

logo

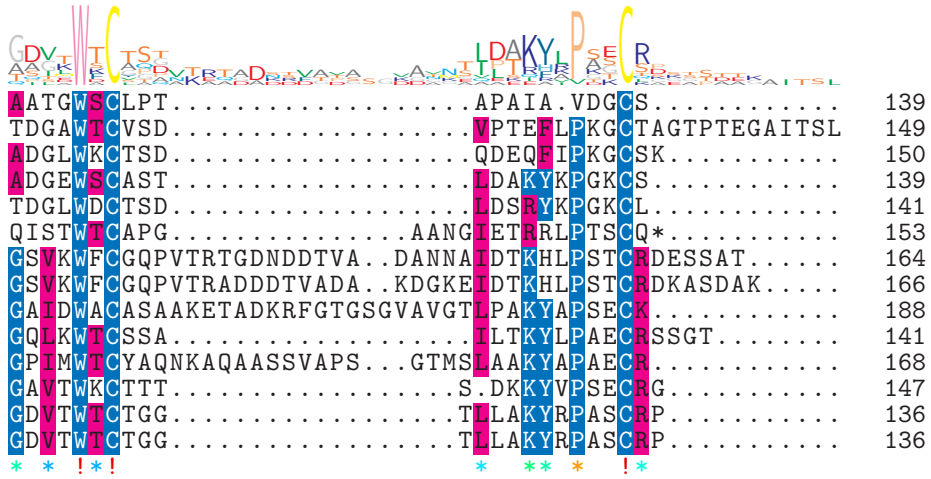
NTDB id 1292 VP RS12235 WP 005479693.1	...MKHSKQKK	QKGFTLIELMIVVAITGV	LAA	AAATPAYQNYVTRSEVTS	GLATVKALITPAELHYQ	ENGIAAAA	71
NTDB id 1038 HON27 RS01510 WP 000993725.1	MNA	QKGFTLIELMIVVAIIGIL	AAIATPQYQNYIAKSQVNS	GFSEISSIKTCYENAVN	EGAVPTKLSDV		69
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKA	QKGFTLIELMIVVAIIGIL	AAIATPQYQNYVARSEGAS	ALASVNPLKTTVEALS	RGWSVKSGTGTED		71
NTDB id 1197 PSJM300 03940 AFN76866.1	MKAQM	QKGFTLIELMIVVAIIGIL	AAIATPAYQDYTVRSN	AAAAALAEITPGKTCF	EQAIN.ECKTPSLTSTDE		72
NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSS	CFGFS	LIELMIVVAIIGIL	AAIATPAYQNYTVRSTAAS	ALAEITPAKAAFEQAIS	EGHTPSLVSSLP	74
NTDB id 1191 AAC28468.1 484..945()	MLKKV	QKGFTLIELMIVTAIIGIL	AAIATPAYQDYTTARAQATE	GFKATAGLQTD	LGAWRADRGSEFP	NAAALVAG	74
NTDB id 1109 OK783 RS10670 WP 157147416.1	MNTL	QKGFTLIELMIVTAIVGIL	AAVALPAYQDYTTARAQVSE	ATLLAEGQKSAVTET	YYLNHGIWPKD	NTSA	71
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTL	QKGFTLIELMIVTAIVGIL	AAVALPAYQDYTTARAQVSE	ATLLAEGQKSAVTET	YYLNHCKWP	ENNTSA	71
NTDB id 1188 NELON RS01755 WP 003772458.1	MKAM	QKGFTLIELMIVTAIIGIL	AAIATPAYQDYTVRSKVSE	ALIAASSPKSALSEAF	QTDGITGMTAAAKAF		73
NTDB id 1101945 Q9279 RS07710 WP 011408068.1	MKK	QKGFTLIELMIVVAITL	AAIATPAYQDYTTRAKLSE	ALTMSAPAKLAVTET	ASSLGGVANVTLAN		70
NTDB id 1424 RS RS02795 WP 011000517.1	MKSMRHLNKR	VQKGFTLIELMIVVAIVGIL	AAIATPAYQDYTVRARVTE	GLSLAAQAKALVSE	NAANAQSDLSV		74
NTDB id 1002 ACIAD RS15100 WP 004923779.1	MNA	QKGFTLIELMIVTAIIGIL	AAIATPAYTDYTVRARVSE	GLTAASSMKTTTVSEN	ILNACALVAG	TPST	70
NTDB id 1274 LPP RS09530 WP 011214168.1	MR	QKGFTLIELMIVVAIIGIL	AAIATPAYQDYTTIRARVTE	GLTLADSAKLAVSET	AITNNALPATQAAT		69
NTDB id 1258 GCO85 RS09745 WP 014844212.1	MR	QKGFTLIELMIVVAIIGIL	AAIATPAYQDYTTIRARVTE	GLTLADSAKLAVSET	AITNNALPATQAAT		69
consensus		* * ! * ! ! ! ! ! ! ! *	* ! ! * * * ! ! ! ! *	* * ! * ! ! ! ! ! *	* * ! * ! ! ! ! ! *	* * ! * ! ! ! ! ! *	* *

logo

NTDB id 1292 VP RS12235 WP 005479693.1	TLAQL	QVDVAANDLG	AIDSA...LAGC	...SPTLTFTF	DADSSID	GGITLFTTRD	119
NTDB id 1038 HON27 RS01510 WP 000993725.1	C	FTAATSSVCATYSIDA	FTETD	...ACKIT	CTL...KG..NPKIATKI	VELTRS	116
NTDB id 1192 PAKAF RS23870 WP 024947914.1	AT...KKEVPL	QVAADANKLG	ITAKP..DPADG	...TADITL	TFT...MGGAGPKNK	CKIITLRTA	128
NTDB id 1197 PSJM300 03940 AFN76866.1	GYIG	ITDSTS.YCDVD	LD..TAADG	...H..IE	CTA...KGGNAGKFD	CKTITLNR	118
NTDB id 1204 PSJM300 03935 AFN76865.1	GFI	GIGASTN.YC	ITLI...ATATC	...S..IT	CTT...QNGMPDRFN	CKTITART	120
NTDB id 1191 AAC28468.1 484..945()	NVIFDSA	NGLDGKYFAPG	GVTVT.PDT	AINLAFDAG	ANACKIT	MTITPTVI	128
NTDB id 1109 OK783 RS10670 WP 157147416.1	G	VASA.STIKGKYVQKV	EVAKG	...VVTAQMAS	...TGVNKEIQDKKLS	LWA.....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	NKTAAAGGGAG	CAAAAGTQHASKYV	KSV.KVTEA	...SPWDIT	VAISATEANGIPTGLD	CNTIRLFPLVNK	148
NTDB id 1101945 Q9279 RS07710 WP 011408068.1	SGY	SFPGATKYV	SNV.TIADC	...TCLVT	VST.V.....PNA	AGTILLTPKD	116
NTDB id 1424 RS RS02795 WP 011000517.1	G	SSVF...TPTKNVANL	TIAGTGTVP	QITVTYT.T	AAGGGT	LALVPTAAGTALPVSSAPS	131
NTDB id 1002 ACIAD RS15100 WP 004923779.1	AGSSCV	GVEISASNATTNVATA	...TCGASSAG	QITVTMDTT	KAKGANI	TLTPT	125
NTDB id 1274 LPP RS09530 WP 011214168.1	GYV	SPAATPNVASI	...AICA..NCVIT	ITY.TA	AAGGGT	IVMTPTL	113
NTDB id 1258 GCO85 RS09745 WP 014844212.1	GYV	SPAATPNVQSI	...AICA..NCVIT	ITY.TA	AAGGGT	IVMTPTL	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 1101945 Q9279 RS07710 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