

logo

NTDB id 460 GPW51 RS00790 WP 011921720.1	MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFFVFNLSQQKDAIQKKGDAAVVKVVESQMELYELEHD.EEATVADLQ	75
NTDB id 502 SM12261 RS00955 WP 000738641.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLLLFVFNLTQKKEAVNDKGKAADVKKVVESQAELYSLDKN.EDASLSKLQ	79
NTDB id 162 SP RS10370 WP 000738627.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKAADVKKVVESQAELYSLEKN.EDASLRKLQ	79
NTDB id 272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKAADVKKVVESQAELYSLEKN.EDASLRKLQ	79
NTDB id 231 SPD RS09885 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKAADVKKVVESQAELYSLEKN.EDASLRKLQ	79
NTDB id 197 SPR RS09420 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKAADVKKVVESQAELYSLEKN.EDASLRKLQ	79
NTDB id 530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLLLLFVFNLTQKKEAVNDKGKAADVKKVVESQAELYSLDKN.EDASLSKLQ	79
NTDB id 431 SGO RS09410 WP 012130940.1	MNKLLKLRVKAFTLVEMLVLLIISVLMLLFVFNLTQKKEAVSDTGNAADVKKVVESQAELYELKNTGDQATLSKLV	76
NTDB id 396 FSA28 RS09395 WP 002263437.1	MKKIKRVSVRFAFTLIEMLVLLIISILLLLFFVFNLTQKQDKVKDTGGSADVKKVVESQAELYELNKT.DSATLSKLV	75
NTDB id 387 SMU RS09025 WP 002263437.1	MKKIKRVSVRFAFTLIEMLVLLIISILLLLFFVFNLTQKQDKVKDTGGSADVKKVVESQAELYELNKT.DSATLSKLV	75
NTDB id 98 BSU 24710 NP 390351.1	MNEKGFTLVEMLIVLFIISILLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQ	70
NTDB id 305327 DWB97 RS06150 WP 037573894.1	...MKLFFKLLKENKGFLLIEMLLVLLIISVLLILIIPNIAKQSEHIQKTGCSAQLKLVESQIEAYTLKFNRKPSTIDELV	77
NTDB id 32 MW RS07995 WP 000472256.1	MFKFLKKTQAFTLIEMLVLLIISLILLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIIEDLI	75
NTDB id 16 SA RS07765 WP 000472256.1	MFKFLKKTQAFTLIEMLVLLIISLILLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIIEDLI	75
consensus	* **** *!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!*!!!!	

logo

NTDB id 460 GPW51 RS00790 WP 011921720.1	AKGYITEKQAKQYATAKK.....	93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ADGRITEEQAKAYKEYHAKQNTSQTIVAD.	107
NTDB id 162 SP RS10370 WP 000738627.1	ADGRITEEQAKAYKEYYNDKNGGANRKVND	108
NTDB id 272 KZH43 RS09255 WP 000738626.1	ADGRITEEQAKAYKEYYHDKNGGANRKVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ADGRITEEQAKAYKEYYHDKNGGANRKVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ADGRITEEQAKAYKEYYHDKNGGANRKVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ADGRITEEQAKAYNEYAKNGGANRKVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	AAGNISQKQADSYPKAYYGKNNSETQAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ASGDITQEQANSYKDYAKHPSKTRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ASGDITQEQANSYKDYAKHPSKTRHVAN	104
NTDB id 98 BSU 24710 NP 390351.1	SEGYVKKD.AVCPNGKRIITIGEVKVEH	98
NTDB id 305327 DWB97 RS06150 WP 037573894.1	QEGYIKENQKSCKNGDTIILRGGEAVVS.	105
NTDB id 32 MW RS07995 WP 000472256.1	ADGFIKEAQKTCKSGETITISNCEAVAN.	103
NTDB id 16 SA RS07765 WP 000472256.1	ADGFIKEAQKTCKSGETITISNCEAVAN.	103
consensus	**! ***** * * *	

- non conserved
- similar
- ≥ 50% conserved