```
logo

NTDB id 667236 MPG79 RS00730 WP 245107551.1  MRKVLCALVGFLLAFSVLKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNVDYFQGQTYKIRLRYAMA   80
NTDB id 1217 CAA10656.1 1440..2420( )        MRKVLYALMGFLLAFSALKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNIDYFQGQTYKIRLRYAMA   80
consensus                                    !!!!!*!!*!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!

logo

NTDB id 667236 MPG79 RS00730 WP 245107551.1  TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDNEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
NTDB id 1217 CAA10656.1 1440..2420( )        TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 667236 MPG79 RS00730 WP 245107551.1  YYSNAFMKPQNKENTLEKAPT...NNNKPLKEEKEET..KEEETITIGDNTNAMKIVKKDIQKGYRALKSSQRKWYCLGI  235
NTDB id 1217 CAA10656.1 1440..2420( )        YYSNAFLKPQNKENVLENALENTPTNNKPLKEKKEETKEKEEETITIGDNTNAMKIVKKDIQKGYKALKSSQRKWYCLGI  240
consensus                                    !!!!!!*!!!!!!!! !! !* *** !!!!!!! !!!!**!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!

logo

NTDB id 667236 MPG79 RS00730 WP 245107551.1  CSKKSKLSLMPEEIFNDKQFTYFKFDKKLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSAKWTLRLGKDYLCIRFI  315
NTDB id 1217 CAA10656.1 1440..2420( )        CSKKSKLSLMPKEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSTKWTLRLGKDYLCIRFV  320
consensus                                    !!!!!!!!!!!! !!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*

   <0

logo

NTDB id 667236 MPG79 RS00730 WP 245107551.1  KKGKDE.   321
NTDB id 1217 CAA10656.1 1440..2420( )        KKGKDE*   326
consensus                                    !!!!!!
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X non conserved
X similar
X ≥ 50% conserved