

[illegible]

NTDB id	602	KW2	RS03060	WP	011834647.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYL	LIHQPMGGAGQG	TQTQTDMAIVAEQLL	KTRKRLEQ	155
NTDB id	611	V4T04	RS07970	WP	003129593.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYL	LIHQPMGGTGQG	TQTQTDMAIVAEQLL	KTRKRLEQ	155
NTDB id	330	STU	RS11365	WP	002949732.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRNNLEQ	152	
NTDB id	298	STER	RS01925	WP	002949732.1	LAIVDTMNFIKS	DVQTI	VMGMAASMG	TIASSGT	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRNNLEQ	152	
NTDB id	282	KZH43	RS03215	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	VIASSG	AKGGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRNTLEK	152	
NTDB id	246	SPD	RS03495	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	VIASSG	AKGGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRNTLEK	152	
NTDB id	244	SPR	RS03300	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	VIASSG	AKGGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRNTLEK	152	
NTDB id	242	SP	RS03655	WP	000613477.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	VIASSG	AKGGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRNTLEK	152	
NTDB id	446	SpyM6JRS4	RS01695	WP	002985850.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	VIASSG	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRHRL	LEK	152
NTDB id	445	SPYM3	RS01720	WP	002985850.1	LAIVDTMNFIKA	DVQTI	VMGMAASMG	VIASSG	TGKGRFMLPNAEYMIHQPMGGTGGGT	TQTQTDMAIAAAEHLL	KTRHRL	LEK	152
NTDB id	376	SMU	RS07595	WP	255262720.1	LAIVDTMNFIKS	DVQTI	VMGTAASMG	TIASSGT	AKGGRFMLPNAEYLIHQPMGGTGGGT	TQS	DMIAIAAEQLL	KTRKKLEK	160
NTDB id	1156379	KJP69	RS17800	WP	003228214.1	MAYIDTMCFIKPKV	STICI	GMAASMG	AFLLAAGE	EKGKRYALPNSEV	MIHQPLGGAQG..	QATIEIEIAAKRI	LLLRDKLNK	152
NTDB id	86	BSU	34540	NP	391334.1	MAYIDTMCFIKPKV	STICI	GMAASMG	AFLLAAGE	EKGKRYALPNSEV	MIHQPLGGAQG..	QATIEIEIAAKRI	LLLRDKLNK	152
NTDB id	1223	Cj0192c	YP	002343650.1	FSAIYDTMNLIKPD	VCTICIGQAASMG	AFLLSGAE	EGRFALPNSRIMIHQP	LGARGAR..	QATDIEIQAKEILRL	KTII	ND	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	ILADNSGQPPIEKVHVDAERDNWMSAQETLEYGFIDEIMTNQLK.	196
NTDB id	298	STER	RS01925	WP	002949732.1	ILADNSGQPPIEKVHVDAERDNWMSAQETLEYGFIDEIMTNQLK.	196
NTDB id	282	KZH43	RS03215	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	246	SPD	RS03495	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	244	SPR	RS03300	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	196
NTDB id	242	SP	RS03655	WP	000613477.1	ILAENSQGSMSEKVHADAERDNWMSAQETLEYGFIDEIMANNSLN.	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	1156379	KJP69	RS17800	WP	003228214.1	VLAERTGGPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	86	BSU	34540	NP	391334.1	VLAERTGGPLEVIERDTDRDNFKSAEEALEYGLIDKILTHTEDKK	197
NTDB id	1223	Cj0192c	YP	002343650.1	ILAKNTIKQKVAKIAKTDTERDFFMSAQEAKFYGLIDKVLEKSFK..	194	
consensus						*!*****!*!!*!!!*!!!!*! !****!!*!*** ** *	

 non conserved
 similar
 $\geq 50\%$ conserved