

logo

NTDB id 1292 VP RS12235 WP 005479693.1	...MKHSKQKKKQKGF	71
NTDB id 1038 HON27 RS01510 WP 000993725.1	MNAKQGF	69
NTDB id 1192 PAKAF RS23870 WP 024947914.1	MKAQKGF	71
NTDB id 1197 PSJM300 03940 AFN76866.1	MKAQMKG	72
NTDB id 1204 PSJM300 03935 AFN76865.1	MKPVKSSCRG	74
NTDB id 1109 OK783 RS10670 WP 157147416.1	MNTLKQGF	73
NTDB id 1119 NGFG RS11465 WP 017147169.1	MNTLKQGF	73
NTDB id 1191 AAC28468.1 484..945()	MLKKVKQGF	74
NTDB id 347675 EYR26 RS17290 WP 053502729.1	MKKQGF	72
NTDB id 1424 RS RS02795 WP 011000517.1	MKSMRHLNKRVKQGF	74
NTDB id 1002 ACIAD RS15100 WP 004923779.1	MNAKQGF	72
NTDB id 1274 LPP RS09530 WP 011214168.1	MRQKGF	71
NTDB id 1258 GC085 RS09745 WP 014844212.1	MRQKGF	71
consensus	*****	

logo

NTDB id 1292 VP RS12235 WP 005479693.1	...TLAQLGVDVAANDLGAIDSA.....LAGGSPTLTF	126
NTDB id 1038 HON27 RS01510 WP 000993725.1	GFTAATSSVCATYSIDA..FTETDGACKIT.CTLKG..NPKIATKIVELT.....RS.TDGAWTC	123
NTDB id 1192 PAKAF RS23870 WP 024947914.1	ATKKEVPLGVAADANKLGTIAIKP....DPADGTADITLTFTMGGAGPKNKGKIITLT.....RTAADGLWKC	135
NTDB id 1197 PSJM300 03940 AFN76866.1	GYIGITDSTS.YCDVDLD....TAADGH..IE.CTAKGGNAGKFDGKTITLN.....RT.ADGEWSC	125
NTDB id 1204 PSJM300 03935 AFN76865.1	GFIGIGASTN.YCTVTLI....ATATGS..IT.CTTQNGMPDRFNGKTTITIA.....RT.TDGLWDC	127
NTDB id 1109 OK783 RS10670 WP 157147416.1	A.....SA.STIKGKYVQKVEVAKG....VVTQAQMASTGVNKEIQDKKLSLW.....AKRQDGSVKWFC	127
NTDB id 1119 NGFG RS11465 WP 017147169.1	A.....SPPTDIKGKYVKEVEVKNG....VVTATMLSSGVNNEIKGKKLSLW.....GRRENCVSKWFC	128
NTDB id 1191 AAC28468.1 484..945()	N.....VIF.DSANGLDGKYFAPGG.VTVTPDTCAINIAFDA....GANAGKTMITITPTVIA.....ASGQISTWTC	135
NTDB id 347675 EYR26 RS17290 WP 053502729.1	Y.....AFPGQ..TKYVSGITIASG....TGKVTVTST....VPNATGDIVLTPTD.....VGGGQLKWTC	123
NTDB id 1424 RS RS02795 WP 011000517.1	GSSVF....TP.TKNVANLTIAGTGTVPGQITVYTT....TAAGGTLALVPTAAGTALPVSSAPSGPIMWTC	138
NTDB id 1002 ACIAD RS15100 WP 004923779.1	SS.CVGVEI.SASNA..TTNVATA..TCGASSAGQIIVTMDT....TKAKGANITLTPT.....YASGAVTWKC	132
NTDB id 1274 LPP RS09530 WP 011214168.1	V.....SPAA..TPNVASI..AIGA..NGVITITY.T....AAAGGGTIVMTPTL.....QANGDVTWTC	120
NTDB id 1258 GC085 RS09745 WP 014844212.1	V.....SPAA..TPNVQSI..AIGA..NGVITITY.T....AAAGGGTIIMTPTL.....QANGDVTWTC	120
consensus	*****	

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logo

NTDB id 1292 VP RS12235 WP 005479693.1	LPT.....APAIA.VDGC	139
NTDB id 1038 HON27 RS01510 WP 000993725.1	VSD.....VPTEFLPKGCTAGTPTEGAITS	149
NTDB id 1192 PAKAF RS23870 WP 024947914.1	TSD.....QDEQFIPKGC	150
NTDB id 1197 PSJM300 03940 AFN76866.1	AST.....LDAKYKPGKCS	139
NTDB id 1204 PSJM300 03935 AFN76865.1	TSD.....LDSRYKPGKCL	141
NTDB id 1109 OK783 RS10670 WP 157147416.1	GQPVTRTGDNDTIVADANNAIDTKHLPSTCRDESSAT.....	164
NTDB id 1119 NGFG RS11465 WP 017147169.1	GQPVTRADDDTIVADAKDGKEIDTKHLPSTCRDKASDAK.....	166
NTDB id 1191 AAC28468.1 484..945()	APGAAN.....GIETRRRLPTSCQ*	153
NTDB id 347675 EYR26 RS17290 WP 053502729.1	SSA.....IATKYLPAECRNSGT	141
NTDB id 1424 RS RS02795 WP 011000517.1	YAQNKAQAAS.SVAPSGTMSLAAKYAPAE	168
NTDB id 1002 ACIAD RS15100 WP 004923779.1	TTT.....S.DKKYVPSECRG	147
NTDB id 1274 LPP RS09530 WP 011214168.1	TGG.....TLLAKYRPASCRP	136
NTDB id 1258 GC085 RS09745 WP 014844212.1	TGG.....TLLAKYRPASCRP	136
consensus	*****	

 non conserved
 similar
 $\geq 50\%$ conserved