

NTDB id 32	MW RS07995 WP 000472256.1	MFKFL..KKTQAFTLIEMLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 16	SA RS07765 WP 000472256.1	MFKFL..KKTQAFTLIEMLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 588	KW2 RS10845 WP 051303741.1	MSKALSLV K IHGRKLCQKKQKAFTLIEMLVLAIIISILILLLFVPNLIKEKAQVKQTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id 460	GPW51 RS00790 WP 011921720.1	MKKLVEMVKV G FTLVEMLVVLGGIISLLLFLFVPNL S QQKDAIQKKGDAAVVKVVSQMELYELEHD EE	68
NTDB id 502	SM12261 RS00955 WP 000738641.1	M K KMMTFLKKA K VKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN ED	72
NTDB id 272	KZH43 RS09255 WP 000738626.1	M K KMMTFLKKA K VKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN ED	72
NTDB id 231	SPD RS09885 WP 000738626.1	M K KMMTFLKKA K VKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN ED	72
NTDB id 197	SPR RS09420 WP 000738626.1	M K KMMTFLKKA K VKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN ED	72
NTDB id 162	SP RS10370 WP 000738627.1	M K KMMTFLKKA K VKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN ED	72
NTDB id 338417	EQH19 RS09655 WP 000738625.1	M K KMMTFLKKA K VKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN ED	72
NTDB id 530	SMSK321 RS09230 WP 000738643.1	M K KMMTFLKKA K VKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKN ED	72
NTDB id 431	SGO RS09410 WP 012130940.1	MNK L KKL R VKAFTLVEMLVLLIISVLM L LFVPNLTKQKEAVSDTGNAAVVKVVSQAELYELKNTGDQ	69
NTDB id 396	FSA28 RS09395 WP 002263437.1	M K K I KRVSV R AFTLIEMLVLLIIS I LLLLFVPNLTKQKDKVKD T GG S AVVKVVSQAELYELNKT DS	68
NTDB id 387	SMU RS09025 WP 002263437.1	M K K I KRVSV R AFTLIEMLVLLIIS I LLLLFVPNLTKQKDKVKD T GG S AVVKVVSQAELYELNKT DS	68
consensus		*** ** ***** ** ** ** **	

NTDB id 32	MW RS07995 WP 000472256.1	SSIEDLIADGF	IKEAQKTKSG	ETITIS.NGEA	VAN	103
NTDB id 16	SA RS07765 WP 000472256.1	SSIEDLIADGF	IKEAQKTKSG	ETITIS.NGEA	VAN	103
NTDB id 588	KW2 RS10845 WP 051303741.1	PNLSELLSAG	MITCKQVTA	YDDYDQNKNE	QRNFDD	116
NTDB id 460	GPW51 RS00790 WP 011921720.1	ATVADLQAKG	YITEKQAKQ	YATAKK.....		93
NTDB id 502	SM12261 RS00955 WP 000738641.1	ASLSKLQADGR	ITEEQAKAYK	EYHAKKNTS	QTVD.	107
NTDB id 272	KZH43 RS09255 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDKNGGAN	NRKVND	108
NTDB id 231	SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDKNGGAN	NRKVND	108
NTDB id 197	SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDKNGGAN	NRKVND	108
NTDB id 162	SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQAKAYK	EYHDKNGGAN	NRKVND	108
NTDB id 338417	EQH19 RS09655 WP 000738625.1	ATLSKLQCGDR	ITEEQVKAYN	EYYTKNGGAN	NRKVND	108
NTDB id 530	SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQAKAYN	EYYAKNGGAN	NRKVND	108
NTDB id 431	SGO RS09410 WP 012130940.1	ATLSKLVAAGN	ISQKKQADS	YKAYYGKNN	SETQAVAN	105
NTDB id 396	FSA28 RS09395 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAKHPSKT	RHVAN	104
NTDB id 387	SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAKHPSKT	RHVAN	104
consensus		***	*!***!	!***!*****	**	* * *

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