

logo

NTDB id 98 BSU 24710 NP 390351.1	MNEKGF TL VEMLIVLFIISILLITIPNVTKHNQTIQKKGCEGLIQNMVKAQMTAFELDHEGQTPSLAD	68
NTDB id 619 LCA RS06470 WP 011374999.1	MKKKRKAFTLIEMVVVLAIIALLILLVAPNLGAQKQRAEKKTDQALVTTLQTQVELASDEIGHQIKSL.D	69
NTDB id 32 MW RS07995 WP 000472256.1	MFKF..LKKTQAF TL IEMLLVLLIISLLLLIIPNTIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIED	73
NTDB id 16 SA RS07765 WP 000472256.1	MFKF..LKKTQAF TL IEMLLVLLIISLLLLIIPNTIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIED	73
NTDB id 113406 MEPL RS06615 WP 015694726.1	MKKR SWH KGF TL LEMLIVLLILSVLILLFVFNLAQKQETIDQKGNEAIVKVI E AQMELYELDKN.VKPSAEQ	71
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS LV KIHGRKILCQKKQKAFTLIEMLIVLAIISILILLFVFNLIK E KAAQVQKTGEAAVVKVVESQAQLYELDHDNDKPNLSE	85
NTDB id 460 GPW51 RS00790 WP 011921720.1	MKKLVEMKVKGF TL VEMLVVLGIISLLLLLFVFNLSQKKDAIQKKGDAAVVKVVESQMELYELEHD.EEATVAD	73
NTDB id 502 SM12261 RS00955 WP 000738641.1	MKKM MT FLKKA V KVKAFTLVEMLVLLIISVLLLLFVFNLTQKKEAVNDKGKA A AVVKVVESQAELYSLDKN.EDASLSK	77
NTDB id 162 SP RS10370 WP 000738627.1	MKKM MT FLKKA V KVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKA A AVVKVVESQAELYSLEKN.EDASLRK	77
NTDB id 272 KZH43 RS09255 WP 000738626.1	MKKM MT FLKKA V KVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKA A AVVKVVESQAELYSLEKN.EDASLRK	77
NTDB id 231 SPD RS09885 WP 000738626.1	MKKM MT FLKKA V KVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKA A AVVKVVESQAELYSLEKN.EDASLRK	77
NTDB id 197 SPR RS09420 WP 000738626.1	MKKM MT FLKKA V KVKAFTLVEMLVLLIISVLFLLFVFNLTQKKEAVNDKGKA A AVVKVVESQAELYSLEKN.EDASLRK	77
NTDB id 530 SMSK321 RS09230 WP 000738643.1	MKKM MT FLKKA V KVKAFTLVEMLVLLIISVLLLLFVFNLTQKKEAVNDKGKA A AVVKVVESQAELYSLDKN.EDASLSK	77
NTDB id 431 SGO RS09410 WP 012130940.1	M N KLKL R VKAFTLVEMLVLLIISVLMLLFVFNLTQKKEAVSDTGNA A AVVKVVESQAELYELKNTGDDQATLSK	74
NTDB id 396 FSA28 RS09395 WP 002263437.1	M K KIK R VS V RAFTLIEMLVLLIISILLLLFVFNLTQKQDKVKDTGG S AVVKVVESQAELYELNKT.DSATLSK	73
NTDB id 387 SMU RS09025 WP 002263437.1	M K KIK R VS V RAFTLIEMLVLLIISILLLLFVFNLTQKQDKVKDTGG S AVVKVVESQAELYELNKT.DSATLSK	73
consensus	* ** *	

logo

NTDB id 98 BSU 24710 NP 390351.1	LQSEGYVKKDAVC.PNGKRIIITGGEVKKVEH	98
NTDB id 619 LCA RS06470 WP 011374999.1	ELTDKYISKDQLKHAK E KGITIEGGTVKQK.	99
NTDB id 32 MW RS07995 WP 000472256.1	LIADGFIKEAQKTCKSGETITI.SNGEAVAN	103
NTDB id 16 SA RS07765 WP 000472256.1	LIADGFIKEAQKTCKSGETITI.SNGEAVAN	103
NTDB id 113406 MEPL RS06615 WP 015694726.1	LLTEKYITKDQYEKYQTAKK.....	91
NTDB id 588 KW2 RS10845 WP 051303741.1	LLSAGMITQKQVTAYDDYYDQNKNEQRNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	LQAKGYITEKQAKQYATAKK.....	93
NTDB id 502 SM12261 RS00955 WP 000738641.1	LQADGRITEEQAKAYKEYHAQNTSQTVAD.	107
NTDB id 162 SP RS10370 WP 000738627.1	LQADGRITEEQAKAYKEYNDKNGGANRKVND	108
NTDB id 272 KZH43 RS09255 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	LQADGRITEEQAKAYNEYAKNGGANRKVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	LVAAGNISQKQADS Y KAYYGKNNSETQAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	LVASGDITQEANSYKDY Y AKHPSKTRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	LVASGDITQEANSYKDY Y AKHPSKTRHVAN	104
consensus	* ** * * * * * * * * * * * * *	

☒ non conserved
☒ similar
☒ ≥ 50% conserved