

NTDB id	32 MW RS07995 WP 000472256.1	MFKFL..KKTKQAFTLIEMLVLLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	16 SA RS07765 WP 000472256.1	MFKFL..KKTKQAFTLIEMLVLLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	588 KW2 RS10845 WP 051303741.1	MSKALSLV K IHGRLCKKKQKAFTL IEMLVLAIIISILIFLFVPNLTK E AQVQKTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	460 GPW51 RS00790 WP 011921720.1	MKKLVEMKVKGFTLVEMLVVLGIISLLLLFVPNLSSQKDIAQCKKGDAAVVKVVSQMELYEHL EE	68
NTDB id	502 SM12261 RS00955 WP 000738641.1	MKKMMTFLKKA V KVAFTLVEMLVLLLIISVLLLFVPNLTKQEAVNDKGA A AVVKVVSQAELYSL E KN ED	72
NTDB id	272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKA V KVAFTLVEMLVLLLIISVLFLLFVPNLTKQEAVNDKGA A AVVKVVSQAELYSL E KN ED	72
NTDB id	231 SPD RS09885 WP 000738626.1	MKKMMTFLKKA V KVAFTLVEMLVLLLIISVLFLLFVPNLTKQEAVNDKGA A AVVKVVSQAELYSL E KN ED	72
NTDB id	197 SPR RS09420 WP 000738626.1	MKKMMTFLKKA V KVAFTLVEMLVLLLIISVLFLLFVPNLTKQEAVNDKGA A AVVKVVSQAELYSL E KN ED	72
NTDB id	162 SP RS10370 WP 000738627.1	MKKMMTFLKKA V KVAFTLVEMLVLLLIISVLFLLFVPNLTKQEAVNDKGA A AVVKVVSQAELYSL E KN ED	72
NTDB id	530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKA V KVAFTLVEMLVLLLIISVLLLFVPNLTKQEAVNDKGA A AVVKVVSQAELYSL E KN ED	72
NTDB id	431 SGO RS09410 WP 012130940.1	MNKLLKLRVKAFTLVEMLVLLLIISVLMLLFVPNLTKQEAVSDTGNA A AVVKVVSQAELYELKNTGDQ	69
NTDB id	1132025 H1W80 RS08450 WP 002946126.1	MKLMLKKLNAV K LRAFTL IEMLVLLLIISILLFFVPNL S KQKDSVKETGNAAVVKVVD S QAELYEL N KN KT	72
NTDB id	396 FSA28 RS09395 WP 002263437.1	MKKIKRVSVRAFTL IEMLVLLLIISILLFFVPNLTKQKDKVKDTGG S AVVKVVSQAELYEL N KT DS	68
NTDB id	387 SMU RS09025 WP 002263437.1	MKKIKRVSVRAFTL IEMLVLLLIISILLFFVPNLTKQKDKVKDTGG S AVVKVVSQAELYEL N KT DS	68
consensus		** * * * * ** * * * * * * * * * * * * * * * * * * * * * * * * *	

NTDB id 32	MW RS07995 WP 000472256.1	SSIEDLIADGFIKEAQKTC	KSGETITI.SNGEAVAN	103
NTDB id 16	SA RS07765 WP 000472256.1	SSIEDLIADGFIKEAQKTC	KSGETITI.SNGEAVAN	103
NTDB id 588	KW2 RS10845 WP 051303741.1	PNLSELLSAGMITCKQVTA	YDDYDQNKNEQRNFDD	116
NTDB id 460	GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQ	YATAKK.....	93
NTDB id 502	SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAY	KEYHAKNNTSQTIVAD.	107
NTDB id 272	KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAY	KEYHDKNGGANRKYND	108
NTDB id 231	SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAY	KEYHDKNGGANRKYND	108
NTDB id 197	SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAY	KEYHDKNGGANRKYND	108
NTDB id 162	SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAY	KEYNDKNGGANRKYND	108
NTDB id 530	SM321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAY	NEYAKNGGANRKYND	108
NTDB id 431	SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADS	YKAYGKNINSETQAVAN	105
NTDB id 1132025	H1W80 RS08450 WP 002946126.1	ASLAALVSEGGITCKQADS	YNDYYAKHGGESRSVAN	108
NTDB id 396	FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQEQANS	YKDYAKHPSKTRHVN	104
NTDB id 387	SMU RS09025 WP 002263437.1	ATLSKLVASGDITQEQANS	YKDYAKHPSKTRHVN	104
consensus		*** * ! * * ! ! * * ! * * * * * *		

- X non conserved
- X similar
- X $\geq 50\%$ conserved