

gi	57012420	ref	NP_001008833.1	SPSTINMGIDVTSVFLGPLAVFVVLAIIGAA--VAVVRRRRRNTGGEGG---DYTPAPGRDSS-----	360
gi	57012348	ref	NP_001008830.1	SPSTINMGIDVTSVVLGAVAVFVVLAIIGAV--VCVVRRRRRRNTGGKG---DYTPAPGRDSSQSSDVSLPDCKA	369
gi	57012402	ref	NP_001008840.1	SPSTDSNMGIDVTSVLPAGAVAVFVVLAIIGAV--VAIVRRR---NTGGKG---DYAPAPGRDSSQSSDVSLPDCKA	365
gi	57012396	ref	NP_001008835.1	PSSINSNMEFNVIYVVLGALS---AVAISGAV-IIGAVLAVVRKRNTGGEG---GDYTPAPDRDSCQSSNVSLPDCKA	371
gi	57012398	ref	NP_001008832.1	PPSTDSNMEFNVIYVVLGALS---AVAIIGAV-IIGAVLAVVRKRNTGRRKQGGKGDYAPAPDRDSCQSSNVSLPDCKA	375
gi	77404172	ref	NP_001029157.1	PPSTDSNMEFNVIYVVFAGVITIIIGAVAIIGAVAIIGVCVRRRRRRNTGGI-----	347
gi	57012400	ref	NP_001008845.1	PPSTDSNMEFNVIYVVLG--AISAVAIIG-AVLAVVRKK---NTGGTGG---DYTPAPGQ-POQL	355
gi	57012418	ref	NP_001008834.1	PPSTDSNMEFNVIYVVGALVITIGAVIIG-AVVAVVRKR---NTGGTGG---DYAPAPGK-ANPNSVSLPDCKA	367
gi	57012424	ref	NP_001008827.1	PPSTDSNMEFNVIYVVLGAVAIIAAVIIG-AVVTIVVRKRRR---NTGGKG---VYTPAPGTDSSQSSDVSLPDCKA	369
gi	57012422	ref	NP_001008841.1	SLSTDSNMEFNVIYVVLG-AIS-ALAIIG-ALVPVVRKRR---NTGGKGG---VYAPAPG-----	352
gi	57012342	ref	NP_001008829.1	SPSTINSNMEFNVIYVVLG-AITGLAIIG-IVVAVVRKRRR---NTGGKG---DYTPAPGRDSSQSSDVSLPDCKA	366
gi	57012426	ref	NP_001008839.1	SPSTDSNMEFNVIYVVLGAVAIIAAVIIV-AVVAVVMKRRR---NTGGKVG---VYAPAPSRDSSQSSDVSLPDCKA	370
gi	57012416	ref	NP_001008836.1	PSSDFNMRII AVLVLGAVAIIG---AVVAVMMKRRR---NTGGTGG---DYTPAPGKASPNKCDLSLPLCKA	365
gi	47777291	ref	NP_996730.1	PPSTDPNMEFNVIYVVGAVVITGAVLMIIGALLIVVRGRKR---NTGGKVG---VYTPAPGKDSQSSDVSLPDCKA	368
gi	57012404	ref	NP_001008842.1	PPGTDNMEFNVIYVVFAGAVAMIGAVAVIGAVVAVRRRRRR-KRNTGGKG---DYAPAPGSDSCQSSDVSLADCEA	373
gi	57012412	ref	NP_001008838.1	PPGTDNMEFNVIYVVLGAVAMIGAVAVIGAVVAVRRRRR-KRNTGGTGG---DYAPAPGSDSCQSSDVSLADCEA	373
gi	57114336	ref	NP_001008843.1	PPSTNSNTGMSVILGVTIIIAVMAIIVVAVIRIVVRKRWI---KGGKGV---YTPAPGRDSSQSSDVSLPDCKA	370
gi	6981494	ref	NP_036777.1	SPSTDSN-----LLLLFLELWQFL-----	319
gi	133922588	ref	NP_001001892.2	PPSTVSNMAIVAVLVLGAAIVTGAVVAF---VMKMRRR---NTGGKGG---DYALAPGS---DTSDLSLPDCKVMVHDPHSLA	369
gi	133778955	ref	NP_034510.3	PPSTDSYMVIVAVLGVLGAMAIIGAVVAF---VMK-RRR---NTGGKGG---DYALAPGS---QSSMSLRDCKA	362
gi	52630342	ref	NP_002108.4	-----SSQPTIPIMGIVAGLAVLVLAVLAVLAVVITAMCRRK---SSGGKGG---SCSQAAACSNAAQSSDESLITCKA	366
gi	60115686	ref	NP_001012441.1	-----SSQPTIPIVGIAAGLAVLAVLAVLAVLAVVAVVMCRRK---SSGGKGG---SCSQAAASSNAAQSSDESLIACKA	366

