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gi 4758258|ref|NP_004086.1| MSGGSSCSQTPSRAIPA-TRRVVLGDGVQLPPGDYSTTPGGTLFSTTPGGTRIIYDRKFLMECRNSPVTKTPPRDLPIPGVTSP-----SSDEPPMEASQSHLRNSPEDKRAGGEESQFEMDI 124
gi 114619677|ref|XP_531121.2| MSGGSSCSQTPSRAIPA-TRRVVLGDGVQLPPGDYSTTPGGTLFSTTPGGTRIIYDRKFLMECRNSPVTKTPPRDLPIPGVTSP-----SSDEPPMEASQSHLRNSPEDKRAGGEESQFEMDI 124
gi 122937368|ref|NP_031944.3| MSAGSSCSQTPSRAIP--TRRVVLDGVDVQLPPGDYSTTPGGTLFSTTPGGTRIIYDRKFLMECRNSPVAKTPPKDLPAIPGVTSP-----TSDEPPMQASQSQLESSEPEDKRAGGEESQFEMDI 124
gi 16758724|ref|NP_446309.1| MSAGSSCSQTPSRAIP--TRRVVLDGVDVQLPPGDYSTTPGGTLFSTTPGGTRIIYDRKFLMECRNSPVAKTPPKDLPIPGVTSP-----TSDEPPMQASQSHLRNSPEDKRAGGEESQFEMDI 124
gi 118150896|ref|NP_001071361.1| MSGGSSCSQTPSRAIPT-TRRVVLADGVQLPPGDYSTTPGGTLFSTTPGGTRIIYDRKFLMECRNSPVTKTPPRDLPIPGVTSP-----TGDEPPTEARONHLRNSPEDKPAAGGEESQFEMDI 124
gi 73979318|ref|XP_549448.2| MSGGSSCSQTPSRAIPA-TRRVVLGDGVQLPPGDYSTTPGGTLFSTTPGGTRIIYDRKFLMECRNSPVTKTPPRDLPIPGVTSP-----ASEEPTTEPSONHLRNSPEDKPAAGGEESQFEMDI 124
gi 50806236|ref|XP_424384.1| MSGRCCQGQTPSRDIPGPGKRLALPDGAPLPPGDYSTTPGGTVFGTTPGGTRIIYDRKFLMECRNSPVAKTPPSDLPIPGVTSP-----IVEELKIEN--NHVQN-VDEKANVGEEEFDMDI 124
gi 41054483|ref|NP_955939.1| MSGM--QKITTSQAIPT-TRRVILNDAHLR-HDYSTTPGGTLFSTTPGGTRIIYDRKFLDLCRSPPLARIPPCCLPDIPGVTSPPSVTVNNEKAYPKPQVNNNSISPPVDKSTGEDAQFEMDI 124
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120....

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