

[illegible]

NTDB id	602	KW2	RS03060	WP	011834647.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYL	LHQPMGGAGC	QGTTQTDMAI	VAEQLL	KTRKRLEQ	155		
NTDB id	611	V4T04	RS07970	WP	003129593.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYL	LHQPMGGTGQ	GTTQTDMAI	VAEQLL	KTRKRLEQ	155		
NTDB id	330	STU	RS11365	WP	002949732.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRNNLEQ	152			
NTDB id	298	STER	RS01925	WP	002949732.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRNNLEQ	152			
NTDB id	282	KZH43	RS03215	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	TIASSGT	AKGGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRNTLEK	152			
NTDB id	246	SPD	RS03495	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	TIASSGT	AKGGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRNTLEK	152			
NTDB id	244	SPR	RS03300	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	TIASSGT	AKGGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRNTLEK	152			
NTDB id	242	SP	RS03655	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	TIASSGT	AKGGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRNTLEK	152			
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRHRLEK	152			
NTDB id	445	SPYM3	RS01720	WP	002985850.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQP	MGGTGGGT	QTTQTDMAIAAAEHLL	KTRHRLEK	152			
NTDB id	376	SMU	RS07595	WP	255262720.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	AKGGRFMLPNAEYL	LHQPMGGTGQ	GSDMAIAAEQ	LLKTRKKLEK	160			
NTDB id	397846	GII83	RS17610	WP	003228214.1	MAIYDTMQFIKPKV	STICIGMAASMG	AFL	LLAAGEKGKRYALPNSEV	MIHQPLGGAQG..QATEIEIAAKRI	LLLLRDKLNK	152					
NTDB id	86	BSU	34540	NP	391334.1	MAIYDTMQFIKPKV	STICIGMAASMG	AFL	LLAAGEKGKRYALPNSEV	MIHQPLGGAQG..QATEIEIAAKRI	LLLLRDKLNK	152					
NTDB id	1223	Cj0192c	YP	002343650.1	FSAIYDTMNYIKP	DVCTICIGQAASMG	AFL	LLSCGAEGKRFALPNSRIMIHQPLGGAR	..QATDIEIQAKEILLRL	ITILND	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	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	397846	GII83	RS17610	WP	003228214.1	VLAERTTGQPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	86	BSU	34540	NP	391334.1	VLAERTTGQPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	1223	Cj0192c	YP	002343650.1	ILAKNITIKQKVAKIAKTDTERDFFMSAQEAKKEYGLIDKVLKSFK..	194	
consensus						*!*****!****!*!!**!!!***!!*!!!!!!*!!*!!*!!*!!*	

 non conserved
 similar
 $\geq 50\%$ conserved