```
logo

NTDB id 1145785 CS888 RS00230 WP 089086476.1  MRKVLYALIGFLLAFSVLKADDFLEEANETAPAHLNHPMQDLNAIQGSFFDKNRSKMSNTLNINYFQGQTYKIRLRYAMA  80
NTDB id 1217 CAA10656.1 1440..2420( )         MRKVLYALMGFLLAFSALKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNIDYFQGQTYKIRLRYAMA  80
consensus                                     !!!!!!!!*!!!!!!! !!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!

logo

NTDB id 1145785 CS888 RS00230 WP 089086476.1  TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
NTDB id 1217 CAA10656.1 1440..2420( )         TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
consensus                                     !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1145785 CS888 RS00230 WP 089086476.1  YYTNAFIKPQKENQENMAENAPKDAQKNNKPLKEEKEETKTPEEEVIIIGDNTNAMKIIKKDIQKGYKALKSSQRKWYCL  240
NTDB id 1217 CAA10656.1 1440..2420( )         YYSNAFLKPQN..KENVLENALENTPTNNKPLKEKKEETKEKEEETIITIGDNTNAMKIVKKDIQKGYKALKSSQRKWYCL  238
consensus                                     !!*!!!*!!! ** !!* !!!*   * !!!!!!! !!!!! *!!! ! !!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1145785 CS888 RS00230 WP 089086476.1  WACSKKSKLSLMPEEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSAKWTLRLGKDYLCIR  320
NTDB id 1217 CAA10656.1 1440..2420( )         GICSKKSKLSLMPKEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSTKWTLRLGKDYLCIR  318
consensus                                       !!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

   <0

logo

NTDB id 1145785 CS888 RS00230 WP 089086476.1  FVKRSKGE.   328
NTDB id 1217 CAA10656.1 1440..2420( )         FVKKGKDE*   326
consensus                                     !!!* ! !
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X non conserved
X similar
X ≥ 50% conserved