

NTDB id	32 MW RS07995 WP 000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	16 SA RS07765 WP 000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	588 KW2 RS10845 WP 051303741.1	MSKALSLV ^K IHGRLCQKKQKAFTLIEMLLVLAIISILILLFVPNLIKEKAQVKGTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	460 GPW51 RS00790 WP 011921720.1	MKKLIVEMKVKGFTLVEMLLVLGIIISLLLLFVPNLSQQKDIAIKKGDAAVVKVVESQMELYELEHDEE	68
NTDB id	502 SM12261 RS00955 WP 000738641.1	MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	231 SPD RS09885 WP 000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	197 SPR RS09420 WP 000738626.1	MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	162 SP RS10370 WP 000738627.1	MKKMMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKAKVKAFTLVEMLVLLIISVLLLFVPNLTKQKEAVNDKGKAAAVKVVSQAELYSLDKNEED	72
NTDB id	431 SGO RS09410 WP 012130940.1	MNKLLKLRVKAFTLVEMLVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVSQAELYELKNITGDQ	69
NTDB id	25888 CW339 RS08970 WP 002946126.1	MKLMLKKLNNAVKLRAFTLIEMLVLLIISILLLLFVPNLSKQKDSVKETGNAAVVKVVDSPAELYEMKNNK.T	72
NTDB id	396 FSA28 RS09395 WP 002263437.1	MKKI ^R KRVSVRAFTLIEMLVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNKTD.S	68
NTDB id	387 SMU RS09025 WP 002263437.1	MKKI ^R KRVSVRAFTLIEMLVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNKTD.S	68
consensus		* * * * *	

NTDB id	32 MW RS07995 WP 000472256.1	SSI E DLIADGFIKEAQKTKSGETIT I .SNGEAVAN	103
NTDB id	16 SA RS07765 WP 000472256.1	SSI E DLIADGFIKEAQKTKSGETIT I .SNGEAVAN	103
NTDB id	588 KW2 RS10845 WP 051303741.1	PNLSELLSACMITQKVTA YDDYDQ NKNEQRNFDD	116
NTDB id	460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB id	502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYHYAKNNTSQTVAD.	107
NTDB id	272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN D	108
NTDB id	231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN D	108
NTDB id	197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVN D	108
NTDB id	162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVN D	108
NTDB id	530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKVN D	108
NTDB id	431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYPKAYYGKNNSETQAVAN	105
NTDB id	25888 CW339 RS08970 WP 002946126.1	ASLAALVSEGGITQKQADSYN D YAKHGGSRSVAN	108
NTDB id	396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQEANSYKDYYAKHPSKTRHVAN	104
NTDB id	387 SMU RS09025 WP 002263437.1	ATLSKLVASGITQEANSYKYDYAKHPSKTRHVAN	104
consensus		*** *! *** ! ****! ***** ** *	

-  non conserved
-  similar
-  > 50% conserved