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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	.EGAV	PTKLSDV..	69
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKA	QKGF	TLIELMIVVAIIGILAATA	IPQYQNYVARSEGASALASVNPLKTTVEEALS	.RGWSVKSGTG	TED		71
NTDB id 1197 PSJM300 03940 AFN76866.1	MKAQM	QKGF	TLIELMIVVAIIGILAATA	IPAYQDYTVRSNAAAAALAEITPGKIGFEQAIN	.EGKT	PSLT	STDE..	72
NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSSC	RGS	LIELMIVVAIIGILAATA	IPAYQNYTVRSTAASALAEITPAKAAFEQAIS	.EGHT	PSLV	SSLP.	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 1076829 ACJGEU RS07755 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	IPAYTDYTVRARVSEGLTAASSMKTIVSENILNAGALVAGTPT	..			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			*****	*****	*****		*	

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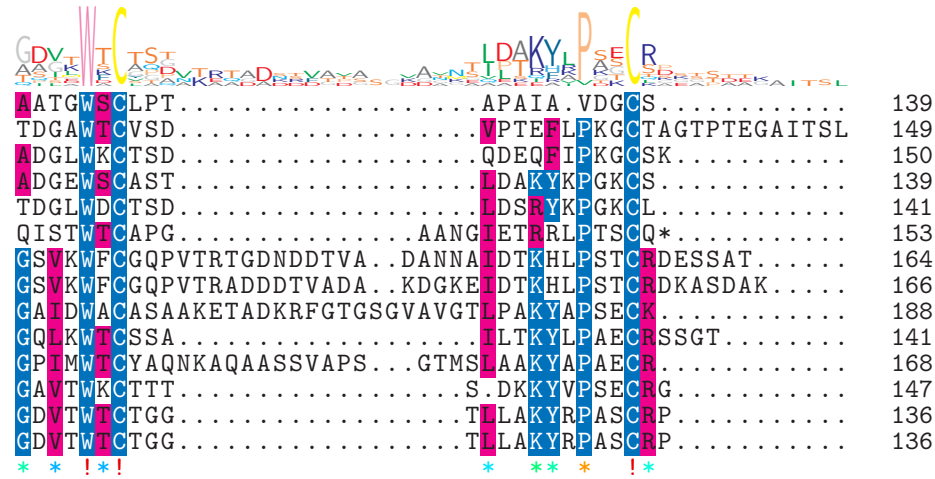
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NTDB id 1038 HON27 RS01510 WP 000993725.1	GFTAATSSVCATYSIDAFTETDG	...AGKIT	.CTL..KG..NPKIATKI	VELTRS				116
NTDB id 1192 PAKAF RS23870 WP 024947914.1	AT...KKEVPL	GVAADANKLG	TIALKP..DPADG	...TADITLTFT	..MGGAGPKNK	GKIITL	TRTA	128
NTDB id 1197 PSJM300 03940 AFN76866.1	GYIGITDSTS	YCDVDLD..TAADG	..H..IE.CTA..KGGNAGKFD	GKTITL	NRRT			118
NTDB id 1204 PSJM300 03935 AFN76865.1	GFIGIGASTN	YCI	VTLI..ATATG	..S..IT.CTT..QNGMPDRFN	GKTIT	IART		120
NTDB id 1191 AAC28468.1 484..945()	NVIFDSA	NGLDGKYFAPG	GVTVT.PDT	CAINIAFDAG	ANAGKTMTITPT	TVI	AASG	128
NTDB id 1109 OK783 RS10670 WP 157147416.1	G	VASA.STIKGKYVQKV	EVAKG	V	TAQMAS..TGVNKEIQDKKLSL	WA	KRQD	120
NTDB id 1119 NGFG RS11465 WP 017147169.1	G	VASPTDIKGYVKEV	EVKNG	V	TATMLS..SGVNNEIKGKKLSL	WG	RREN	121
NTDB id 1188 NELON RS01755 WP 003772458.1	NTKTAAGGGGAG	CAAAAGTQHASKYVKS	VKVTEA..SPWDIT	VAISATEANGIPTGLD	GN	TIRLFPLV	KNK.AVP.AATST	148
NTDB id 1076829 ACJGEU RS07755 WP 011408068.1	SGY	SFPGATKYV	SNV.TIADG	...TGLVT	VTST.V.....PNA	AGTILLTP	PKD.....VGG	116
NTDB id 1424 RS RS02795 WP 011000517.1	GSSVF	...TPTKN	VANL.TIAGTGTVP	QGITV	TYT.T.....AAG	GGTLALVP	TAAGTALPVSSAPS	131
NTDB id 1002 ACIAD RS15100 WP 004923779.1	AGSSCV	GVQEISASNATTN	VATA...TCGASSAGQ	IIVTMDTT	KAK	GANITLT	PT	125
NTDB id 1274 LPP RS09530 WP 011214168.1	GYV	SPAATPN	VASI...AIGA	..NGVITITY.TA	AAG	GGTIVM	TP	113
NTDB id 1258 GC085 RS09745 WP 014844212.1	GYV	SPAATPN	VQSI...AIGA	..NCVITITY.TA	AAG	GGTIIM	TP	113
consensus			*	*	*	*	*	

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NTDB id 1038 HON27 RS01510 WP 000993725.1
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NTDB id 1274 LPP RS09530 WP 011214168.1
NTDB id 1258 GC085 RS09745 WP 014844212.1
consensus

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- non conserved
- similar
- $\geq 50\%$ conserved