

NTDB id	602	KW2	RS03060	WP	011834647.1	MGYLVPTVIEQSSRGERAYDIYSRLKDKRIIMLTGPVEDGMANSIIAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	75
NTDB id	611	V4T04	RS07970	WP	003129593.1	MGYLVPTVIEQSSRGERAYDIYSRLKDKRIIMLTGPVEDGMANSIIAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	75
NTDB id	330	STU	RS11365	WP	002949732.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPIEDNMANSIIAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	72
NTDB id	298	STER	RS01925	WP	002949732.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPIEDNMANSIIAQLFLDQAQDNTKDIYLYVNTPGGSVSAG	72
NTDB id	282	KZH43	RS03215	WP	000613477.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPVEDNMANSVIAQLFLDQAQDS TKDIYLYVNTPGGSVSAG	72
NTDB id	246	SPD	RS03495	WP	000613477.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPVEDNMANSVIAQLFLDQAQDS TKDIYLYVNTPGGSVSAG	72
NTDB id	244	SPR	RS03300	WP	000613477.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPVEDNMANSVIAQLFLDQAQDS TKDIYLYVNTPGGSVSAG	72
NTDB id	242	SP	RS03655	WP	000613477.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPVEDNMANSVIAQLFLDQAQDS TKDIYLYVNTPGGSVSAG	72
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPVEDNMANSVIAQLFLDQAQDN TKDIYLYVNTPGGSVSAG	72
NTDB id	445	SPYM3	RS01720	WP	002985850.1	MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPVEDNMANSVIAQLFLDQAQDN TKDIYLYVNTPGGSVSAG	72
NTDB id	376	SMU	RS07595	WP	255262720.1	MELKEKYS MIPVVIEQTSGERSYDIYSRLKDKRIIMLTGPVEDNMANSIIAQLFLDQAQDN TKDIYLYINS PGGSVSAG	80
NTDB id	1223	Cj0192c	YP	002343650.1	MFIPYVIEKSSRGERSYDIYSRLKDKRIIMLSGEIHDELAASIVAQLLFLEAEDPTKDIYLYINSPGGVITSG	73	
NTDB id	979702	AADZ70	RS12025	WP	003151513.1	MNLIPTVIEQTNRGERAYDIYSRLKDKRIIMLGS AIDNVANSIVS QLLFLEAEDPEKDIST IYN SPGGS ITAG	74
NTDB id	86	BSU	34540	NP	391334.1	MNL IPTVIEQTNRGERAYDIYSRLKDKRIIMLGS AIDNVANSIVS QLLFLA EAED PEKEIS L YIN SPGGS ITAG	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	ILADNSGQPTEKVVHDAERDNWMSAQETLEYGFIDEIMTNQLK..	196
NTDB id	298	STER	RS01925	WP	002949732.1	ILADNSGQPTEKVVHDAERDNWMSAQETLEYGFIDEIMTNQLK..	196
NTDB id	282	KZH43	RS03215	WP	000613477.1	ILAENSQGSMKEKVHADAEARNWMSAQETLEYGFIDEIMANNSLN..	196
NTDB id	246	SPD	RS03495	WP	000613477.1	ILAENSQGSMKEKVHADAEARNWMSAQETLEYGFIDEIMANNSLN..	196
NTDB id	244	SPR	RS03300	WP	000613477.1	ILAENSQGSMKEKVHADAEARNWMSAQETLEYGFIDEIMANNSLN..	196
NTDB id	242	SP	RS03655	WP	000613477.1	ILAENSQGSMKEKVHADAEARNWMSAQETLEYGFIDEIMANNSLN..	196
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	ILAAQNAGKTIKQIHKAERDYWMSAEEETLAYGFIDEIMENNELK..	196
NTDB id	445	SPYM3	RS01720	WP	002985850.1	ILAAQNAGKTIKQIHKAERDYWMSAEEETLAYGFIDEIMENNELK..	196
NTDB id	376	SMU	RS07595	WP	255262720.1	ILSDNSGKTIKQIHKAERDYWMDAKETLKYGFIDEIMENNELK..	204
NTDB id	1223	Cj0192c	YP	002343650.1	ILAKNTTKQKVAKIAKDTERDFFMSAQEAKLEYGLIDKVLKSFK..	194	
NTDB id	979702	AADZ70	RS12025	WP	003151513.1	VLAERTTGQPLEVIERDTDRDNFCKSADEALEYGLIDKVLTRNTEDQK	198
NTDB id	86	BSU	34540	NP	391334.1	VLAERTTGQPLEVIERDTDRDNFCKSADEALEYGLIDKILTHTEDKK.	197
consensus						*!***** *!*!!*!!!*!!!!!!*!!*!!!*!!*!!*!!*	

 non conserved
 similar
 $\geq 50\%$ conserved