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logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1       MFKIILLGWIAGIAMMGRTLECIEPFTCITGILTFLIWAIYFYKKRKHQIGSGLRALVILISASFLFGIGAQYADSALEE    80
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1   MLKIILLGWIAGIAIMGFHLDFIELTWWIWLAIAILIFSFGIYKKEFYFYRPFYKALILISASWGLFTTGYHFADSALEQ    80
NTDB id 1070 ABD1 RS13145 WP 000472713.1        MFKIILLGWIGGIALMGIDFPLIMQYEKVGEALLLLAFIFYLYKRPMFVDRPFLKAVFCLLCTTSLFMVGYHYAEKALIE    80
NTDB id 1083 HON27 RS04190 WP 168726476.1       MFKIILLGWIGGIALMGIDFPLIMQYEKVGEALLLLAFIFYLYKRPMFVDRPFLKAVFCLLCTTSLFMVGYHYAEKALIE    80
consensus                                       !*!!!!!!!!*!!!*!!** **! ***********! *****!***********!*********!!**!***!* !! *

logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1       RLYYKIMHSDTHEVIIYVHQMGESSDKGIKQVAEVLNIHPDVVNWVFYIDQSL.LGAEKSSSLELGQYYRITAKTKPIHS   159
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1   RLQLKEFESQSYEAIVYIKNIDELSDDGHKQIAEVLNRHKQSVNWLLYLKQDK.YLLNQKQSLELGHYYRVYGQIKPVHS   159
NTDB id 1070 ABD1 RS13145 WP 000472713.1        RLEQRETDTRNLDIIVYINRLSEEKDNKVQQTAQVLNLSKEPVNWLLYLKSNNQNLLKNNQNLELGHYYRISGKTRPAHS   160
NTDB id 1083 HON27 RS04190 WP 168726476.1       RLEQRETDTRNLDIIVYINRLSEEKDNKVQQTAQVLNLSKEPVNWLLYLKSNNQNLLKNNQNLELGHYYRISGKTRPAHS   160
consensus                                       !!*************!*!*****!* !* * !*! !!!*****!!!**!** ********** !!!!*!!!******!*!!

logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1       YALEGAFDVEKWFVQQNWQATAQVYRIQTLSKQEVMSAGFYLHVKQQQTWRAQFLQSVEKLRLKFRNYFIDSSLSHHALL   239
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1   YAVAGAFDQEKWFLQQNIMSAFSVQSIQPLSFDEIYHLGYQGHLKQQQSFSAKFLLSVEMKRLAFRQMLQTSKLQHKGLL   239
NTDB id 1070 ABD1 RS13145 WP 000472713.1        YATPGAFDQEKWFIQRNIMSGFNVRYIEPLSLDEIYRLGYQQHLKEQQSFSNSFRLNIEKLRLTFRQILNSSSLQQKGLI   240
NTDB id 1083 HON27 RS04190 WP 168726476.1       YATPGSFDQEKWFIQRNIMSGFNVRYIEPLSLDEIYRLGYQQHLKEQQSFSSSFRLNIEKLRLTFRQILNSSSLQQKGLI   240
consensus                                       !!**!*!!*!!!!*! !******!**! *!!**!****!***!*! !!******! * *!**!!*!!*****!*!****!*

logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1       LALLTGDRSLLNPELEQQFQRFGISHLLAISGPHVLIFAAMVSWIFTRLIHRFKPEIYLSIPRQKLVLLPFLMCVILYCA   319
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1   LALLTGDESLLSNTLKQQFQQLGISHLLAISGPHVLIFAFMLSWLIQRLIQHYCPHLYLWQPRQILLLLPFLLSVLVYVA   319
NTDB id 1070 ABD1 RS13145 WP 000472713.1        LALLTGDESLLSDETQLQFKQLGISHLLAISGPHVLIFAIMLSWAYHQFISRYYPQIYLWKPKQVLIAVPCCLGVLIYTA   320
NTDB id 1083 HON27 RS04190 WP 168726476.1       LALLTGDESLLSDETQLQFKQLGISHLLAISGPHVFIFAIMLSWACHQFISRYYPKIYLWKPKQILMAVPCCLGVLIYTA   320
consensus                                       !!!!!!!*!!!*** * !! **!!!!!!!!!!!!!*!!!*!*!!*** !****! *!!**!*!*!* *!****!**!*!

logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1       FVGFEIPALRTLFTVMLGSILLLIRHHLSAFSLLVLSASLLLWMDPFSILSAAFWLSYGACFILIRIYQDMTQAQQFHQT   399
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1   FVGFEIPAIRTLLTALLASIFLFFRQSIRPFSLLIYSASILLLFDPFSILSAGFWLSYGACFILLRVYQTIAQLPQDQPI   399
NTDB id 1070 ABD1 RS13145 WP 000472713.1        FVGFEIPALRTFLSVLIFIGFLLLKQPIKPFTLLVYSASLLLLMDPFSVLSAGFWLSYGACFILLRIYQTIAQLPEQHFL   400
NTDB id 1083 HON27 RS04190 WP 168726476.1       FVGFEIPALRTLLSTLIFIGFLLLKQPIKPFTLLVYSASLLLLMDPFSVLSAGFWLSYGACFILIRIYQTIAQLPEQHFL   400
consensus                                       !!!!!!!!*!!******* *!********!*!!**!!!*!!**!!!!*!!!*!!!!!!!!!!!!*!*!!***!** ****

logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1       RFTRGM.QILKALVRSQWKIFVALMPLTLIIFKQISWIAPLSNLIAIPILGGIIVPLDIMAGCLWLFIPSLGNLLFYLNN   478
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1   TWTQKIFLAAKALVESQSKIFIALLPLVLIFFQQVSWIAPVSNLIAIPLLGGVVVPLDILAACIWLIIPALGQFLFQINND   479
NTDB id 1070 ABD1 RS13145 WP 000472713.1        SLSSKMIFMSKVLIESQGKIFIALSPLTLLFFQQISWVAPLTNIIAVPIVGSVIVPLNIMAACAWFVIKPFGNMLFHFND   480
NTDB id 1083 HON27 RS04190 WP 168726476.1       SLSSKMIFMSKVLIESQGKIFIALSPLTLLFFQRISWVAPLTNIIAVPIVGSVIVPLNIIAACAWFVVKPFGNMLFHFND   480
consensus                                       **********! !**!!*!!!*!!*!!*!**!***!!*!!**!*!!*!**! **!!! !*!*!*! **** !**!!**!*
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logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1      SLLELFIGILKGLDHLFNPQLQYYAFTPLLLICLTLAIVILFLPKGVVPKFWSVLCFVPLLIPSKMIRPTQISILDVGQG   558
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1 FVLSILMSILDILQ.NIAPHLYGVSYTPLMMVSLILGMLILFMPRGTIPKAWALICFLPICIGVKT.QPTVLNILDVGQG   557
NTDB id 1070 ABD1 RS13145 WP 000472713.1       MLLSILLSCLGLLE.KLSLTLQGISLTPLSLLAISCAIIILFLPKGILPKTWGILCCLPLVIMNKTSQQIQLNILDVGQG   559
NTDB id 1083 HON27 RS04190 WP 168726476.1      MLLSILLSCLGLLE.KLSLPLQGISLTPLSLLAISCAIIILFLPKGILPKTWGILCCLPLVMMNKTSQQIQLNILDVGQG   559
consensus                                      **!******!**!* **** !*****!!!**********!!!*!*!**!!*!***!**!*****!**** ***!!!!!!!
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logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1      QSIFIQDQTQRILLDTGGYYDESKFSIGKQVVVPFLKSQGVHQLDRVILTHLDQDHSGGLPAVLQLIPTRMIQANEEI..   636
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1 QAVFLQHPEQIFMIDTGGSYNENKFSLGERVVIPFIRQQGVRHIDHVILSHLDQDHSGAFPIIQKNFEISQVQSNEYDQK   637
NTDB id 1070 ABD1 RS13145 WP 000472713.1       QAIFLQHPEQNWLIDTGGSYDEKIFSIGQNVVVPFLRQQGVRQLDHVVLSHLDQDHSGAFPLIQQEIPVKQLISNEQL.P   638
NTDB id 1083 HON27 RS04190 WP 168726476.1      QAIFLQHPQQNWLIDTGGSYDEKIFSIGQNVVVPFLRQQGVRQLDHVVLSHLDQDHSGAFPIIQQEIPVKQLISNEQL.P   638
consensus                                      !**!*!***!****!!!!*!*!* !!*!**!!*!!***!!!***!*!*!*!!!!!!!!**!*********** *!!** *
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logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1      KDLSIPFSLCHSGQTWETSELKIQILSPKLENLQSTVFNRNEASCVMYLQLKQASDLKYFLMMGDAGWETEYQLLQQYPD   716
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1 MQFKNNFSFCQQGQHWSYENLKVEILSPKADELIHAKDQQNEYSCVVYLQFLNAQPYQNFLIMGDAGWEAEYKLMQQYPD   717
NTDB id 1070 ABD1 RS13145 WP 000472713.1       NDLKQPFQYCHQGQQWHYPELDIQILWPKEKDLAFVTSNQNQYSCVVYLQFKKVGGYQNFLIMGDAGWEAEYELLKDYPN   718
NTDB id 1083 HON27 RS04190 WP 168726476.1      NDLKQPFQYCHQGQQWHYSELDIQILWPKEKDLAFVASNQNQYSCVVYLQFKKVGGYQNFLIMGDAGWEAEYELLKDYPN   718
consensus                                      ******! *!**!!*!****! **!! !!****!*** ***! *!!!*!!!*** *****!!*!!!!!!!*!!*!*  !!
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logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1      LKVDVLILGHHGSQHSSSYDFLKRYHPKIAIASAGFENGYGHPSAEVSNRLQDLNIPLLSTTDKCAIHFKFDSGNEQTEF   796
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1 LKVDVLILGHHGSKHSSAYDFLATLKPKLAIASAGFDNRYGHPSQELKNRLQALHIPLLTTVEAGTIRFIFENGQVELRQ   797
NTDB id 1070 ABD1 RS13145 WP 000472713.1       LKIDVLVLGHHGSKHSSAYDFLATLKPKLAIASAGFDNRYGHPSQQVIARLKALHIPLKSTVEQGTLSFVLENHKIVLHD   798
NTDB id 1083 HON27 RS04190 WP 168726476.1      LKIDVLVLGHHGSKHSSAYDFLATLKPKLAIASAGFDNRYGHPSQQVIARLKALHIPLKSTLEQGTLSFVLENHKLVLHD   798
consensus                                      !!*!!!*!!!!!!!*!!!*!!!!****!!*!!!!!!!*!*!!!!!* ** !! *!*!!! *!***!***!* ************
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logo
NTDB id 1015 ACIAD RS12005 WP 004928865.1      YRQSYKWLQR..   806
NTDB id 1006845 ABLB98 RS09390 WP 348882113.1  QRQQIKWLERK.   808
NTDB id 1070 ABD1 RS13145 WP 000472713.1       RRLDRLWLSRGF   810
NTDB id 1083 HON27 RS04190 WP 168726476.1      RRLDRLWLSRGF   810
consensus                                      *! ** !!*!**
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X non conserved

X similar

X $\geq$ 50% conserved