

NTDB id 602 KW2 RS03060 WP 011834647.1	MGYLVPTVIEQSSRGERAYDIYSRLKDR IIMLTGTPVEDGMANS	IAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	75
NTDB id 611 V4T04 RS07970 WP 003129593.1	MGYLVPTVIEQSSRGERAYDIYSRLKDR IIMLTGTPVEDGMANS	IAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	75
NTDB id 330 STU RS11365 WP 002949732.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPIEDNMANS	IAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	72
NTDB id 298 STER RS01925 WP 002949732.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPIEDNMANS	IAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	72
NTDB id 282 KZH43 RS03215 WP 000613477.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDSTKDIYLYVNTPGGSVSAG	72
NTDB id 246 SPD RS03495 WP 000613477.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDSTKDIYLYVNTPGGSVSAG	72
NTDB id 244 SPR RS03300 WP 000613477.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDSTKDIYLYVNTPGGSVSAG	72
NTDB id 242 SP RS03655 WP 000613477.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDSTKDIYLYVNTPGGSVSAG	72
NTDB id 342534 ETT44 RS07705 WP 002985850.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	72
NTDB id 446 SpyM6JRS4 RS01695 WP 002985850.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	72
NTDB id 445 SPYM3 RS01720 WP 002985850.1	MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	72
NTDB id 376 SMU RS07595 WP 255262720.1	MELKEKYS MIPVVIEQTSRGRERSYDIYSRLKDR IIMLTGTPVEDNMANS	IAQLFLDQAQDNTKDIYLYINSPGGSVSAG	80
NTDB id 1223 Cj0192c YP 002343650.1	MFIPYVIEKSSRGRERSYDIYSRLKDR IIMLTSGEIHDELAAS	IVAQLFLFEAEDPTKDIYLYINSPGGVITSG	73
NTDB id 86 BSU 34540 NP 391334.1	MNLIPTVIEQTNRGERAYDIYSRLKDR IIMLTGSAIDNNVANS	IVSQLLFLAAEDPEKEISLYINSPGGISITAG	74
consensus	*****	*****	

[illegible]

NTDB id 602 KW2 RS03060 WP 011834647.1	ILADNSNRSLQIHKDAERDHWMDAKETLEYGFIDEIMENNSLK.	199
NTDB id 611 V4T04 RS07970 WP 003129593.1	ILADNSNRSLQIHKDAERDHWMDAKETLEYGFIDEIMENNSLK.	199
NTDB id 330 STU RS11365 WP 002949732.1	ILADNSGQPIEKVHVDAERDNWMSAQETLEYGFIDEIMTNNQLK.	196
NTDB id 298 STER RS01925 WP 002949732.1	ILADNSGQPIEKVHVDAERDNWMSAQETLEYGFIDEIMTNNQLK.	196
NTDB id 282 KZH43 RS03215 WP 000613477.1	ILAE NSGQSM EK V HAD AERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 246 SPD RS03495 WP 000613477.1	ILAE NSGQSM EK V HAD AERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 244 SPR RS03300 WP 000613477.1	ILAE NSGQSM EK V HAD AERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 242 SP RS03655 WP 000613477.1	ILAE NSGQSM EK V HAD AERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 342534 ETT44 RS07705 WP 002985850.1	ILAQNA GKT I K Q I H K D A E R D Y W M S A E E T L A Y G F I D E I M E N N E L K .	196
NTDB id 446 SpyM6JRS4 RS01695 WP 002985850.1	ILAQNA GKT I K Q I H K D A E R D Y W M S A E E T L A Y G F I D E I M E N N E L K .	196
NTDB id 445 SPYM3 RS01720 WP 002985850.1	ILAQNA GKT I K Q I H K D A E R D Y W M S A E E T L A Y G F I D E I M E N N E L K .	196
NTDB id 376 SMU RS07595 WP 255262720.1	ILSDNSGKTIKQIHKDAERDYWMDAKETLKYGFIDEIMENNELK.	204
NTDB id 1223 Cj0192c YP 002343650.1	ILAKNTKQKVAKIAKDTERDFFMSAQEAK EYGLIDKVL EKSFK.	194
NTDB id 86 BSU 34540 NP 391334.1	VLAERTGQPLEVIERDTRDNEKSAEEALEYGLIDKILTHTEDKK	197
consensus	*!*****! ****! ****! !*****! !*!***** ** *	

 non conserved
 similar
 $\geq 50\%$ conserved