```
logo
NTDB id 667729 MPG67 RS00725 WP 245072922.1  MRKVLCTLVGFLLVSSALKADDFLEEANETAPAHLNHPMQDLNAIQGSFFDKNRSKMSNTLNVDYFQGQTYKIRLRYAMA   80
NTDB id 1217 CAA10656.1 1440..2420( )        MRKVLYALMGFLLAFSALKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNIDYFQGQTYKIRLRYAMA   80
consensus                                    !!!!!* !*!!!!  !!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!

logo
NTDB id 667729 MPG67 RS00725 WP 245072922.1  TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDNEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
NTDB id 1217 CAA10656.1 1440..2420( )        TLLFFSKPISDFVLGDKVGFDAKILESNDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKN  160
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 667729 MPG67 RS00725 WP 245072922.1  YYSNAFMKPQNKENTLEKAPTN...NNKSLKEEKEE..TKEEETITIGDNTNAMKIVKKDIQKGYRALKSSQRKWYCLGI  235
NTDB id 1217 CAA10656.1 1440..2420( )        YYSNAFLKPQNKENVLENALENTPTNNKPLKEKKEETKEKEEETITIGDNTNAMKIVKKDIQKGYKALKSSQRKWYCLGI  240
consensus                                    !!!!!!*!!!!!!!! !! !* !***!!!*!!! !!!** !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 667729 MPG67 RS00725 WP 245072922.1  CSKKSKLSLMPEEIFNDKQFTYFKFDKKLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSAKWTLRLGKDYLCIRFV  315
NTDB id 1217 CAA10656.1 1440..2420( )        CSKKSKLSLMPKEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSTKWTLRLGKDYLCIRFV  320
consensus                                    !!!!!!!!!!!! !!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!
```