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NTDB id 1127834 FGL21 RS00860 WP 003055259.1	MKNQWKPLRQYQLQAFTLLVLLVLLVLSVLMLLFVPNLSKQKDRVTETGNAAVVKLVENQAELYELSQG.SK	72
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALSLVKIHGRKLCQKKQKAFTLLVLAIIISLILLFVPNLTKKKAQVQKTCEAAVVKVSVESQAQLYELDHDNDK	80
NTDB id 460 GPW51 RS00790 WP 011921720.1	MKKLVEMKVKGFTLVEMLVVLLGIISLILLFVPNLSQKKDAIQKKGDAAVVKVSVESQMELEYELDHD.EE	68
NTDB id 502 SM12261 RS00955 WP 000738641.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 231 SPD RS09885 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 197 SPR RS09420 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 162 SP RS10370 WP 000738627.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 431 SGO RS09410 WP 012130940.1	MNKLLKLRLVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTCNAAVVKVSVESQAELYELKNTGDDQ	69
NTDB id 396 FSA28 RS09395 WP 002263437.1	MKKTKRVSVRAFTLLVEMLVVLLIISLILLFVPNLTKQKDKVKDTGSAVVKVSVESQAELYELNKT.DS	68
NTDB id 387 SMU RS09025 WP 002263437.1	MKKTKRVSVRAFTLLVEMLVVLLIISLILLFVPNLTKQKDKVKDTGSAVVKVSVESQAELYELNKT.DS	68
consensus	*** * *** **** ! ! ! * ! ! ! * ! * * * ! ! * ! * ! ! ! ! ! ! * * * ! * * * * * ! * ! ! ! ! * ! * ! * ! ! * ! * *	

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NTDB id 1127834 FGL21 RS00860 WP 003055259.1	PSLSQLKTDGSIITDKQVKAYTDYYGKHRDETQKIQP	108
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELLSAGMITQKQVTAYDDYYDQNKNEGRNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYYHAKQNTSQTIVAD.	107
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKYND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKYND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKYND	108
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYYNDKNGGANRKYND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKYND	108
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYKAYYGKMNSETQAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASCDITGEQANSYKDYAKHPSKTRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASCDITGEQANSYKDYAKHPSKTRHVAN	104
consensus	***** ! *** ! ! *** ! *** ! *** * * * * * * * * * *	

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved