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logo
NTDB id 1010 ACIAD RS15195 WP 004923715.1     MLRLYRRPKKGLVGVDISSTSVKVLELSVKNNQYRVESYGLVPLLENSVVEKNILNPEAVADALTRSINLANPQAINAAI   80
NTDB id 1054 ABD1 RS15870 WP 002014066.1      MLRLYRKPNKGLMGVDISSTSVKLLELSVKNGKYWVESYALMPLPENSVVEKNILNPEAVAEALERAMNLANPQTTHAAI   80
NTDB id 1084 FDQ49 RS11070 WP 004711654.1     MLRLYRKPNKGLMGVDISSTSVKLLELSVKNGKYWVESYALMPLPENSVVEKNILNPEAVAEALERAMNLANPQTTNAAI   80
NTDB id 396446 GH742 RS10600 WP 203454937.1   MLKLFKPKSRSIIGIDITSSSVKVLEISSADNKYRIEGYGAEPLPANAIEGNTVKDVEAVAMCIKRVLSKAHLSGKHAVV   80
NTDB id 1253 GC085 RS04835 WP 011215094.1     MLKLLKPKHRSILGIDISSSAVKILEISGSGEDICVEGYGREEIPASALEGNMIKDVDVVANCIKKLVERLKFTSKNVAL   80
consensus                                     !!*!**** ****!*!!*!**!!*!!*!*** ** *!*!*********************!!*** * ******* ****

logo
NTDB id 1010 ACIAD RS15195 WP 004923715.1     AVPTSMVVHKIIEMDADMTDDEREVQIRMDAERYIPFPLDEVSLDFEVMSEPLANPNRVSVLIAATRTENVDTRIEVLEL  160
NTDB id 1054 ABD1 RS15870 WP 002014066.1      AVPTSTVIHKTIEMDADMSDDEREVQIRVDAEQYIPFPLDEVSLDFEVLPDRLANPNRVNVLLVATRTENVETRVEVLEL  160
NTDB id 1084 FDQ49 RS11070 WP 004711654.1     AVPTSTVIHKTIEMDADMSDDEREVQIRVDAEQYIPFPLDEVSLDFEVLPDRLLNPNRVNVLLVATRTENVETRVEVLEL  160
NTDB id 396446 GH742 RS10600 WP 203454937.1   AVPDSAVISKVVQINDGLNDAEIEELIVMEADKYIPYPIDEINIDFEVLGHSPKNSAMLDVLIVASRAENVGSRVDAVNR  160
NTDB id 1253 GC085 RS04835 WP 011215094.1     AVPDSAVISKVVQINEGLTDLEMEELVVIEADKYIPYPIDEINLDFEVLGHSAKSSSLLDVLIVASRAENVNTRVEAAHR  160
consensus                                     !!!*! !**!*********!*!*!******!**!!!*!*!!***!!!!* * * ***** !!**!*!*!!!**!******

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NTDB id 1010 ACIAD RS15195 WP 004923715.1     AGLTPKVADVESYAMERAFSVFADTLPMGV..NTVGILDIGHTMTTLSVMQKGKIIYTREQVFGGKQLTQEVQKRYGMSY  238
NTDB id 1054 ABD1 RS15870 WP 002014066.1      ADLNPKLADVESYAVERAFSVFADSLPMGA..NTIGILDIGHTMTTLSVMQNGKIIYTREQVFGGKQLTLEIQSRYGLSL  238
NTDB id 1084 FDQ49 RS11070 WP 004711654.1     ADLNPKLADVESYAVERAFSVFADSLPMGA..NTIGILDIGHTMTTLSVMQNGKIIYTREQVFGGKQLTLEIQSRYGLSL  238
NTDB id 396446 GH742 RS10600 WP 203454937.1   AGLEAKIVDVESFAVERAAHLLAKELPAQGQDKIVAVIDIGALYTHLFVLHGMKIIFTREEEFGGKQLIDSIASQYQVSF  240
NTDB id 1253 GC085 RS04835 WP 011215094.1     AGLEPKVVDVESYAVERSVQQIAQELPASGQDKIVAIVDIGANYTHLFVLHGMRLIFSREEKFGGMQLVEAIAEHYHMNL  240
consensus                                     !*! *!**!!!!*!*!!*****!**!!*** ******!!!***!*!*!** ***!**!!**!!!*!!* *****!****

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NTDB id 1010 ACIAD RS15195 WP 004923715.1     EEAGRAKKDGSLPEDFTQEVFEPYLEALVQQAARSLQFFFSSSQYNEIDHILLAGGNANILGLSRLLQQKLGYRVTIANP  318
NTDB id 1054 ABD1 RS15870 WP 002014066.1      EEASRAKKDRSLPDDYEIEVLDPFLDAVVQQAARSLQFFFSSSQFNEIDHILLAGGNANIPGLAKLLQQKLGYRVTIANP  318
NTDB id 1084 FDQ49 RS11070 WP 004711654.1     EEASRAKKDRTLPDDYEIEVLDPFLDAVVQQAARSLQFFFSSSQFNEIDHILLAGGNANIPGLAKLLQQKLGYRVTIANP  318
NTDB id 396446 GH742 RS10600 WP 203454937.1   EQAALMKAEGNLPENYEEEVLEPFIEMILLQVKRTLQFFFSTSHHGFVDHIILAGGVARLPNLASLVQEKTGIPTSVADP  320
NTDB id 1253 GC085 RS04835 WP 011215094.1     EQAIALKNQGKLPEDYESAVLEPFKDMILLQIKRTLQFFYSTSQHGFVDHILLAGGMARQPGLAALIQEQLGVTTTVANP  320
consensus                                     !*! **!***!!**** *!**!******!**!*!!!!*!*!*****!!!*!!!!*!****!**!*!***!*****!*!

logo
NTDB id 1010 ACIAD RS15195 WP 004923715.1     FLQMEFSPQVDIKKVENDAASLMVACGLALRSFDE   353
NTDB id 1054 ABD1 RS15870 WP 002014066.1      FLQMGFSPQVDVQKIENDASSLMVACGLALRSFD.   352
NTDB id 1084 FDQ49 RS11070 WP 004711654.1     FLQMGFSPQVDVQKIENDASSLMVACGLALRSFD.   352
NTDB id 396446 GH742 RS10600 WP 203454937.1   LHEMDFGKYVNRDEVISQAPSLLVACGLALRIVG.   354
NTDB id 1253 GC085 RS04835 WP 011215094.1     LSHMSFGKKVNLDVINSDAPTLMVACGLALRHVE.   354
consensus                                     ***! !***!** *****! *!*!!!!!!!!***
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X non conserved
X similar
X $\geq$ 50% conserved