

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 149585 ACU13 RS15965 WP 047340102.1	MKKKQGF	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	MRKQGF	71
NTDB id 1258 GC085 RS09745 WP 014844212.1	MRKQGF	71
consensus	*****	

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NTDB id 1292 VP RS12235 WP 005479693.1	...TLAQLGV..DVAANDLGAIDSA.....LAGGSPTL	124
NTDB id 1038 HON27 RS01510 WP 000993725.1	GF..TAATSSVCATYSIDA..FTETDGACKIT	121
NTDB id 1192 PAKAF RS23870 WP 024947914.1	ATKKEVPLGV..AADANKLGTIAIKP...DPADGTAD	133
NTDB id 1197 PSJM300 03940 AFN76866.1	GYIGI..TDSTS.YCDVDLD....TAADGH..IE	123
NTDB id 1204 PSJM300 03935 AFN76865.1	GFIGI..GASTN.YCTVTLI....ATATGS..IT	125
NTDB id 1109 OK783 RS10670 WP 157147416.1	ASA.STIKGKYVQKVEVAKG....VVTQAQMASTG	125
NTDB id 1119 NGFG RS11465 WP 017147169.1	ASPPTDIKGKYVKEVEVKNG....VVTATMLSSG	126
NTDB id 1191 AAC28468.1 484..945()	N.....VIFDSANGLDGKYFAPGG.VTVTPDTC	133
NTDB id 149585 ACU13 RS15965 WP 047340102.1	Y....A.....FPGQ..TKYVSAVTIASG....TGKVT	121
NTDB id 1424 RS RS02795 WP 011000517.1	GSSVF...TP..TKNVANLTIAGTGTVP	136
NTDB id 1002 ACIAD RS15100 WP 004923779.1	S.....SCVGVQEISASNA..TTNVATA..TCGASSAG	130
NTDB id 1274 LPP RS09530 WP 011214168.1	V.....SPAA..TPNVASI..AIGA..NGVITITY.T...	118
NTDB id 1258 GC085 RS09745 WP 014844212.1	V.....SPAA..TPNVQSI..AIGA..NGVITITY.T...	118
consensus	*****	

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NTDB id 1292 VP RS12235 WP 005479693.1	SCLPT.....APAIA.VDGC	139
NTDB id 1038 HON27 RS01510 WP 000993725.1	TCVSD.....VPTFLPKGCTAGTPTEGAITS	149
NTDB id 1192 PAKAF RS23870 WP 024947914.1	KCTSD.....QDEQFIPKGC	150
NTDB id 1197 PSJM300 03940 AFN76866.1	SCAST.....LDAKYKPGK	139
NTDB id 1204 PSJM300 03935 AFN76865.1	DCTSD.....LDSRYKPGKCL	141
NTDB id 1109 OK783 RS10670 WP 157147416.1	FCGQPVRTGDNDTIVADANNAIDTKHLPSTCRDESSAT	164
NTDB id 1119 NGFG RS11465 WP 017147169.1	FCGQPVRTADDDTIVADAKDGKEIDTKHLPSTCRDKASDAK	166
NTDB id 1191 AAC28468.1 484..945()	TCAPGAAN.....GIETRLPTSCQ*	153
NTDB id 149585 ACU13 RS15965 WP 047340102.1	TCSSA.....IATKYLPAECRNSGT	141
NTDB id 1424 RS RS02795 WP 011000517.1	TCYAQNKAQAAS.SVAPSGTMSLAAKYAPAE	168
NTDB id 1002 ACIAD RS15100 WP 004923779.1	KCTTT.....S.DKKYVPSECRG	147
NTDB id 1274 LPP RS09530 WP 011214168.1	TCTGG.....TLLAKYRPASCRP	136
NTDB id 1258 GC085 RS09745 WP 014844212.1	TCTGG.....TLLAKYRPASCRP	136
consensus	*****	

 non conserved
 similar
 $\geq 50\%$ conserved