

[illegible]

NTDB id	602	KW2	RS03060	WP	011834647.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYL	LHQPMGGAG	QGTTQQTDMAI	VAEQLLKTRKRLEQ	155
NTDB id	611	V4T04	RS07970	WP	003129593.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYL	LHQPMGGTG	QGTTQQTDMAI	VAEQLLKTRKRLEQ	155
NTDB id	330	STU	RS11365	WP	002949732.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRNNLEQ	152		
NTDB id	298	STER	RS01925	WP	002949732.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRNNLEQ	152		
NTDB id	282	KZH43	RS03215	WP	000613477.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	AKGKRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRNTLEK	152		
NTDB id	246	SPD	RS03495	WP	000613477.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	AKGKRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRNTLEK	152		
NTDB id	244	SPR	RS03300	WP	000613477.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	AKGKRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRNTLEK	152		
NTDB id	242	SP	RS03655	WP	000613477.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	AKGKRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRNTLEK	152		
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRHRLEK	152		
NTDB id	445	SPYM3	RS01720	WP	002985850.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGTQQTDMAIAAAEHLLKTRHRLEK	152		
NTDB id	376	SMU	RS07595	WP	255262720.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	AKGKRFMLPNAEYL	LHQPMGGTGGGTQQSDMAIAAEQLLKTRKKLEK	160		
NTDB id	1157774	KJP47	RS18615	WP	003228214.1	MAIYDTMQFIKPKVST	ICIGMAASMGAFLLAAG	EKGKRYALPNSEVMIHQPLGGAQG..	QATEIEIAAKRIILLRLDKLNK	152				
NTDB id	86	BSU	34540	NP	391334.1	MAIYDTMQFIKPKVST	ICIGMAASMGAFLLAAG	EKGKRYALPNSEVMIHQPLGGAQG..	QATEIEIAAKRIILLRLDKLNK	152				
NTDB id	1223	Cj0192c	YP	002343650.1	FSAIYDTMNYIKPDVCT	ICIGQAASMGAFLLSCG	AECKRFALPNSRIMIHQPLGGARG..	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	ILAENSQGSMSEKVVHADAEARNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	246	SPD	RS03495	WP	000613477.1	ILAENSQGSMSEKVVHADAEARNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	244	SPR	RS03300	WP	000613477.1	ILAENSQGSMSEKVVHADAEARNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	242	SP	RS03655	WP	000613477.1	ILAENSQGSMSEKVVHADAEARNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	ILAAQNAGKTIKQIHKAEDYWMSEAETLAYGFIDEIMENNELK.	196
NTDB id	445	SPYM3	RS01720	WP	002985850.1	ILAAQNAGKTIKQIHKAEDYWMSEAETLAYGFIDEIMENNELK.	196
NTDB id	376	SMU	RS07595	WP	255262720.1	ILSDNSGKTIKQIHKAEDYWMDAKETLKYGFIDEIMENNELK.	204
NTDB id	1157774	KJP47	RS18615	WP	003228214.1	VLAERTTGQPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	86	BSU	34540	NP	391334.1	VLAERTTGQPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	1223	Cj0192c	YP	002343650.1	ILAKNITIKQKVAKIAKTDTERDFFMSAQAKEYGLIDKVLKSFK..	194	
consensus						*!*****!****!*!!**!!!***!!*!!!!!!*!!*!!*!!*!!*	

 non conserved
 similar
 $\geq 50\%$ conserved