

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	MSKALSLVKKIHGRKLCKKQKAFTLIELVLVAIISILILLFVPNLIKEKAQVKGTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	460	GPW51	RS00790	WP	011921720.1	MKKLVEMKVKCFTLVEMLVLVGIIISLLLLLFVPNLSQQKDIAIKKGDAAVVKVVSQMELYELEHDEE	68
NTDB id	502	SM12261	RS00955	WP	000738641.1	MKKMMTFLKKA KVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKNED	72
NTDB id	272	KZH43	RS09255	WP	000738626.1	MKKMMTFLKKA KVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKNED	72
NTDB id	231	SPD	RS09885	WP	000738626.1	MKKMMTFLKKA KVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKNED	72
NTDB id	197	SPR	RS09420	WP	000738626.1	MKKMMTFLKKA KVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKNED	72
NTDB id	162	SP	RS10370	WP	000738627.1	MKKMMTFLKKA KVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKNED	72
NTDB id	403443	GL189	RS09670	WP	000738625.1	MKKMMTFLKKA KVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKNED	72
NTDB id	530	SMSK321	RS09230	WP	000738643.1	MKKMMTFLKKA KVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVSQAELYSLDKNED	72
NTDB id	431	SGO	RS09410	WP	012130940.1	MNKLKCLR VKAFTLVEMLVLLIISVLM LLLFVPNLTKQKEAVSDTGNAAVVKVVSQAELYELKNTGDQ	69
NTDB id	396	FSA28	RS09395	WP	002263437.1	MKKITKRVS VRAFTLIEMLVLLIIS ILLLLFVPNLTKQKDKVKDTGGS AVVKVVSQAELYELNKTD DS	68
NTDB id	387	SMU	RS09025	WP	002263437.1	MKKITKRVS VRAFTLIEMLVLLIIS ILLLLFVPNLTKQKDKVKDTGGS AVVKVVSQAELYELNKTD DS	68
consensus						** * * * * ** * * * * *	

NTDB	id	32	MW	RS07995	WP	000472256.1	SSTEDLIADGFIKEAQKTKSGETITIS.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	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	231	SPD	RS09885	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	197	SPR	RS09420	WP	000738626.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	162	SP	RS10370	WP	000738627.1	ASLRKLQADGRITEEQAKAYKEYYHDKNGGANRKKVND	108
NTDB	id	403443	GL189	RS09670	WP	000738625.1	ATLSKLQGDGRITEEQVQKAYNEYYTNGGGANRKKVND	108
NTDB	id	530	SMSK321	RS09230	WP	000738643.1	ASLSKLQADGRITEEQAKAYNEYYYAKNGGANRKKVND	108
NTDB	id	431	SGO	RS09410	WP	012130940.1	ATLSKLVAAGNISQKQADSYKAYYGGKNNSSETQAVAN	105
NTDB	id	396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGDITQEQANSYKDYAKHPSKTRHVN	104
NTDB	id	387	SMU	RS09025	WP	002263437.1	ATLSKLVASGDITQEQANSYKDYAKHPSKTRHVN	104
consensus							*** *!***! *!***!*** ** * *	

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