

logo

NTDB id 1381 A4U84 RS02010 WP 021113424.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAPRGYTTIEQPARIALDLPGVQNKLGTKNRE	0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 25178 ECH74115 RS23365 WP 000815971.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1193 PA5040 NP 253727.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAPRGYTTIEQPARIALDLPGVQNKLGTKNRE	80
NTDB id 1186 GTF74 RS01320 WP 000788426.1		0
NTDB id 1175 A1552VC RS12105 WP 000788426.1		0
NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
consensus		

logo

NTDB id 1381 A4U84 RS02010 WP 021113424.1	LSVGNTSRVTVWEKMDKRRITINLRASLITTRVEGPNFFESVLRXAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 25178 ECH74115 RS23365 WP 000815971.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1193 PA5040 NP 253727.1	LSVGNTSRVTVVEAKDRTRLIINLTALSSYTTTRVEGNNLFVVVGNSPAGASVASAAPVKASPAPASYAQPIKPKPYVPAG	160
NTDB id 1186 GTF74 RS01320 WP 000788426.1	MKNGLKT..YVAQW.....LTLWVGLALCAS.....S....MVFSAESATA	36
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQW.....LTLWVGLALCAS.....S....MVFSAESATA	36
NTDB id 1408 DSB67 RS13980 WP 010643500.1	MKMRHGLTLTPRKLTGL.....FSLWFSLLAP.....FVAAEEVQS	37
consensus		

logo

NTDB id 1381 A4U84 RS02010 WP 021113424.1	NQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	0
NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 25178 ECH74115 RS23365 WP 000815971.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1193 PA5040 NP 253727.1	RAIRNIDFQRGEKGEKNVVIDLSDPITLSPDIQEQQGKIRLDFAKTQLPDALRVRLDVKDFATPVQFVNASDAQSDRTSITI	240
NTDB id 1186 GTF74 RS01320 WP 000788426.1	NQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1175 A1552VC RS12105 WP 000788426.1	NQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
NTDB id 1408 DSB67 RS13980 WP 010643500.1	NSLKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSIDLLNTSVKDDQLYLLDVKDFSTVVESVEVFRDTSITRLVA	117
consensus		

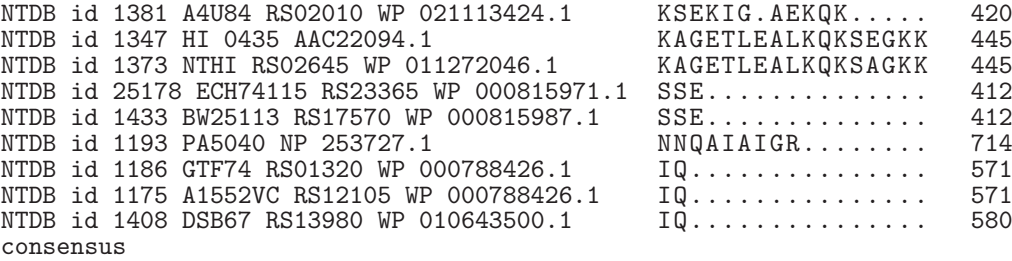
logo

NTDB id 1381 A4U84 RS02010 WP 021113424.1	TVVDGEYQHDYTLKGBYLVLSPLK ADEKPKPKPKS VLEKEGKL ISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLT	55
NTDB id 1347 HI 0435 AAC22094.1	MRYLFLFFAT.....FPVLANQQISLSIRNAPTAEIISYLAETGKNITISDEIKDTKN	67
NTDB id 1373 NTHI RS02645 WP 011272046.1	MKKYFLKCGYFLVCFCLPLI.....VFANPKTDNERFFIRLSQAPLAQTLEQLAFQDDVNLVIGDILENKIS	67
NTDB id 25178 ECH74115 RS23365 WP 000815971.1	MKKYFLKCGYFLVCFCLPLI.....VFANPKTDNECFIRLSQAPLAQTLEQLAFQDDVNLVIGDILENKIS	59
NTDB id 1433 BW25113 RS17570 WP 000815987.1	MKQWIAALIL.ML.....IPGVQAAPKPKVTLMVDDVPVAQVLQALAEQEKLNLVSPDVSQTVS	59
NTDB id 1193 PA5040 NP 253727.1	EPGSLYDYLQYQTDNRITVSIK.PMTTEDAERRKKDNFAYTGEKLSINFQDIDVRSVLQLIADFTDLNLVASDVTQGNIT	319
NTDB id 1186 GTF74 RS01320 WP 000788426.1	TVVDGEYQHDYTLKGBYLVLSPLK ADEKPKPKPKS VLEKEGKL ISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLT	194
NTDB id 1175 A1552VC RS12105 WP 000788426.1	TVVDGEYQHDYTLKGBYLVLSPLK ADEKPKPKPKS VLEKEGKL ISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLT	194
NTDB id 1408 DSB67 RS13980 WP 010643500.1	YIDEDYSHDYRLTGRYLEVKVG.KLK.ANEKAPDKSILDKEGKLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLT	195
consensus	* * ** *** *	

logo



logo



X non conserved
X similar
X ≥ 50% conserved