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                                   .  .*:** ::**::*** ***:**: **   *  **:. * :****:.:*:. * ***::*: ** . :  *  ::::.:::** ::*.***:** *:**::*::::*.**:*                .:  :
gi|23943892|ref|NP_703152.1|      --------------MAGSQDIFDAIVMADERFHGEGYREGYEEGSSLGVMEGRQHGTLHGAKIGSEIGCYQGFAFAWKCLLHSCTTEKDSRKMKVLESLIGMIQKFPYDDPTYDKLHEDLDKIRGKFKQ--------------FCSLLNV   122
gi|114639006|ref|XP_508610.2|     --------------MAGSQDIFDAIVMADERFHGEGYREGYEEGSSLGVMEGRQHGTLHGAKIGSEIGCYQGFAFAWKCLLHSCTTEKDSRKMKVLESLIGMIQKFPYDDPTYDKLHEDLDKIRGKFKQTGLSPSAVIVRPPSYVELAVV   136
gi|31980738|ref|NP_082460.2|      MKGMRGPGESRDTAMAVEQDIFDAVVMADERFHGEGYQEGYEEGSSLGIVEGKRYGMVHGAKIGSEIGCYRGFALAWKCLLHSGAGEKDSRKMKVVEALIALLQDFPYDDPTYEKLHEDLDRIRGKFRQ--------------LCSLLNV   136
gi|109459696|ref|XP_219536.3|     MTGVRGPGESRDTAMAMEQDIFDAVVMADERFHGEGYQEGYEEGSSLGIVEGKQYGVLHGAKIGSEIGCYRGFALAWKCLLHSGAGEKE-RKMKVVEALITLLQDFPYDDPTYEKLHEDLERIRGKFRQ--------------LCSLLNV   135
gi|71894711|ref|NP_001026357.1|   -----------MAAPSAGSDMFDEIVMAEDRFHGEGYQEGYAEGSHAGAAEGRRCGALHGAKIGSEIGSYLGFALTWQHLLPKCTDEKDSKKMRALDSLIGMIQKFPYEDPTYEKLQDDLEKIRGKFKQ--------------VCSMLNI   125
gi|134133281|ref|NP_001077021.1|  -----------MAFEPNNDDLFDSIIMADDRFHVEGYQEGFDEGVRQGTIEGRNHGRLHGAKLSAEVSFYYGFALAWKCLLQSSTDVKARKRLKAMESLIGVIQNFPYEDPQYEKLQEDMERVRAKFRQ--------------VCSLLNV   125
                                  1.......10........20........30........40........50........60........70........80........90.......100.......110.......120.......130.......140.......150
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                                   .           *::                .:::
gi|23943892|ref|NP_703152.1|      QP-------------DFKISAE----------GSGLSF--------------------------------   137
gi|114639006|ref|XP_508610.2|     QPETHRPSRPPPRPSDLRVTGQTLLCPVHLPPAFGLSIHTTSRTCALAPSRQTKRPSRPPRGSDRRRSQDG   207
gi|31980738|ref|NP_082460.2|      QP-------------DFKVTPG----------GSGLAF--------------------------------   151
gi|109459696|ref|XP_219536.3|     QP-------------DFKVTAG----------GSGLAF--------------------------------   150
gi|71894711|ref|NP_001026357.1|   RS-------------DFRISNE----------RSSLTF--------------------------------   140
gi|134133281|ref|NP_001077021.1|  ST-------------DFREYLSG---------STSMSF--------------------------------   141
                                  .......160.......170.......180.......190.......200.......210.......220.
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