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NTDB	id	120921 DK93 RS00660 WP 002986552.1	MINQWNNLRHKKLLKGFTLLEMLLVILVISVLMFLFVPNLSKKQDRVTETGNAAVVKLVENQAELYELSQG.SKPSSLSQ	77
NTDB	id	588 KW2 RS10845 WP 051303741.1	MSKALSLVKIIGHRKLIKCKKKQKAFTLIEMLIIVLAIIISILLIFLPVNLIKBAKVQKTGEAAVVKVVESQAQLYELDHDNDPKPNLSE	85
NTDB	id	460 GPW51 RS00790 WP 011921720.1	MKKKLVEMKVKVGFTLEVEMLVVLGGIISLLLLLFVPNLSQQKDIAIQKGDAAVVKVVESQMELYELEHDD.EBATVAD	73
NTDB	id	502 SM1261 RS00955 WP 000738641.1	MKKMMMTFLKKAQVKAFTLVEMLVVLIIISVLFLLFVPNLTKQKEAENVDKGAADVVKVVESQAELYSLEKN.DEASLSK	77
NTDB	id	272 KZH43 RS09255 WP 000738626.1	MKKMMMTFLKKAQVKAFTLVEMLVVLIIISVLFLLFVPNLTKQKEAENVDKGAADVVKVVESQAELYSLEKN.DEASLRK	77
NTDB	id	231 SPD RS09885 WP 000738626.1	MKKMMMTFLKKAQVKAFTLVEMLVVLIIISVLFLLFVPNLTKQKEAENVDKGAADVVKVVESQAELYSLEKN.DEASLRK	77
NTDB	id	197 SPR RS09420 WP 000738626.1	MKKMMMTFLKKAQVKAFTLVEMLVVLIIISVLFLLFVPNLTKQKEAENVDKGAADVVKVVESQAELYSLEKN.DEASLRK	77
NTDB	id	162 SP RS10370 WP 000738627.1	MKKMMMTFLKKAQVKAFTLVEMLVVLIIISVLFLLFVPNLTKQKEAENVDKGAADVVKVVESQAELYSLEKN.DEASLRK	77
NTDB	id	530 SMSK321 RS09230 WP 000738643.1	MKKMMMTFLKKAQVKAFTLVEMLVVLIIISVLFLLFVPNLTKQKEAENVDKGAADVVKVVESQAELYSLEKN.DEASLSK	77
NTDB	id	431 SGO RS09410 WP 012130940.1	MNKLLKKLRVKAFTLVEMLVVLIIISVLMFLFVPNLTKQKEAVSDTGNAADVVKVVESQAELYELKNTGDAQATLSK	74
NTDB	id	396 FSA28 RS09395 WP 002263437.1	MKKIKRVSVRRAFTLIEMLVVLIIISILLIFLPVNLTQKQDKVKDTGGSADVVKVVESQAELYELNKTD.SSATLSK	73
NTDB	id	387 SMU RS09025 WP 002263437.1	MKKIKRVSVRRAFTLIEMLVVLIIISILLIFLPVNLTQKQDKVKDTGGSADVVKVVESQAELYELNKTD.SSATLSK	73
consensus			** * ** * **** * * * * ** * * * * *	

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NTDB id	120921 DK93 RS00660 WP 002986552.1	LKADGSI TEKQEKAYQDYDKHKNEKARISN	108
NTDB id	588 KW2 RS10845 WP 051303741.1	LLSAGMITGKKVTAYYDDYDGNKNEQRNFDD	116
NTDB id	460 GPW51 RS00790 WP 011921720.1	LQAACKYTEKQAKQYATAKK.....	93
NTDB id	502 SM12261 RS00955 WP 000738641.1	LQADGRITEEQAKAYKEYHAKNGNSTVADV.	107
NTDB id	272 KZH43 RS09255 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id	231 SPD RS09885 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id	197 SPR RS09420 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id	162 SP RS10370 WP 000738627.1	LQADGRITEEQAKAYKEYNDKNGGANRKVND	108
NTDB id	530 SMSK321 RS09230 WP 000738643.1	LQADGRITEEQAKAYNEYAKNGGANRKVND	108
NTDB id	431 SGO RS09410 WP 012130940.1	LVAAGNISQKQADSYPKAYYGKNSETQAVAN	105
NTDB id	396 FSA28 RS09395 WP 002263437.1	LVASGDITTEQANSYKYDYAKHPSKTRHVAN	104
NTDB id	387 SMU RS09025 WP 002263437.1	LVASGDITTEQANSYKYDYAKHPSKTRHVAN	104
consensus		! * * * ! ! * * * ! * * * ! * * * * * * * *	

- X non conserved
- X similar
- X >50% conserved