

logo

NTDB id 1292 VP RS12235 WP 005479693.1	...MKHSKQKKQKGF	TLIELMIVVAIIGV	LAAAAIPAYQNYVT	RSEVTSGLATVKALI	TPAELHYQ	.ENGIAAAA.....	71
NTDB id 1038 HON27 RS01510 WP 000993725.1	MNA	QKGF	TLIELMIVVAIIGILAATA	IPQYQNYIAKSQVNSGFSEISSI	KTGYENAVN	.EGAVPTKLSDV..	69
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKA	QKGF	TLIELMIVVAIIGILAATA	IPQYQNYVARSEGASALASVNPLKTTVEEALS	.RGWSVKSGTGTED		71
NTDB id 1197 PSJM300 03940 AFN76866.1	MKAQM	QKGF	TLIELMIVVAIIGILAATA	IPAYQDYTVRSNAAAAALAEITPGKIGFEQAIN	.EGKTPSLTSTDE.		72
NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSSC	RGS	LIELMIVVAIIGILAATA	IPAYQNYTVRSTAASALAEITPAKAAFEQAIS	.EGHTPSLVSSLP.		74
NTDB id 1191 AAC28468.1 484..945()	MLKKV	QKGF	TLIELMIVIAIIGILAATA	IPAYQDYTARAQATEGFKATAGLQTD	LIGAWRADRG	SFPNAAALVAG	74
NTDB id 1109 OK783 RS10670 WP 157147416.1	MNTL	QKGF	TLIELMIVIAIVGILAAV	ALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGIWPKDNTSA..			71
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTL	QKGF	TLIELMIVIAIVGILAAV	ALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHKGWPNNTSA..			71
NTDB id 1188 NELON RS01755 WP 003772458.1	MKAM	QKGF	TLIELMIVIAIIGILAATA	IPAYQDYTVRSKVSEALIAASSPKSALSEAFQTDGITGMTAAAKAF			73
NTDB id 1036168 ACA165 RS16415 WP 011408068.1	MKK	QKGF	TLIELMIVVAIIAILAAIA	IPAYQDYTTTRAKLSEALTMSAPAKLAVTETASSLGGVANVTLAN..			70
NTDB id 1424 RS RS02795 WP 011000517.1	MKSMRHLNKR	V	QKGF	TLIELMIVVAIVGILAATA	IPAYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSV.....		74
NTDB id 1002 ACIAD RS15100 WP 004923779.1	MNA	QKGF	TLIELMIVIAIIGILAATA	IPAYTDYTVRARVSEGLTAASSMKTIVSENILNAGALVAGTPTST..			70
NTDB id 1274 LPP RS09530 WP 011214168.1	MR	QKGF	TLIELMIVVAIIGILAATA	IPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT..			69
NTDB id 1258 GC085 RS09745 WP 014844212.1	MR	QKGF	TLIELMIVVAIIGILAATA	IPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAAT..			69
consensus	*****						

logo

NTDB id 1292 VP RS12235 WP 005479693.1	TLAQL	GVDVAANDLGAIDSA...	L	AGC...	SPTLTFTF.....	DADSSIDGGILTFTRD.....	119
NTDB id 1038 HON27 RS01510 WP 000993725.1	GFTAATSSVCATYSIDAFTETDG...	AGKIT..CTL...	KG..	NPKIATKIVELTRS			116
NTDB id 1192 PAKAF RS23870 WP 024947914.1	AT...KKEVPL	GVAADANKLG	TIALKP..DPADG...	TADITLTFT..	MGGAGPKNK	GKIITLTRA.....	128
NTDB id 1197 PSJM300 03940 AFN76866.1	GYIGITDSTS.YCDVDLD..	TAADG...	H..	IE.CTA...	KGGNAGKFD	GKTITLNRIT.....	118
NTDB id 1204 PSJM300 03935 AFN76865.1	GFIGIGASTN.YC	IVTLI...	ATATG...	S..	IT.CTT...	QNGMPDRFNGKTTITART.....	120
NTDB id 1191 AAC28468.1 484..945()	NVIFDSA.....	NGLDGKYFAPG.GVTVT.PDT	CAINIAFDAG.....	ANAGKMTITPTVI.....	AASG		128
NTDB id 1109 OK783 RS10670 WP 157147416.1	G	VASA.STIKGKYVQKV.EV	AKG.....	V	TAQMAS...	TGVNKEIQDKKLSLWA.....	120
NTDB id 1119 NGFG RS11465 WP 017147169.1	G	VASPTDIKGYVKEV.EVKNG....	V	TATMLS...	SGVNNEIK	GKKLSLWG.....	121
NTDB id 1188 NELON RS01755 WP 003772458.1	NTKTAAAGGGAG	CAAAAGTQHASKYVKS	VKVTEA..	SPWDITVAISATEANGIPTGLD	GNITIRLFPLVNK.AVP.AATST		148
NTDB id 1036168 ACA165 RS16415 WP 011408068.1	SGY.....	SFPGATKYVSNV.TI	ADG...	TGLVTVTST.V.....	PNAAGTILLTPKD.....	VGG	116
NTDB id 1424 RS RS02795 WP 011000517.1	GSSVF...	TPTKNVANL.TI	AGTGTVPQ	ITVITYT.T.....	AAGGGTLALVPTAAGTALPVSSAPS		131
NTDB id 1002 ACIAD RS15100 WP 004923779.1	AGSSCV	GVQEISASNATTNVATA...	TCGASSAGQ	IIVTMDTT.....	KAKGANITLTPT.....	YAS	125
NTDB id 1274 LPP RS09530 WP 011214168.1	GYV.....	SPAATPNVASI...	AIGA..	NGVITITY.TA.....	AAGGGTIVMTPTL.....	QAN	113
NTDB id 1258 GC085 RS09745 WP 014844212.1	GYV.....	SPAATPNVQSI...	AIGA..	NCVITITY.TA.....	AAGGGTIIMTPTL.....	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 1036168 ACA165 RS16415 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

