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NTDB id	977804	MGAS37834	RS00670	WP	002986552.1	MINQWNNLRHKKLKGFTLLEMLLVILVISVLMLLFVPNLSKQKDRIETGNAAVVKLVENQAELYELSGG.SK	72
NTDB id	588	KW2	RS10845	WP	051303741.1	MSKALSLVKIIGHGRKLCCQKKQKAFTLIEMLIVLAITISILILLFVPNLIKKEKAQVCKTGEEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	MKKLVEMKVGFTLVEMVLVLGIISLLLLLFVPNLSQQKDAIQKKGDAADVKKVESQMELYELEHDEE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	MKKMMTFLKKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVKKVESQAELYSLDKNEED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVKKVESQAELYSLDKNEED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVKKVESQAELYSLDKNEED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVKKVESQAELYSLDKNEED	72
NTDB id	162	SP	RS10370	WP	000738627.1	MKKMMTFLKKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVKKVESQAELYSLDKNEED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	MKKMMTFLKKAKVKAFTLVEMLVLLIIISVLFLLFVPNLTKQKEAVNDKGKAADVKKVESQAELYSLDKNEED	72
NTDB id	431	SGO	RS09410	WP	012130940.1	MNKKLKLVRVKAFTLVEMLVLLIIISVLMLLFVPNLTKQKEAVSDTGNAAVVKKVESQAELYELKNTGDQ	69
NTDB id	396	FSA28	RS09395	WP	002263437.1	MKKIKRVSVRRAFTLIEMLVLLIIISILLLLFPVNLTQKQDKVKDTGGSAVVKKVESQAELYELNKTD.SS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKKIKRVSVRRAFTLIEMLVLLIIISILLLLFPVNLTQKQDKVKDTGGSAVVKKVESQAELYELNKTD.SS	68
consensus						** * ** * **** * ** * ** * ** * ** * ** * ** * ** * ** *	

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NTDB id 977804	MGAS37834	RS00670	WP 002986552.1	PSLSQLKADGS	ITEKQE	KAYQ	DYYD	KHKNEKAR	LSN	108
NTDB id 588	KW2	RS10845	WP 051303741.1	PNLSELLS	SAGM	ITCKQVT	AYDD	DYYDQ	NKNEQRNFDD	116
NTDB id 460	GPW51	RS00790	WP 011921720.1	ATVADLQAKGY	ITEKQAKQY	ATAKK				93
NTDB id 502	SM12261	RS00955	WP 000738641.1	ASLSKLQADGR	ITEEQAKAYK	EYHAK	QNTS	QTVAD		107
NTDB id 272	KZH43	RS09255	WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDK	KNGGAN	NRKVND		108
NTDB id 231	SPD	RS09885	WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDK	KNGGAN	NRKVND		108
NTDB id 197	SPR	RS09420	WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDK	KNGGAN	NRKVND		108
NTDB id 162	SP	RS10370	WP 000738627.1	ASLRKLQADGR	ITEEQAKAYK	EYHDK	KNGGAN	NRKVND		108
NTDB id 530	SMK321	RS09230	WP 000738643.1	ASLSKLQADGR	ITEEQAKAYN	EYYAK	KNGGAN	NRKVND		108
NTDB id 431	SGO	RS09410	WP 012130940.1	ATLSKLVAAGN	ISQKQAD	SYK	AYYG	KNNSETQ	AVAN	105
NTDB id 396	FSA28	RS09395	WP 002263437.1	ATLSKLVASGD	ITQE	EQAN	SYK	DYYAK	HPSKTRH	104
NTDB id 387	SMU	RS09025	WP 002263437.1	ATLSKLVASGD	ITQE	EQAN	SYK	DYYAK	HPSKTRH	104
consensus				*****!*****!	*****!*****!	*****!	*****!	*****!	*****!	

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