

[illegible]

NTDB id	602	KW2	RS03060	WP	011834647.1	LAIVDTMNFIKS	DVQTIVMGMAASMGTTIASSGT	TGKGRFMLPNAEYLIHQPMGGAGQG	TQQTDMIAIVAEQLLKRKRLEQ	155
NTDB id	611	V4T04	RS07970	WP	003129593.1	LAIVDTMNFIKS	DVQTIVMGMAASMGTTIASSGT	TGKGRFMLPNAEYLIHQPMGGTGQG	TQQTDMIAIVAEQLLKRKRLEQ	155
NTDB id	330	STU	RS11365	WP	002949732.1	LAIVDTMNFIKS	DVQTIVMGMAASMGTTIASSGT	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRNNLEQ	152
NTDB id	298	STER	RS01925	WP	002949732.1	LAIVDTMNFIKS	DVQTIVMGMAASMGTTIASSGT	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRNNLEQ	152
NTDB id	282	KZH43	RS03215	WP	000613477.1	LAIVDTMNFIKA	DVQTIVMGMAASMGTVIASSG	AKGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRNTLEK	152
NTDB id	246	SPD	RS03495	WP	000613477.1	LAIVDTMNFIKA	DVQTIVMGMAASMGTVIASSG	AKGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRNTLEK	152
NTDB id	244	SPR	RS03300	WP	000613477.1	LAIVDTMNFIKA	DVQTIVMGMAASMGTVIASSG	AKGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRNTLEK	152
NTDB id	242	SP	RS03655	WP	000613477.1	LAIVDTMNFIKA	DVQTIVMGMAASMGTVIASSG	AKGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRNTLEK	152
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	LAIVDTMNFIKA	DVQTIVMGMAASMGTVIASSG	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRHRLEK	152
NTDB id	445	SPYM3	RS01720	WP	002985850.1	LAIVDTMNFIKA	DVQTIVMGMAASMGTVIASSG	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQQTDMIAAAEHLLKTRHRLEK	152
NTDB id	376	SMU	RS07595	WP	255262720.1	LAIVDTMNFIKS	DVQTIVMGTAASMGTTIASSG	AKGKRFMLPNAEYLIHQPMGGTGGGT	TQQSDMAIAAEQLLKRKKLEK	160
NTDB id	319214	D9C09	RS22465	WP	003228214.1	MAIYDTMQFIKPKVST	ICIGMAASMGAFLLAAGEK	GKRYALPNSEVMIHQPLGGAQG..	QATEIEIAAKRIILLRLDKLNK	152
NTDB id	86	BSU	34540	NP	391334.1	MAIYDTMQFIKPKVST	ICIGMAASMGAFLLAAGEK	GKRYALPNSEVMIHQPLGGAQG..	QATEIEIAAKRIILLRLDKLNK	152
NTDB id	1223	Cj0192c	YP	002343650.1	FSAIYDTMNYIKP	DVCTICIGQAASMGAFLLSCG	AECKRFALPNSRIMIHQPLGGARG.	QATDIEIQAKEILLRLKITLND	151	
consensus						**!*****!	*!*!!!!!!*	****!!!**!!	*****!*****!	*

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 NSGQSMEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 246 SPD RS03495 WP 000613477.1	ILAE NSGQSMEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 244 SPR RS03300 WP 000613477.1	ILAE NSGQSMEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 242 SP RS03655 WP 000613477.1	ILAE NSGQSMEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id 446 SpyM6JRS4 RS01695 WP 002985850.1	ILAQNAKGTIKQIHKDAERDYWMSAEETLAYGFIDEIMENNELK.	196
NTDB id 445 SPYM3 RS01720 WP 002985850.1	ILAQNAKGTIKQIHKDAERDYWMSAEETLAYGFIDEIMENNELK.	196
NTDB id 376 SMU RS07595 WP 255262720.1	ILSDNSGKTIKQIHKDAERDYWMDAKETLKYGFIDEIMENNELK.	204
NTDB id 319214 D9C09 RS22465 WP 003228214.1	VLAERTGQPLEVIERDTRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id 86 BSU 34540 NP 391334.1	VLAERTGQPLEVIERDTRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id 1223 Cj0192c YP 002343650.1	ILAKNTIKQKVAKIAKDTERDFFMSAQEAKYGLIDKVLKSFK..	194
consensus	*!*****!*****!*****!*****!*****!*****!*****	

 non conserved
 similar
 $\geq 50\%$ conserved