```
logo
NTDB id 1367 NTHI RS01945 WP 011271971.1      .MTKKLFYYQGSNALNQKQKGSIIADTKQQAHFQLISRGLTHIKLQQNWQFGAKPKNSEISELLNQLATLLQSVIPLKNS   79
NTDB id 1356 HI 0297 AAC21961.1               .MTKKLFYYQASNPLNQKQKGSIIADTKQQAHFQLISRGLTHIKLQQNWQFGAKPKNSEISELLNQLATLLQSAIPLKNS   79
NTDB id 244631 CJD39 RS01715 WP 111750325.1   MNNIYLFQWKGRNRLGQKQTGQRIATSRAVLEQQLQQKGYRQLRISRQFVLSTAPKKEEINRYIQQLAMLVQASIPLKQC   80
NTDB id 1385 A4U84 RS08520 WP 005711777.1     MNNIYEFQWKGRNRFGQKQTGRQLAESREVLEKRLQQKGYSQLSISRHFALPTAPKKEEINQFMQQLALLIQAKIPLKQS   80
consensus                                     *    *! * * !** !!! !***!****  ***!  *! *****  *** *  !!  !! ****!!!*!*!  !!!!**

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NTDB id 1367 NTHI RS01945 WP 011271971.1      LQILQQNCTQIMLNKWLERLLQSIESGLAFSQAIEQQGKYLTQQEIQLIQVGEMTGKLAVVCKKIATHRSQSLALQRKLQ   159
NTDB id 1356 HI 0297 AAC21961.1               LQILQQNCTQIVLNEWLERLLQSIESGLAFSQAIEQQGKYLTQQEIQLIQVGEMTGKLAVVCKKIATHRSQSLALQRKLQ   159
NTDB id 244631 CJD39 RS01715 WP 111750325.1   LSMLLESCQNLSFYRWQQETIRLLEAGFAFSSALEKQGKYLNAQEIQLIKMAETSGNLGIILSNIAQTREKSEKLTKKVK   160
NTDB id 1385 A4U84 RS08520 WP 005711777.1     LVMLLETCQNRTFYRWQQETIRLIEAGFALSVAFKKQGKYINPQEIQLIKMAETSGNLGLILTNIAQRREKSEKLAKKIK   160
consensus                                     !**!  *! **   *!    *  *! ! !*!*!** !!!!* *!!!!!! **! *! !***** !! *!  !  !**!*

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NTDB id 1367 NTHI RS01945 WP 011271971.1      KIMLYPSMVLGISLLLTLALLLFIAPQFAEMYSGNNAELPTITAILLSISNFLKQNIGILLFFVLSFFLFYYFYLKRQTW   239
NTDB id 1356 HI 0297 AAC21961.1               KIMLYPSMVLGISLLLTLALLLFIVPQFAEMYSGNNAELPTITAILLSISNFLKQNIGILLFFVLSFFLFYYFYLKRQTW   239
NTDB id 244631 CJD39 RS01715 WP 111750325.1   KILFYPLFILTISITLSILLLIFIVPQFAELYGSKGKSLPLITNVLFILSNFLQQNIIHLILFIIISLLIISFLNKKTTI   240
NTDB id 1385 A4U84 RS08520 WP 005711777.1     KILFYPVFILAISITLSILLLLFIVPQFAELYGSKGKSLPLITEILFSLSHFLQHSILTLIILCVLCFLLIHILNKKTDL   240
consensus                                     !!* !!* *!*!!* !** !!*!!*!!!!!*!       !! !!**! **!*!! **!**!********!* **  !* **

logo
NTDB id 1367 NTHI RS01945 WP 011271971.1      FHQKKNQLISITPIFGTIQKLSRLVNFSQSLQIMLQAGVPLNQALDSFLPRTQTWQTKKTLVNDMVLDKEVRSILQWVSQ   319
NTDB id 1356 HI 0297 AAC21961.1               FYQKKNQLISITPIFGTIQKLSRLVNFSQSLQIMLQAGVPLNQALDSFLPRTQTWQTKKTLVNDIVLDKEVRSILQWVSQ   319
NTDB id 244631 CJD39 RS01715 WP 111750325.1   IKKVKLLILSQLPIFKGIIQDSRIIFFCQNSSLMLASHIRLDSVLHSFLANK.........SDDILLQHSLASTLTFLKQ   311
NTDB id 1385 A4U84 RS08520 WP 005711777.1     IKRLKFIILSHIPILKGIIQYSRIIFFCQNSSLMLASHIRLDTVLHSFLANK.........SDDILLQGSLATTLTYLKQ   311
consensus                                      ***!****!**!!*  !  *!!** !*!   *!!  ***! * !*!!!*  *********  !**! * * * ! ** !

logo
NTDB id 1367 NTHI RS01945 WP 011271971.1      GYAFSNSVSSDLFPMEAQQMLQIGEQSGKLALMLEHIADNYQEKLNHQIDLLSQMLEPLMMVIIGSLIGIIMMGMYLPIF   399
NTDB id 1356 HI 0297 AAC21961.1               GYAFSNSVSSDLFPMEAQQMLQIGEQSGKLALMLEHIAENYQEKLNHQIDLLSQMLEPLMMVIIGSLIGIIMMGMYLPIF   399
NTDB id 244631 CJD39 RS01715 WP 111750325.1   GYRLADSLDPKLFSNDVLQMIAIGEQSGKLSVMLKQISDIYQQRLDYQIDLLSQLLEPMLMLLMGIIVGTILVGLYLPIF   391
NTDB id 1385 A4U84 RS08520 WP 005711777.1     GYRLADSLDPKLFPTDMLQMIAIGEQSGKLSPMLQQISDGYQQRLDYQIDILSQLLEPMLMVLMGIIVGTILVGLYLPIF   391
consensus                                     !!    !* * !!**** !!* !!!!!!!! *!!**! **!! *! *!!!*!!!*!!!**!***! **! !**!*!!!!!

logo
NTDB id 1367 NTHI RS01945 WP 011271971.1      NMGSVIQ   406
NTDB id 1356 HI 0297 AAC21961.1               NMGSVIQ   406
NTDB id 244631 CJD39 RS01715 WP 111750325.1   DMGAMIE   398
NTDB id 1385 A4U84 RS08520 WP 005711777.1     DMGAMIE   398
consensus                                      !! *!
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X non conserved
X similar
X $\geq$ 50% conserved