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gi 6981010|ref|NP_037228.1| MVL$ADKTNIKNCWGI$G$G$EY$G$EAL$QRMFAAF$TKTYF$SHIDV$SPGSAQVKAHGKKVADALAKAADHVEDLPGALS$TLDLHAHKL$RVD$PVN$FKFL$SHCLLVTLACHHPGDFTPAMHASLDRFLASVSTVLT$SKYR 142
gi 60678292|ref|NP_001007723.1| MVL$AADKTNIKNCWGI$G$G$EY$G$EAL$QRMFAAF$TKTYF$SHIDV$SPGSAQVKAHGKKVADALAKAADHVEDLPGALS$TLDLHAHKL$RVD$PVN$FKFL$SHCLLVTLACHHPGDFTPAMHASLDRFLASVSTVLT$SKYR 142
gi 6680175|ref|NP_032244.1| MVL$G$EDK$SNIKAAWGI$G$G$AEY$G$AEALERMFL$F$P$TKTYF$PHFDV$SHGSAQVKGHGKKVADALASAAGHLDDLPGALS$ALSDLHAHKL$RVD$PVN$FKLL$SHCLLVTLASHHPADFTPAVHASLDRFLASVSTVLT$SKYR 142
gi 4504345|ref|NP_000508.1| MVL$PADKTNVKAW$G$V$GAHAGEY$G$AEALERMFL$F$P$TKTYF$PHFDL$SHGSAQVKGHGKKVADALANAVAHVDDMPNALS$ALSDLHAHKL$RVD$PVN$FKLL$SHCLLVTLAAHLPAEFTPAVHASLDRFLASVSTVLT$SKYR 142
gi 4504347|ref|NP_000549.1| MVL$PADKTNVKAW$G$V$GAHAGEY$G$AEALERMFL$F$P$TKTYF$PHFDL$SHGSAQVKGHGKKVADALANAVAHVDDMPNALS$ALSDLHAHKL$RVD$PVN$FKLL$SHCLLVTLAAHLPAEFTPAVHASLDRFLASVSTVLT$SKYR 142
gi 116812902|ref|NP_001070890.2| MVL$AADKGNVKAW$G$V$G$HAAEY$G$AEALERMFL$F$P$TKTYF$PHFDL$SHGSAQVKGHGAKVAAAL$KAVEHLLDDLPGALS$ELSDLHAHKL$RVD$PVN$FKLL$SHLLVTLASHLPSDFTPAVHASLDRFLANVSTVLT$SKYR 142
gi 194678485|ref|XP_001788728.1| MVL$AADKGNVKAW$G$V$G$HAAEY$G$AEALERMFL$F$P$TKTYF$PHFDL$SHGSAQVKGHGAKVAAAL$KAVEHLLDDLPGALS$ELSDLHAHKL$RVD$PVN$FKLL$SHLLVTLASHLPSDFTPAVHASLDRFLANVSTVLT$SKYR 142
gi 52138655|ref|NP_001004376.1| MVL$AADKNNVKGIFTKIAGHAAEY$G$AEALERMFL$F$P$TKTYF$PHFDL$SHGSAQIKGHGKKVVAALIEAANHIDDIAAGTSLKSLSDLHAHKL$RVD$PVN$FKLL$GQCFLVVAIHH$PAAL$PEVHASLDRFLCAVGTVLT$AKYR 142
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140..

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