

logo

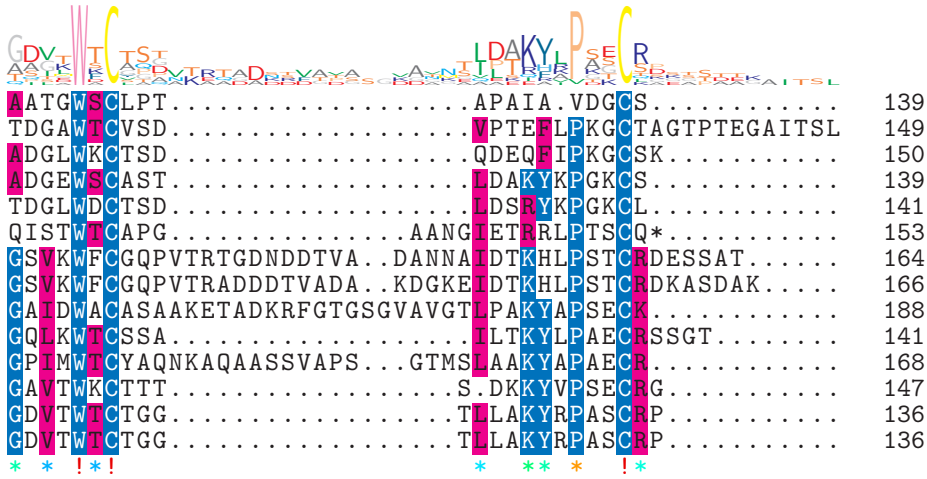
NTDB id 1292 VP RS12235 WP 005479693.1	...MKHSKQKKQKGF	TLIELMIVVAIIGV	LAAAAIPAYQNYVT	RSEVTSGLATVKALI	TPAELHYQ	.ENGIAAAA.....	71
NTDB id 1038 HON27 RS01510 WP 000993725.1	MNA	KGF	TLIELMIVVAIIGILAATAIP	QYQNYIAKSQVNSGFSEISSI	KTGYENAVN	.EGAVPTKLSDV..	69
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKA	KGF	TLIELMIVVAIIGILAATAIP	QYQNYVARSEGASALASVNPLK	TTVEEALS	.RGWSVKSGTGTED	71
NTDB id 1197 PSJM300 03940 AFN76866.1	MKAQM	KGF	TLIELMIVVAIIGILAATAIP	AYQDYTVRSNAAAA	ALAEITPGKIGFEQAIN	.EGKTPSLTSTDE.	72
NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSSC	RGS	LIELMIVVAIIGILAATAIP	AYQNYTVRSTAASA	LAIEITPAKAAFEQAIS	.EGHTPSLVSSLP.	74
NTDB id 1191 AAC28468.1 484..945()	MLKKV	KGF	TLIELMIVIAIIGILAATAIP	AYQDYTARAQATEGFKATAGLQ	TDIGAWRADRG	SFPNAAALVAG	74
NTDB id 1109 OK783 RS10670 WP 157147416.1	MNTL	KGF	TLIELMIVIAIVGILA	AAVALPAYQDYTARAQV	SEAILLAEGQKSAVTE	EYYLNHGIWPKDNTSA..	71
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTL	KGF	TLIELMIVIAIVGILA	AAVALPAYQDYTARAQV	SEAILLAEGQKSAVTE	EYYLNHKGWPNNTSA..	71
NTDB id 1188 NELON RS01755 WP 003772458.1	MKAM	KGF	TLIELMIVIAIIGILAATAIP	AYQDYTVRSKVSEALIA	AASSPKSALSEAFQTDG	ITGMTAAAKAF	73
NTDB id 413812 IX01104 RS17600 WP 011408068.1	MKK	QKGF	TLIELMIVVAIIAILAAIAIP	AYQDYTTTRAKLSEALTMSAPA	KLAVTETASSLGGVANVTLAN..		70
NTDB id 1424 RS RS02795 WP 011000517.1	MKSMRHLNKR	V	KGF	TLIELMIVVAIVGILAATAIP	AYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSV.....		74
NTDB id 1002 ACIAD RS15100 WP 004923779.1	MNA	KGF	TLIELMIVIAIIGILAATAIP	AYTDYTVRARVSEGLTAASSMKT	TVSENILNAGALVAGTPTST..		70
NTDB id 1274 LPP RS09530 WP 011214168.1	MR	KGF	TLIELMIVVAIIGILAATAIP	AYQDYTIRARVTEGLTLADSA	KLAVSETAITNNALPATQAAT..		69
NTDB id 1258 GC085 RS09745 WP 014844212.1	MR	KGF	TLIELMIVVAIIGILAATAIP	AYQDYTIRARVTEGLTLADSA	KLAVSETAITNNALPATQAAT..		69
consensus			*****	*****	*****	*	*

logo

NTDB id 1292 VP RS12235 WP 005479693.1	TLAQL	GVDVAANDLGAIDSA...	L	AGG	...	SPTL	TFTF	DADSSID	GGIL	TFTRD		119	
NTDB id 1038 HON27 RS01510 WP 000993725.1	GFTAATSSVCATYSIDAFTETDG	...	AGKIT	.CTL..KG..NPKIATKI	VELTR	S						116	
NTDB id 1192 PAKAF RS23870 WP 024947914.1	AT...KKEVPL	GVAADANKLG	TIALKP..DPADG	...	TADIT	L	TFT	..MGGAGPKNK	GKIIT	L	TRTA		128
NTDB id 1197 PSJM300 03940 AFN76866.1	GYIGITDSTS.YCDVDLD	..TAADG	...	H..IE.CTA..KGGNAGKFD	GKTIT	LNRT						118	
NTDB id 1204 PSJM300 03935 AFN76865.1	GFIGIGASTN.YC	IVT	LI...ATATG	...	S..IT	.CTT	..QNGMPDRFN	GKTIT	I	ART		120	
NTDB id 1191 AAC28468.1 484..945()	NVIFDSA	NGLDGKYFAPG.GVTVT	.PDT	CAIN	I	AFDAG	ANAGKTMTIT	PTVI	AASG			128	
NTDB id 1109 OK783 RS10670 WP 157147416.1	G	VASA.STIKGKY	VQKV.EV	AKG	V	TAQMAS	..TGVNKEIQDKKLSL	WA	KRQD			120	
NTDB id 1119 NGFG RS11465 WP 017147169.1	G	VASPTDIKGYVKEV.EV	KNG	V	TATMLS	..SGVNNEIKGKKLSL	WG	RREN				121	
NTDB id 1188 NELON RS01755 WP 003772458.1	NKTAAGGGGAG	CAAAAGTQHASKYVKS	VKVTEA	..SPWDIT	VAISATEANGIPTGLD	GN	TIRLFPLVNK	AVP	..AATST			148	
NTDB id 413812 IX01104 RS17600 WP 011408068.1	SGY	SFGAT	KYVSNV.TI	ADG	...	TGL	VT	ST.V	PNA	AGTILLTPKD	VGG	116	
NTDB id 1424 RS RS02795 WP 011000517.1	GSSVF	..TPTKNVANL.TI	AGTGTVP	Q	ITV	TYT.T	AAGGGT	LALVPTAAGTALPVSSAPS				131	
NTDB id 1002 ACIAD RS15100 WP 004923779.1	AGSSCV	GVQEISASNATTN	VATA	..TCGASSAGQ	I	IVTMDTT	KAKGANIT	L	TPT	YAS		125	
NTDB id 1274 LPP RS09530 WP 011214168.1	GYV	SPAATPNVASI	...	AIGA	..NGV	ITITY.TA	AAGGGT	I	MTPTL	QAN		113	
NTDB id 1258 GC085 RS09745 WP 014844212.1	GYV	SPAATPNVQSI	...	AIGA	..NCV	ITITY.TA	AAGGGT	I	MTPTL	QAN		113	
consensus		*	*	*	*	*	*	*	*	*	*		

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 413812 IX01104 RS17600 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