839|ref|NP_036978.1|      MAEGEITTFAALTERFNLPLGNYKKPKLLYCSNGGHFLRILPDGTVDGTRDRSDQHIQLQLSAESAGEVYIKGTETGQYLAMDTEGLLYGSQTPNEECLFLERLEENHYNTYTSKKHAEKN-WFVGLKKNGSCKRGPRTHYGQKAILFLP   149
gi|45383992|ref|NP_990511.1|     MAEGEITTFTALTERFGLPLGNYKKPKLLYCSNGGHFLRILPDGKVDGTRDRSDQHIQLQLSAEDVGEVYIKSTASGQYLAMDTNGLLYGSQLPGEECLFLERLEENHYNTYISKKHADKN-WFVGLKKNGNSKLGPRTHYGQKAILFLP   149
gi|27806633|ref|NP_776480.1|     MAEGETTTFTALTEKFNLPLGNYKKPKLLYCSNGGYFLRILPDGTVDGTKDRSDQHIQLQLCAESIGEVYIKSTETGQFLAMDTDGLLYGSQTPNEECLFLERLEENHYNTYISKKHAEKH-WFVGLKKNGRSKLGPRTHFGQKAILFLP   149
gi|41152157|ref|NP_957054.1|     MTEADIAVKSSPRD--------YKKLTRLYCMNGGFHLQILADGTVAGAADEN-TYSILRIKATSPGVVVIEGSETGLYLSMNEHGKLYASSLVTDESYFLEKMEENHYNTYQSQKYGEN--WYVGIKKNGKMKRGPRTHIGQKAIFFLP   139
gi|157426953|ref|NP_001098748.1| MTERDFTAFALLSSTTDNRDTKHNSLIKLHCRNGGFHLRILPSGTVDASRQDNDINTLLKVKAVKAGVVAIRGHETGLYLAMDKCGRLYGTAVLNEECFFIEKMEENHYNTYRSQRYQDNGDWFVGIRKNGRTKDGSRTHKGQNAVYFLP   150
                                 1.......10........20........30........40........50........60........70........80........90.......100.......110.......120.......130.......140.......150
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                                   *. .
gi|4503697|ref|NP_000791.1|      LPVSSD--   155
gi|114602469|ref|XP_001153487.1| LPVSSD--   155
gi|73949448|ref|XP_849274.1|     LPVSSD--   155
gi|6753850|ref|NP_034327.1|      LPVSSD--   155
gi|6978839|ref|NP_036978.1|      LPVSSD--   155
gi|45383992|ref|NP_990511.1|     LPVSAD--   155
gi|27806633|ref|NP_776480.1|     LPVSSD--   155
gi|41152157|ref|NP_957054.1|     RQVEQEED   147
gi|157426953|ref|NP_001098748.1| IPVDGSVQ   158
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