

logo

NTDB id 1292 VP RS12235 WP 005479693.1	...MKHSKQKK	QKGFTLIELMIVVAITGV	LAA	AAATPAYQNYVTRSEVTS	GLATVKALITPAELHYQ	ENGIAAAA	71
NTDB id 1038 HON27 RS01510 WP 000993725.1	MNA	QKGFTLIELMIVVAIIGILA	AAIATPQYQNYIAKSQVNS	GFSEISSIKTCYENAVN	EGAVPTKLSDV		69
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKA	QKGFTLIELMIVVAIIGILA	AAIATPQYQNYVARSEGAS	ALASVNPLKTTVEALS	RGWSVKSGTGTED		71
NTDB id 1197 PSJM300 03940 AFN76866.1	MKAQM	QKGFTLIELMIVVAIIGILA	AAIATPAYQDYTVRSN	AAAAALAEITPGKTCF	EQAIN.ECKTPSLTSTDE		72
NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSS	CRGFS	LIELMIVVAIIGILA	AAIATPAYQNYTVRSTAAS	ALAEITPAKAAFEQAIS	EGHTPSLVSSLP	74
NTDB id 1191 AAC28468.1 484..945()	MLKKV	QKGFTLIELMIVTAIIGILA	AAIATPAYQDYTTARA	QATEGFKATAGLQTD	IGAWRADRGSEFP	NAAALVAG	74
NTDB id 1109 OK783 RS10670 WP 157147416.1	MNTL	QKGFTLIELMIVTAIVGILA	AAVALPAYQDYTTARA	QVSEATLLAEGQKSAVT	EYYLNHGIWPKD	NTSA	71
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTL	QKGFTLIELMIVTAIVGILA	AAVALPAYQDYTTARA	QVSEATLLAEGQKSAVT	EYYLNHCKWP	ENNTSA	71
NTDB id 1188 NELON RS01755 WP 003772458.1	MKAM	QKGFTLIELMIVTAIIGILA	AAIATPAYQDYTVRSK	VSEALIAASSPKSAL	SEAFQTDGITGMTAA	AKAF	73
NTDB id 404615 IX0842 RS07630 WP 011408068.1	MKK	QKGFTLIELMIVVAIT	LAAIATPAYQDYTTRAK	LSEALTMSAPAKLA	VTE	TASSLGGVANVTLAN	70
NTDB id 1424 RS RS02795 WP 011000517.1	MKSMRHLNKR	VQKGFTLIELMIVVAIVGILA	AAIATPAYQDYTVRAR	VTEGLSLAAQAKALV	SENAANAQSDLSV		74
NTDB id 1002 ACIAD RS15100 WP 004923779.1	MNA	QKGFTLIELMIVTAIIGILA	AAIATPAYTDYTVRAR	VSEGLTAASSMKTTV	SENILNACALVAG	TPST	70
NTDB id 1274 LPP RS09530 WP 011214168.1	MR	QKGFTLIELMIVVAIIGILA	AAIATPAYQDYTTIRAR	VTEGLTLADSAKLAV	SETAITNNALPAT	QAAT	69
NTDB id 1258 GCO85 RS09745 WP 014844212.1	MR	QKGFTLIELMIVVAIIGILA	AAIATPAYQDYTTIRAR	VTEGLTLADSAKLAV	SETAITNNALPAT	QAAT	69
consensus		* * ! * ! ! ! ! ! ! ! ! *	* ! ! * * * ! ! ! ! ! *	* * ! ! ! ! ! ! ! ! *	* * ! ! ! ! ! ! ! ! *	* * ! ! ! ! ! ! ! ! *	* *

logo

NTDB id 1292 VP RS12235 WP 005479693.1	TLAQL	QVDVAANDLG	AIDSA...LAGC	...SPTLTFTF	DADSSID	GGILFTFRD	119
NTDB id 1038 HON27 RS01510 WP 000993725.1	C	FTAATSSVCATYSIDA	FTETDC...ACKIT	CTL...KG..NPKIATKI	VELTRS		116
NTDB id 1192 PAKAF RS23870 WP 024947914.1	AT...KKEVPL	QVAADANKLG	ITAKP..DPADG	...TADITLTFT	...MGGAGPKNK	CKIITLRTA	128
NTDB id 1197 PSJM300 03940 AFN76866.1	GYIG	ITDSTS.YCDVD	LD...TAADG	...H..IE.CTA...KGGNAGK	FDCKTITLNR	T.....	118
NTDB id 1204 PSJM300 03935 AFN76865.1	GFI	GIGASTN.YC	ITLI...ATATC	...S..IT.CTT...QNGMPDR	FNCKTITART		120
NTDB id 1191 AAC28468.1 484..945()	NVIFDSA	NGLDGKYFAPG	GVTVT.PDT	AINLAFDAG	ANACKITMTITPT	VI.....AASG	128
NTDB id 1109 OK783 RS10670 WP 157147416.1	G	VASA.STIKGKYVQKV	EVAKG	VVTAQMAS...TGVNKEI	QDKKLSLWA	KRQD	120
NTDB id 1119 NGFG RS11465 WP 017147169.1	G	VASPTDIKGKYVKEV	EVKNG	VVTA	MLS...SGVNNEIK	CKKLSLWG	121
NTDB id 1188 NELON RS01755 WP 003772458.1	NKTAAAGGGGAG	CAAAAGTQHASKYV	KSV.KVTEA	...SPWDITVAISATEANGIPT	GLDGNTRIFPLVNK	AVP.AATST	148
NTDB id 404615 IX0842 RS07630 WP 011408068.1	SGY	SFPGATKYV	SNV.TIADC	...TCLVTVTST.V	PNAAGTILLTPKD	VGG	116
NTDB id 1424 RS RS02795 WP 011000517.1	G	SSVF...TPTKNVANL	TIAGTGTVP	QITVTYT.T	AAGGGTLALVPT	TAAGTALPVSSAPS	131
NTDB id 1002 ACIAD RS15100 WP 004923779.1	AGSSCV	GVQEISASNATTN	VATA...TCGASSAG	QITVTMDTT	KAKGANITLTPT	YAS	125
NTDB id 1274 LPP RS09530 WP 011214168.1	GYV	SPAATPNV	VASI...AICA	..NCVITITY.TA	AAGCGTIVMTPT	TL.....QAN	113
NTDB id 1258 GCO85 RS09745 WP 014844212.1	GYV	SPAATPNV	QSI...AICA	..NCVITITY.TA	AAGCGTIVMTPT	TL.....QAN	113
consensus		*	* *	* *	* ****	* ***** *	

<0
logo

NTDB id 1292 VP RS12235 WP 005479693.1
NTDB id 1038 HON27 RS01510 WP 000993725.1
NTDB id 1192 PAKAF RS23870 WP 024947914.1
NTDB id 1197 PSJM300 03940 AFN76866.1
NTDB id 1204 PSJM300 03935 AFN76865.1
NTDB id 1191 AAC28468.1 484..945()
NTDB id 1109 OK783 RS10670 WP 157147416.1
NTDB id 1119 NGFG RS11465 WP 017147169.1
NTDB id 1188 NELON RS01755 WP 003772458.1
NTDB id 404615 IX0842 RS07630 WP 011408068.1
NTDB id 1424 RS RS02795 WP 011000517.1
NTDB id 1002 ACIAD RS15100 WP 004923779.1
NTDB id 1274 LPP RS09530 WP 011214168.1
NTDB id 1258 GC085 RS09745 WP 014844212.1
consensus



X non conserved
X similar
X ≥ 50% conserved