

NTDB id	602 KW2 RS03060 WP 011834647.1	MGYLVPVTVIEQSSRGERAYDIYSRLKDRIIIMLTGPPVEDGMANSIIAQLFLDQAQN	TCKDIYLYVNTPGGSVSAG	75
NTDB id	611 V4T04 RS07970 WP 003129593.1	MGYLVPVTVIEQSSRGERAYDIYSRLKDRIIIMLTGPPVEDGMANSIIAQLFLDQAQN	TCKDIYLYVNTPGGSVSAG	75
NTDB id	330 STU RS11365 WP 002949732.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPTEEDNMANSIIAQLFLDQAQN	TCKDIYLYVNTPGGSVSAG	72
NTDB id	298 STER RS01925 WP 002949732.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPTEEDNMANSIIAQLFLDQAQN	TCKDIYLYVNTPGGSVSAG	72
NTDB id	282 KZH43 RS03215 WP 000613477.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPPVEDNMANSVIAQLFLDAQDS	TCKDIYLYVNTPGGSVSAG	72
NTDB id	246 SPD RS03495 WP 000613477.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPPVEDNMANSVIAQLFLDAQDS	TCKDIYLYVNTPGGSVSAG	72
NTDB id	244 SPR RS03300 WP 000613477.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPPVEDNMANSVIAQLFLDAQDS	TCKDIYLYVNTPGGSVSAG	72
NTDB id	242 SP RS03655 WP 000613477.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPPVEDNMANSVIAQLFLDAQDS	TCKDIYLYVNTPGGSVSAG	72
NTDB id	446 SpyM6JRS4 RS01695 WP 002985850.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPPVEDNMANSVIAQLFLDQAQN	TCKDIYLYVNTPGGSVSAG	72
NTDB id	445 SPYM3 RS01720 WP 002985850.1	MIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPPVEDNMANSVIAQLFLDQAQN	TCKDIYLYVNTPGGSVSAG	72
NTDB id	376 SMU RS07595 WP 255262720.1	MELKEKYSMIPVVIEQTSGRSERSYDIYSRLKDRIIIMLTGPPVEDNMANSIIAQLFLDQAQN	TCKDIYLYINSPGGSVSAG	80
NTDB id	56599 BSU6051 RS17875 WP 003228214.1	MNLIPTVIEQTNRGERAYDIYSRLKDRIIIMLGSAIDDNVANISVSQLLFLAAEPEKEIISLINSPPGGSITAG	74	
NTDB id	86 BSU 34540 NP 391334.1	MNLIPTVIEQTNRGERAYDIYSRLKDRIIIMLGSAIDNVANISVSQLLFLAAEPEKEIISLINSPPGGSITAG	74	
NTDB id	1223 Cj0192c YP 002343650.1	MFIPPYVEIKSRGRSYDIYSRLKDRIIIMLSGEIHDELAASIVAQFLFLEADEPKETCKDIYLYINSPGGVLITSG	73	
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	ILADNSGQPPIEKVHVDAERDNWMSAQETLEYGFIDEIMTNQLK.	196
NTDB id	298	STER	RS01925	WP	002949732.1	ILADNSGQPPIEKVHVDAERDNWMSAQETLEYGFIDEIMTNQLK.	196
NTDB id	282	KZH43	RS03215	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	246	SPD	RS03495	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	244	SPR	RS03300	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	242	SP	RS03655	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	ILAAQNAGKTIKQIHKAERDYWMSAEETLAYGFIDEIMENNELK.	196
NTDB id	445	SPYM3	RS01720	WP	002985850.1	ILAAQNAGKTIKQIHKAERDYWMSAEETLAYGFIDEIMENNELK.	196
NTDB id	376	SMU	RS07595	WP	255262720.1	ILSDNSGKTIKQIHKAERDYWMDAKETLKYGFIDEIMENNELK.	204
NTDB id	56599	BSU6051	RS17875	WP	003228214.1	VLAERTGGPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	86	BSU	34540	NP	391334.1	VLAERTGGPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	1223	Cj0192c	YP	002343650.1	ILAKNITIKQKVAKIAKTDTERDFFMSAQEAKKEYGLIDKVLEKSFK..	194	
consensus						*!*****!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*	

 non conserved
 similar
 $\geq 50\%$ conserved