

logo

NTDB id 1253 GC085 RS04835 WP 011215094.1	MLRLYRKPNKGLMGVDISSTSVKLELSVKNKYNVESYALMPLPENSWEKNILNPEAVAEALERAMNLANPQTTHAAI	80
NTDB id 1102112 Q9261 RS05550 WP 011407795.1	MLKLLPKHRSITIGIDISSSAVKILEISGSGEDICVVGYYGREETPASALEGNMICKVIVVANCCKKLVERLKFTSKNVAI	79
NTDB id 1010 ACIAD RS15195 WP 004923715.1	.MGLLPKSSQSPILIGVDISSTAVKLLQLSRSGNRFRVEHYVVEPLPPNAVVEKNIVEVEAVGEATRRATNRSKSKAKNAAA	80
NTDB id 1084 FDQ49 RS11070 WP 004711654.1	MLRLYRBPKKKGLMGVDISSTSVKLELSVKNKYNQYRVESYGLVPLLENSVVEKNILNPEAVADALTRSLNLANPQAINAAI	80
NTDB id 1054 ABD1 RS15870 WP 002014066.1	MLRLYRKPNKGLMGVDISSTSVKLELSVKNKGYWVESYALMPLPENSVEKNILNPEAVAEALERAMNLANPQTTHAAI	80
consensus	***!*****!*!!!!*!!	

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NTDB id 1253 GC085 RS04835 WP 011215094.1	AVPTSAVVIHKLEMDADMDDERFVQIRYDAEYIPEDLDEVSDFEVL	160
NTDB id 1102112 Q9261 RS05550 WP 011407795.1	AVPDSAVISKVVQINEGLTDLMEELVITBADKYIPYPIDEINLDFEVLGHSAKSSSLDVLIVASRAENVNTRVEAAHR	159
NTDB id 1010 ACIAD RS15195 WP 004923715.1	AVAGSAVITIKLIPMPADLDDSDMEAQVELEATNYIPYPIEBEVNLDFEVLGMPNPSFDMVQVLLAASRSENVELRQSALEL	160
NTDB id 1084 FDQ49 RS11070 WP 004711654.1	AVPTSMVHKTIEMDADMDDEREVQIRMDAERYIPFPLDEVSLDFEVLSPILANPNRVSVLTAATRTENVDTREVLLEL	160
NTDB id 1054 ABD1 RS15870 WP 002014066.1	AVPTSTVIHKTIEMDADMDDEREVQIRVDAEQYIPFPLDEVSLDFEVLSPDRLLNPNRVNVLLVATRTENVETRVEVLEL	160
consensus	!!*!!	

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NTDB id 1253 GC085 RS04835 WP 011215094.1	AGLTPKVAQDVESYAVERAFAVSFAVDSLP	240
NTDB id 1102112 Q9261 RS05550 WP 011407795.1	AGLEPKVVDVESYAVERSVQQIAQELPASGQKQIVAIIVDIGANYTHLFLVHGMRLIFSREEKFQGMQIVEATAEYHNNL	238
NTDB id 1010 ACIAD RS15195 WP 004923715.1	CGLTAKVMDVEAFAVENAFATVASELPV.AADAVVALVDICATMTTSLVLRSCSLYSREQMFGGKQLTDEVMMRYGLTY	238
NTDB id 1084 FDQ49 RS11070 WP 004711654.1	AGLTPKVADVESYAMERAFSVFADTLPM.GVNT.VGILDIGHTMTTSLVSMQKGIITYTREQVFGGKQLTQEVQKRYCMSY	238
NTDB id 1054 ABD1 RS15870 WP 002014066.1	ADLNPKITADVESYAVERAFAVSFADSLPM.GANT.TGILDIGHTMTTSLVSMQNGKIIITYTREQVFGGKQLTLEIQSRYGLSL	238
consensus	*!!	

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NTDB id 1253 GC085 RS04835 WP 011215094.1	EEASRAKKDGS	320
NTDB id 1102112 Q9261 RS05550 WP 011407795.1	EEAGLAKRQGGLPESYEVEVLEPPFKDMLILQIKRTLQFFYSTSQHGFVDHILLAGCMARQPGLAALIQEQLGVTTVANP	318
NTDB id 1010 ACIAD RS15195 WP 004923715.1	EEAGRAKKDGS	318
NTDB id 1084 FDQ49 RS11070 WP 004711654.1	EEASRAKKDRITLPDDYEIEVLDPFLDAVQQAARSLQFFFSSSQFNEIDHILLAGGNANIPGLAKLLQQKLGYRVTIANP	318
NTDB id 1054 ABD1 RS15870 WP 002014066.1	EEASRAKKDRSLPDDYEIEVLDPFLDAVQQAARSLQFFFSSSQFNEIDHILLAGGNANIPGLAKLLQQKLGYRVTIANP	318
consensus	!!*!!	

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NTDB id 1253 GC085 RS04835 WP 011215094.1	LSHMSFGKKVNLDVINSDAPTLMVACGLALRHVE.	354
NTDB id 1102112 Q9261 RS05550 WP 011407795.1	LAQMTLGPKVQAHVIALDAPALMTATGLALRSFD.	352
NTDB id 1010 ACIAD RS15195 WP 004923715.1	FLQMEFSPQVDIKKVENDAAASLMVACGLALRSFDE	353
NTDB id 1084 FDQ49 RS11070 WP 004711654.1	FLQMGFSPQVDVQKIENDASSLMVACGLALRSFD.	352
NTDB id 1054 ABD1 RS15870 WP 002014066.1	FLQMGFSPQVDVQKIENDASSLMVACGLALRSFD.	352
consensus	***!*****!*!!!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!	

 non conserved
 similar
 $\geq 50\%$ conserved