```
logo
NTDB id 667428 MPG13 RS00715 WP 180602210.1  MRKVLYALVGFLLVFSALKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNIDYFQGQTYKIRLRYAMA   80
NTDB id 1217 CAA10656.1 1440..2420( )        MRKVLYALMGFLLAFSALKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNIDYFQGQTYKIRLRYAMA   80
consensus                                    !!!!!!!!*!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 667428 MPG13 RS00715 WP 180602210.1  TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDNEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
NTDB id 1217 CAA10656.1 1440..2420( )        TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 667428 MPG13 RS00715 WP 180602210.1  YYSNAFMKPQNKENTL....EKAPTNNKPLKEEKEETK..EEEAITIGDNTNAMKIVKKDIQKGYRALKSSQRKWYCLWI  234
NTDB id 1217 CAA10656.1 1440..2420( )        YYSNAFLKPQNKENVLENALENTPTNNKPLKEKKEETKEKEEETITIGDNTNAMKIVKKDIQKGYKALKSSQRKWYCLGI  240
consensus                                    !!!!!!*!!!!!!!! !****!  !!!!!!!!! !!!!!**!!! !!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!! !

logo
NTDB id 667428 MPG13 RS00715 WP 180602210.1  CSKKSKLSLMPKEIFNDKQFTYFKFDKKLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSAKWTLRLGKDYLCIRFI  314
NTDB id 1217 CAA10656.1 1440..2420( )        CSKKSKLSLMPKEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSTKWTLRLGKDYLCIRFV  320
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*

   <0
logo
NTDB id 667428 MPG13 RS00715 WP 180602210.1  KKGKDE.   320
NTDB id 1217 CAA10656.1 1440..2420( )        KKGKDE*   326
consensus                                    !!!!!!
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X non conserved
X similar
X ≥ 50% conserved