

NTDB id 32	MW RS07995 WP 000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 16	SA RS07765 WP 000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 588	KW2 RS10845 WP 051303741.1	MSKALSLV K IHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVKQTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id 460	GPW51 RS00790 WP 011921720.1	MKKLVEMVKV G FTLVEMLVVLGIISLLLFLFVPNL S QKQDAIQKKGDAAVVKVVSQMELYEL HD .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 403367	GL188 RS09785 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 K R VS V RAFTLIEMLVLLIISILL L LFVPNLTKQKDKVKD T GG S AVVKVVSQAELYELNKT.DS	68
NTDB id 387	SMU RS09025 WP 002263437.1	M K K I K R VS V RAFTLIEMLVLLIISILL L LFVPNLTKQKDKVKD T GG S AVVKVVSQAELYELNKT.DS	68
consensus		*** ** ***** ** ** ** **	

NTDB id 32	MW RS07995 WP 000472256.1	SSIEDLIADGFIKEAQKTKSGETITIS.NGEAVAN	103
NTDB id 16	SA RS07765 WP 000472256.1	SSIEDLIADGFIKEAQKTKSGETITIS.NGEAVAN	103
NTDB id 588	KW2 RS10845 WP 051303741.1	PNLSELLSAGMITCKQVTAAYDDYDQNKNEQRNFDD	116
NTDB id 460	GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB id 502	SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKKNTSQTVD.	107
NTDB id 272	KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRNVND	108
NTDB id 231	SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRNVND	108
NTDB id 197	SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRNVND	108
NTDB id 162	SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRNVND	108
NTDB id 403367	GL188 RS09785 WP 000738625.1	ATLSKLQCGDGRITEEQVKAYNEYYTNGGANRNVND	108
NTDB id 530	SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYYYAKNGGANRNVND	108
NTDB id 431	SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYKAYYGGKNNSSETQAVAN	105
NTDB id 396	FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVN	104
NTDB id 387	SMU RS09025 WP 002263437.1	ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVN	104
consensus		*** * ! *** ! ! *** ! **** * * * *	

- X non conserved
- X similar
- X > 50% conserved