

[illegible]

NTDB id	602	KW2 RS03060 WP	011834647.1	LAIVDTMNFIKS	DVQTI VMGMAASMGTTIASSGT	TGGKRFMLPNAEYL L IHQPMGGAAGQG	TQQTDMAIVAEQLLKTRKRLEQ	155
NTDB id	611	V4T04 RS07970 WP	003129593.1	LAIVDTMNFIKS	DVQTI VMGMAASMGTTIASSGT	TGGKRFMLPNAEYL L IHQPMGGTGQG	TQQTDMAIVAEQLLKTRKRLEQ	155
NTDB id	330	STU RS11365 WP	002949732.1	LAIVDTMNFIKS	DVQTI VMGMAASMGTTIASSGT	TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRNNLEQ	152
NTDB id	298	STER RS01925 WP	002949732.1	LAIVDTMNFIKS	DVQTI VMGMAASMGTTIASSGT	TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRNNLEQ	152
NTDB id	282	KZH43 RS03215 WP	000613477.1	LAIVDTMNFIKA	DVQTI VMGMAASMGTVIASSG	A TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRNTLEK	152
NTDB id	246	SPD RS03495 WP	000613477.1	LAIVDTMNFIKA	DVQTI VMGMAASMGTVIASSG	A TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRNTLEK	152
NTDB id	244	SPR RS03300 WP	000613477.1	LAIVDTMNFIKA	DVQTI VMGMAASMGTVIASSG	A TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRNTLEK	152
NTDB id	242	SP RS03655 WP	000613477.1	LAIVDTMNFIKA	DVQTI VMGMAASMGTVIASSG	A TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRNTLEK	152
NTDB id	446	SpyM6JRS4 RS01695 WP	002985850.1	LAIVDTMNFIKA	DVQTI VMGMAASMGTVIASSG	A TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRHRLK	152
NTDB id	445	SPYM3 RS01720 WP	002985850.1	LAIVDTMNFIKA	DVQTI VMGMAASMGTVIASSG	A TGGKRFMLPNAEYMIHQPMGGTGGGT	TQQTDMAIAAAEHLLKTRHRLK	152
NTDB id	376	SMU RS07595 WP	255262720.1	LAIVDTMNFIKS	DVQTI VMTAASMGTTIASSG	A TGGKRFMLPNAEYL L IHQPMGGTGGGT	TQQSDMAIAAEQLLKTRKKLEK	160
NTDB id	398331	GII77 RS18020 WP	003228214.1	MAIYDTMQFIKPVKVSTICIGMAASMGAFLLAAGEKGKRYALPNSEVMIHQPLGGAQG..QATEIEIAAKRIILLRDKLNK				152
NTDB id	86	BSU 34540 NP	391334.1	MAIYDTMQFIKPVKVSTICIGMAASMGAFLLAAGEKGKRYALPNSEVMIHQPLGGAQG..QATEIEIAAKRIILLRDKLNK				152
NTDB id	1223	Cj0192c YP	002343650.1	FSAIYDTMNVIKPDVCTTCIGQAASMGAFLLSCGAEGKRFALPNSRIMIHQPLGLGAR.G.QATDIEIQAKEILRLKITLND				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	ILADNSGQPTEKVVHDAERDNWMSAQETLEYGFIDEIMTNQLK.	196
NTDB id	298	STER	RS01925	WP	002949732.1	ILADNSGQPTEKVVHDAERDNWMSAQETLEYGFIDEIMTNQLK.	196
NTDB id	282	KZH43	RS03215	WP	000613477.1	ILAENSQGSMSEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	246	SPD	RS03495	WP	000613477.1	ILAENSQGSMSEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	244	SPR	RS03300	WP	000613477.1	ILAENSQGSMSEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	242	SP	RS03655	WP	000613477.1	ILAENSQGSMSEKVVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	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	398331	GII77	RS18020	WP	003228214.1	VLAERTTGQPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	86	BSU	34540	NP	391334.1	VLAERTTGQPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	1223	Cj0192c	YP	002343650.1	ILAKNITKQKVAKIAKTDTERDFFMSAQAKEYGLIDKVLEKSFK..	194	
consensus						*!*****!****!*!!**!!!***!!*!!!!!!*!!*!!*!!*	

 non conserved
 similar
 $\geq 50\%$ conserved