

logo

NTDB id 587 KW2 RS10850 WP 042211510.1	MQT	DILRLILRPKGGKLLALIKQAKLIQLMGNLLNNGFNLG	EVINFLALSKLVEKEFVFKM	59
NTDB id 49498 SALIVA RS09445 WP 148263031.1	MPVKISKATRQPAGTSSWKAWLNK	DISLKGISKGKKLQINQQLKVIQLFKQLLKAGFTL	TEIVAFLEERSHLLKESSL	80
NTDB id 395 FSA28 RS09400 WP 257638827.1	MTFLQQ	DISLKEKHKQKKLKIKKQKVIQLFNNLFTSGFN	LTEIIDFLERSHLLAEIYTKM	62
NTDB id 386 SMU RS09030 WP 255262721.1	MTFLQQ	DISLKEKHKQKKLKIKKQKVIQLFNNLFTSGFN	LTEIIDFLERSHLLAEIYTKM	62
NTDB id 430 SGO RS09415 WP 012130941.1	MISFLQQ	DISILSKQKQKKLGTSSKQKQVIELFNNLFSSG	FHLAETVDFLGRSALLEKNYVQQM	63
NTDB id 271 KZH43 RS09260 WP 074196785.1		MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSG	FHLVETISFLDRSALLDKQCVTQM	57
NTDB id 230 SPD RS09890 WP 074196785.1		MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSG	FHLVETISFLDRSALLDKQCVTQM	57
NTDB id 196 SPR RS09425 WP 074196785.1		MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSG	FHLVETISFLDRSALLDKQCVTQM	57
NTDB id 161 SP RS10375 WP 074196785.1		MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSG	FHLVETISFLDRSALLDKQCVTQM	57
NTDB id 529 SMSK321 RS09235 WP 080550783.1		MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSG	FHLVETISFLDRSALLDKQCVTQM	57
NTDB id 501 SM12261 RS00950 WP 004238896.1		MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSG	FHLVETISFLDRSALLDKQCVTQM	57
consensus		*****	*****	

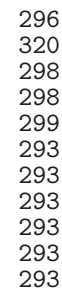
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NTDB id 587 KW2 RS10850 WP 042211510.1	HQGLSSGKSLSEILESLSFSKNVVTQLALVEVHGNLVG	TMQLVELHLKKQLKVKKKLIEVATYPVVLVFLIGIMWGLKN	139
NTDB id 49498 SALIVA RS09445 WP 148263031.1	KASLMRGDRLDQMFASVGFSDNIIVTQIALADKHGNLLG	SLTKIETYMLRMTKVRKKLMEVATYPIILLGFLILIMLGLKN	160
NTDB id 395 FSA28 RS09400 WP 257638827.1	RQGLLNLGLHLAEIMSSSLGFSDDVVTQLALSEVHGNT	TQKSLLKIESYLNLLILVKKKLVEVITYPIILLIFLVVIMLGLKN	142
NTDB id 386 SMU RS09030 WP 255262721.1	RQGLLNLGLHLAEIMSSSLGFSDDVVTQLALSEVHGNT	TQKSLLKIESYLNLLILVKKKLVEVITYPIILLIFLVVIMLGLKN	142
NTDB id 430 SGO RS09415 WP 012130941.1	RQGLANGQAFSEIMASLGFSDAVVTQLSLAELHGNLSL	ALLKIEEYLDNLAKVKKKLIEVATYPMMLLGFLVLIMIGLRN	143
NTDB id 271 KZH43 RS09260 WP 074196785.1	RVGLSQGKSFSEMMESLGCSSAIVTQLSLAEVHGNLHLS	LGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
NTDB id 230 SPD RS09890 WP 074196785.1	RVGLSQGKSFSEMMESLGCSSAIVTQLSLAEVHGNLHLS	LGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
NTDB id 196 SPR RS09425 WP 074196785.1	RVGLSQGKSFSEMMESLGCSSAIVTQLSLAEVHGNLHLS	LGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
NTDB id 161 SP RS10375 WP 074196785.1	RVGLSQGKSFSEMMESLGCSSAIVTQLSLAEVHGNLHLS	LGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
NTDB id 529 SMSK321 RS09235 WP 080550783.1	RTDLSQGKSFSEMMESLGFSSAIVTQLSLAEVHGNLHLS	LGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLSIMLGLRN	137
NTDB id 501 SM12261 RS00950 WP 004238896.1	RAGLSQGKSFSEMMESLGFSSAIVTQLSLAEVHGNLHLS	LGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
consensus	* * ! *		

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NTDB id 430 SGO RS09415 WP 012130941.1	YLLPQLSS....QNFATQLIGHLPTIFLLTVMLLGL	TGAIYLVFKGQKRIPVYSFLARLPFVGSFV	RIYLTAYYAREWG	219
NTDB id 271 KZH43 RS09260 WP 074196785.1	YLLPQLDS....SNIATQIIGNLPQIFLGMVGLVSVL	ALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWG	213
NTDB id 230 SPD RS09890 WP 074196785.1	YLLPQLDS....SNIATQIIGNLPQIFLGMVGLVSVL	ALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWG	213
NTDB id 196 SPR RS09425 WP 074196785.1	YLLPQLDS....SNIATQIIGNLPQIFLGMVGLVSVL	ALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWG	213
NTDB id 161 SP RS10375 WP 074196785.1	YLLPQLDS....SNIATQIIGNLPQIFLGMVGLVSVL	ALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWG	213
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consensus	!!!!!!**	*!*!******!***!*****	*****	*****!*****!*****!*****!*****!

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- X non conserved
- X similar
- X $\geq 50\%$ conserved