

logo

NTDB id 98 BSU 24710 NP 390351.1	MNEKCGFTLVEMLIVLFIISILLITIPNVTKHNTTQKKGCEGLQNMVKAQMTAFELDHGQQT	63
NTDB id 126950 SPS5912 RS05965 WP 015729165.1	MIQQLKLFNKKKCGFTLIEMLLVLLIISVLLILIIPNIAKQSDHIQKTGCAAQLKMDVDSQIEAYALKFNHNP	72
NTDB id 32 MW RS07995 WP 000472256.1	MFKFLKKTQAFTLIEMLLVLLIISILLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 16 SA RS07765 WP 000472256.1	MFKFLKKTQAFTLIEMLLVLLIISILLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS LVKIHGRKLCQKQKQKAFITLIEMLIVLAIISILILLFVFNLIKQKQVQKTGEAAVVKVSVESQAQLYELDHDNDK	80
NTDB id 460 GPW51 RS00790 WP 011921720.1	MKKLVEMKVKCGFTLVEMLVVLGIISLILLFVFNLSQKKDAIQKKGDAAVVKVSVESQMELEYELHDE.E	68
NTDB id 431 SGO RS09410 WP 012130940.1	MKNLKKLRVKAFTLVEMLVLLIISVLMLLFVFNLTQKQKQVSDTGNAAVVKVSVESQAELYELKNTGDQ	69
NTDB id 502 SM12261 RS00955 WP 000738641.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFVFNLTQKQKQVNDKGAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 162 SP RS10370 WP 000738627.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFVFNLTQKQKQVNDKGAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFVFNLTQKQKQVNDKGAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 231 SPD RS09885 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFVFNLTQKQKQVNDKGAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 197 SPR RS09420 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFVFNLTQKQKQVNDKGAAVVKVSVESQAELYSLDKNE.D	72
NTDB id 530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLFVFNLTQKQKQVNDKGAAVVKVSVESQAELYSLDKNE.D	72
consensus	* *** *****!!!!!!*****!! *****!!***** !* *	

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NTDB id 98 BSU 24710 NP 390351.1	PSLADLQSEGYVKKDA.VCPNGKRIIITGGEVKKVEH	98
NTDB id 126950 SPS5912 RS05965 WP 015729165.1	NTIDDLVREGYIKESQKTCKSGEEITIRDCE.AVAS	107
NTDB id 32 MW RS07995 WP 000472256.1	SSIEDLIADGFIKEAQKTCKSGETITISNCE.AVAN	103
NTDB id 16 SA RS07765 WP 000472256.1	SSIEDLIADGFIKEAQKTCKSGETITISNCE.AVAN	103
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELLSAGMITCKQVTAYDDYVDQNKNEQRNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYKAYYGKNNSETQAVAN	105
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVD.	107
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYNDKNGGANRKYND	108
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKYND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKYND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKYND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKYND	108
consensus	*** !***! *** * * * * *	

- [X] non conserved
- [X] similar
- [X] ≥ 50% conserved