

logo

NTDB id 994256 V6C35 RS04555 WP 000959282.1
NTDB id 26 SA RS04860 WP 000959282.1
NTDB id 42 MW RS04740 WP 000959293.1
consensus

MLVALNEEKERV LATTALRKTQYFCPVC GKQVILKRGLKVISHF AHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV 80
MLVALNEEKERV LATTALRKTQYFCPVC GKQVILKRGLKVISHF AHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV 80
MLVALNEEKERV LATTALRKTQYFCPVC GKQVILKRGLKVISHF AHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV 80

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[illegible]

logo

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NTDB id 26 SA RS04860 WP 000959282.1
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consensus
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HTFNLKKQIMMFQQIQYLGGHKYVAEKRNAKI^SELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ 240
HTFNLKKQIMMFQQIQYLGGHKYVAEKRNAKI^SELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ 240
HTFNLKKQIMMFQQIQYLGGHKYVAEKRNAKI^SELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ 240

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LTDQEVVNYGYIFPEQIYIENHPIEWQLQVDLWLKNGKSKLVNDNLNYFKLKKFIVALESKTAIEKLINNYLNICSDR 320
 LTDQEVVNYGYIFPEQIYIENHPIEWQLQVDLWLKNGKSKLVNDNLNYFKLKKFIVALESKTAIEKLINNYLNICSDR 320
 LTDQEVVHNYGYIFPEQIYIKNHPIEWQLQVDLWLKNGKSKLVNDNLNYFKLKKFIVGLESKTAIEKLINNYLNICSDR 320

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The diagram illustrates a protein structure with a sequence logo and a sequence alignment. The sequence logo at the top shows the amino acid sequence GNDVQILF, with each letter color-coded: G (grey), N (blue), D (red), V (green), Q (light blue), I (dark green), L (dark blue), and F (blue). Below the logo, a blue box contains the sequence GNDVQILF repeated three times. To the right of the box, the number 328 is displayed three times, indicating the position of the sequence in the protein. At the bottom, a row of red exclamation marks (!) is shown, likely representing a specific feature or mutation.

X	non conserved
X	similar
X	≥ 50% conserved