

logo

NTDB id 587 KW2 RS10850 WP 042211510.1	MQT	DILRLILRPKGKKLALIKQAKLIQLMGNLLNNGFNLGEVINFLALSKLVEKEFVFKM	59
NTDB id 240577 CG712 RS05270 WP 157769911.1	MPEKISKTI	RRPTGISSWVWLNKDASLRGISKGGKKLKI	80
NTDB id 395 FSA28 RS09400 WP 257638827.1	MTFLQ	QDISLKEKHKQKKLKIKKQKVIQLFNNLFTSGFNLTEIIDFLERSHLLAEIYTKKM	62
NTDB id 386 SMU RS09030 WP 255262721.1	MTFLQ	QDISLKEKHKQKKLKIKKQKVIQLFNNLFTSGFNLTEIIDFLERSHLLAEIYTKKM	62
NTDB id 430 SGO RS09415 WP 012130941.1	MISFLQ	QDISILSKQKQKKLGTSSKQKQVIELFNNLFSSGFHLAETVDFLGRSALLEKNYVQQM	63
NTDB id 271 KZH43 RS09260 WP 074196785.1	MDISQ	VFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 230 SPD RS09890 WP 074196785.1	MDISQ	VFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 196 SPR RS09425 WP 074196785.1	MDISQ	VFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 161 SP RS10375 WP 074196785.1	MDISQ	VFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 529 SMSK321 RS09235 WP 080550783.1	MDISQ	VFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 501 SM12261 RS00950 WP 004238896.1	MDISQ	VFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
consensus		*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!	

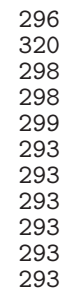
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NTDB id 587 KW2 RS10850 WP 042211510.1	HQGLSSGKSLSEIL	ESLSFSKNVVTQLALVEVHGNLVGTMQLVELHLKKQLKVKKKLIEVATYPVVLVFLIGIMWGLKN	139
NTDB id 240577 CG712 RS05270 WP 157769911.1	KESLIRGDRLDQML	ALVGFSDNIIVTQISLADKHGNLLGSLTKIETYMLRMTKVRKKLMEVATYPIILLGFLILIMLGLKN	160
NTDB id 395 FSA28 RS09400 WP 257638827.1	RQGLLNGHLHLAEI	MSSSLGFSDDVVTQLALSEVHGNLTQKSLKIESYLNLLILVKKKLVEVITYPIILLIFLVVIMLGLKN	142
NTDB id 386 SMU RS09030 WP 255262721.1	RQGLLNGHLHLAEI	MSSSLGFSDDVVTQLALSEVHGNLTQKSLKIESYLNLLILVKKKLVEVITYPIILLIFLVVIMLGLKN	142
NTDB id 430 SGO RS09415 WP 012130941.1	RQGLANGQAFSEI	MASLGFSDAVVTQLSLAELHGNLISLALLKIEEYLDNLAKVKKKLIEVATYPMMLLGFLVLIMIGLRN	143
NTDB id 271 KZH43 RS09260 WP 074196785.1	RVGLSQGKSFSEMM	ESLGCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
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NTDB id 196 SPR RS09425 WP 074196785.1	RVGLSQGKSFSEMM	ESLGCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
NTDB id 161 SP RS10375 WP 074196785.1	RVGLSQGKSFSEMM	ESLGCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRN	137
NTDB id 529 SMSK321 RS09235 WP 080550783.1	RTDLSQGKSFSEMM	ESLGFSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPIILLGFLLSIMLGLRN	137
NTDB id 501 SM12261 RS00950 WP 004238896.1	RAGLSQGKSFSEMM	ESLGFSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPIILLGFLLLIMLGLRN	137
consensus		* *!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!	

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NTDB id 587 KW2 RS10850 WP 042211510.1	YLLPQLDKG...	NNFATLLINHLPLVFFSVGALIVLLTSLSIIIFKHISALTNFTFLVKVPLVSSFIRLYLTAFFSREW	216
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NTDB id 386 SMU RS09030 WP 255262721.1	YLLPQLER...	GNFATRLISNFPILFLLGCCFFILLVGIPTFFFSRKISRLHLIGFCSHLPFAGKYLQLYLTAFFSREW	218
NTDB id 430 SGO RS09415 WP 012130941.1	YLLPQLSS...	QNFATQLIGHLPTIFLLTVMLLGLTGAIVLVFKGQKRIPVYSFLARLPFVGSFVRIYLTAYYAREWG	219
NTDB id 271 KZH43 RS09260 WP 074196785.1	YLLPQLDS...	SNIATQIIGNLPQIFLGMVGLVSVLALLALTFYKRSSKMSVFSILARLPFIFIGFVQTYLTAYYAREWG	213
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NTDB id 196 SPR RS09425 WP 074196785.1	YLLPQLDS...	SNIATQIIGNLPQIFLGMVGLVSVLALLALTFYKRSSKMSVFSILARLPFIFIGFVQTYLTAYYAREWG	213
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consensus		!!!!!!* *!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!	

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