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NTDB id	587 KW2 RS10850 WP 042211510.1	MQT	DILRLLRPKGKKLALIKQAKLIQLMGNLLNNGFNLGVEVINFLALS	SKLVEKEFVFKMHQGLSSGKSLSEILSLSFSKN	81		
NTDB id	22778 SPS RS00585 WP 011054115.1	MISFLRQ	DISIPSKLSLKKLSSKHQHKFIQLLANLLSTGFSAEVI	AFLKRSQQLQLLDYVLKMEESLLKGQGLADMLSGLGFSDA	85		
NTDB id	395 FSA28 RS09400 WP 257638827.1	.MTFLQQ	DISLKEKHKQKKLKIKQRKV	IQLFNNLFTSGFNLTEIIDFLERSHLLAEIYTKKMRQG	LLNGLHLAEIMSSSLGFSDD	84	
NTDB id	386 SMU RS09030 WP 255262721.1	.MTFLQQ	DISLKEKHKQKKLKIKQRKV	IQLFNNLFTSGFNLTEIIDFLERSHLLAEIYTKKMRQG	LLNGLHLAEIMSSSLGFSDD	84	
NTDB id	430 SGO RS09415 WP 012130941.1	MISFLRQ	DISILSKKQKQKLLGTSKQKQVIELFNNLFSSGFHLAET	VDFLGRSALLEKNYVQKMRQGLANGQA	FSEIMASLGFSDA	85	
NTDB id	271 KZH43 RS09260 WP 074196785.1		MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETIS	SFLDRSALLDKQCVTQMRVGLSQGKS	FSEMMESELGCSSA	79	
NTDB id	230 SPD RS09890 WP 074196785.1		MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETIS	SFLDRSALLDKQCVTQMRVGLSQGKS	FSEMMESELGCSSA	79	
NTDB id	196 SPR RS09425 WP 074196785.1		MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETIS	SFLDRSALLDKQCVTQMRVGLSQGKS	FSEMMESELGCSSA	79	
NTDB id	161 SP RS10375 WP 074196785.1		MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETIS	SFLDRSALLDKQCVTQMRVGLSQGKS	FSEMMESELGCSSA	79	
NTDB id	529 SMSK321 RS09235 WP 080550783.1		MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETIS	SFLDRSALLDKQCVTQMRTDLSQGKS	FSEMMESELGFSSA	79	
NTDB id	501 SM12261 RS00950 WP 004238896.1		MDISQVFRLKRKKLATAKQKNIITLFNNLFSSGFHLVETIS	SFLDRSALLDKQCVTQMRA	GLSQGKS	FSEMMESELGFSSA	79
consensus			* ! ! * * * * * * * * ! ! ! * * * * ! * * * ! * * * ! ! * * * ! ! * * * ! * * * ! ! * * ! * ! * * * * * * * ! * * ! * *				

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NTDB id	587 KW2 RS10850 WP 042211510.1	GALIVLLTSLSIIFKHI	SALTNFTFLVKVPLVSSFI	RLYLTAYFSREWGNLI	AQGVELRQTIELMKKQKSRI	FAEVGIKLDLGL	251
NTDB id	22778 SPS RS00585 WP 011054115.1	CSSLILLFGMVWLRWRSQ	SRLKLYSRLSRYPFLGKLL	KQYLTSYAREWGTLIGQLD	LMTILDIMAIETKSSLMKE	LAEDIRMSL	254
NTDB id	395 FSA28 RS09400 WP 257638827.1	CCFFILLVGIPTTFFSRKI	SRLHLIGFCSHLPFVGKYL	QLYLTAYYAREWGNLIGQGV	EAMQIVRLMQDKQSQLFREI	ICYDMEKAL	253
NTDB id	386 SMU RS09030 WP 255262721.1	CCFFILLVGIPTTFFSRKI	SRLHLIGFCSHLPFAGKYL	QLYLTAYYAREWGNLIGQGV	EAMQIVRLMQDKQSQLFREI	ICYDMEKAL	253
NTDB id	430 SG0 RS09415 WP 012130941.1	VLMILGLTGAIYLVFKGQKR	IPVVSFLARLPFVGSFVR	IYLTAYYAREWGNMIGGLE	LSQIFQMIEEQRSVLFEQE	IGQDLQGAL	254
NTDB id	271 KZH43 RS09260 WP 074196785.1	VGLVSVLALLALTFYKRS	SKMSVFSILARLPFIGIFV	QTYLTAYYAREWGNMISQGM	MELTQIFQMMEEQGSQLFKEV	QQDLAQTL	248
NTDB id	230 SPD RS09890 WP 074196785.1	VGLVSVLALLALTFYKRS	SKMSVFSILARLPFIGIFV	QTYLTAYYAREWGNMISQGM	MELTQIFQMMEEQGSQLFKEV	QQDLAQTL	248
NTDB id	196 SPR RS09425 WP 074196785.1	VGLVSVLALLALTFYKRS	SKMSVFSILARLPFIGIFV	QTYLTAYYAREWGNMISQGM	MELTQIFQMMEEQGSQLFKEV	QQDLAQTL	248
NTDB id	161 SP RS10375 WP 074196785.1	VGLVSVLALLALTFYKRS	SKMSVFSILARLPFIGIFV	QTYLTAYYAREWGNMISQGM	MELTQIFQMMEEQGSQLFKEV	QQDLAQTL	248
NTDB id	529 SMSK321 RS09235 WP 080550783.1	VGLVSLTLLALTFYKRS	SKMRVFSILARIPLGIFV	QTYLTAYYAREWGNMISQGM	MELTQIFQMIEEQGSQLFKEI	QMDLAQSL	248
NTDB id	501 SM12261 RS00950 WP 004238896.1	VGFLVSGVLLALTFYKRG	SKMRVFSILARIPLGIFV	QTYLTAYYAREWGNMISQGM	MELTQIFQMIEEQGSQLFKEI	IQDMAHAL	248
consensus		*****	*****	***	*****!	*!!!!!!*!*	!*!!!!!!*!*

