```
logo

NTDB id 1145886 CS886 RS00230 WP 089086476.1  MRKVLYALIGFLLAFSVLKADDFLEEANETAPAHLNHPMQDLNAIQGSFFDKNRSKMSNTLNINYFQGQTYKIRLRYAMA   80
NTDB id 1217 CAA10656.1 1440..2420( )         MRKVLYALMGFLLAFSALKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNIDYFQGQTYKIRLRYAMA   80
consensus                                     !!!!!!!!*!!!!!!! !!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!

logo

NTDB id 1145886 CS886 RS00230 WP 089086476.1  TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
NTDB id 1217 CAA10656.1 1440..2420( )         TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
consensus                                     !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1145886 CS886 RS00230 WP 089086476.1  YYTNAFIKPQKENQENMAENAPKDAQKNNKPLKEEKEETKTPEEEVIIIGDNTNAMKIIKKDIQKGYKALKSSQRKWYCL  240
NTDB id 1217 CAA10656.1 1440..2420( )         YYSNAFLKPQN..KENVLENALENTPTNNKPLKEKKEETKEKEEETITIGDNTNAMKIVKKDIQKGYKALKSSQRKWYCL  238
consensus                                     !!*!!!*!!! ** !!* !!!*   * !!!!!!! !!!!! *!!! ! !!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1145886 CS886 RS00230 WP 089086476.1  WACSKKSKLSLMPEEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSAKWTLRLGKDYLCIR  320
NTDB id 1217 CAA10656.1 1440..2420( )         GICSKKSKLSLMPKEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSTKWTLRLGKDYLCIR  318
consensus                                       !!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

   <0

logo

NTDB id 1145886 CS886 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