

logo

NTDB id 1121208 EL139 RS02665 WP 111714771.1	MKNQWKPLRQYQLQAFTLLEMLLVLLVLSVLMLLFVPNLSKQKDRVTETCNAAVVKLVENQAELYELSQG.SK	72
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALSLVKIHGRKLCQKKQKAFTLLEMLLVLAIIISLLELLFVPNLTKKKAQVQKTCEAAVVKVSVESQAQLYELDHDNDK	80
NTDB id 460 GPW51 RS00790 WP 011921720.1	MKKLVEMKVKGFTLVEMLVVLGIISLLELLFVPNLSQKKDAIQKKGDAAVVKVSVESQMELEYELDHD.EE	68
NTDB id 502 SM12261 RS00955 WP 000738641.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLELLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLELLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 231 SPD RS09885 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLELLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 197 SPR RS09420 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLELLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 162 SP RS10370 WP 000738627.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLELLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKAQVKAFTLVEMLVVLLIISVLELLFVPNLTKQKEAVNDKCKAAVVKVSVESQAELYSLDKN.ED	72
NTDB id 431 SGO RS09410 WP 012130940.1	MNKLLKLLRKAFTLVEMLVVLLIISVLMLELLFVPNLTKQKEAVSDTCNAAVVKVSVESQAELYELKNTGDDQ	69
NTDB id 396 FSA28 RS09395 WP 002263437.1	MKKTKRVSVRAFTLLEMLVVLLIISLLELLFVPNLTKQKDKVKDTGSAVVVKVSVESQAELYELNKT.DS	68
NTDB id 387 SMU RS09025 WP 002263437.1	MKKTKRVSVRAFTLLEMLVVLLIISLLELLFVPNLTKQKDKVKDTGSAVVVKVSVESQAELYELNKT.DS	68
consensus	*** * *** **** ! ! ! * ! ! ! * ! * * * ! ! * ! * ! ! ! ! ! * * * ! * * * * * ! * ! ! ! ! * ! * ! ! ! * ! *	

logo

NTDB id 1121208 EL139 RS02665 WP 111714771.1	PSLSQLKTDGSIIDKQVKAYTDYYEKHTNEKSSI.KP	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	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKYVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKYVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKYVND	108
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYYNDKNGGANRKYVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKYVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYKAYYGKNMSETQAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASCDITQEQANSYKDYAKHPSKTRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASCDITQEQANSYKDYAKHPSKTRHVAN	104
consensus	***** ! *** ! ! *** ! *** ! *** * * * * * * * *	

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