```
logo

NTDB id 1080788 ACK2WG RS12280 WP 014114401.1   MKQIRKTWPLKDQAQLLKRLGEMTEGGYTLLDGIRLMELQMNKRQAADLTDAITRLREGAPFYQVLKGLSFHKEAVGICY   80
NTDB id 97 BSU 24720 NP 390352.1                ......................MTAGGYTLLDGLRLMELQMNKRQAADLTDSVTCLREGAPFYQVLKSLSFHKEAVGICY   58
consensus                                       ************************!! !!!!!!!!*!!!!!!!!!!!!!!!!! *!*!!!!!!!!!!!! !!!!!!!!!!!!

logo

NTDB id 1080788 ACK2WG RS12280 WP 014114401.1   FAETHGELPASMIQSGELLERKIAQADQLKRVLRYPLFLIFTVAVMFYMLQAIIIPQFSGIYQSMNMETTRSTDMLFAFF   160
NTDB id 97 BSU 24720 NP 390352.1                FAETHGELPASMIQSGELLERKIAQADQLKRVLRYPLFLIFTVAVMFYMLQSIIIPQFSGIYQSMNMETSRSTDMLFAFF   138
consensus                                       !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!*!!!!!!!!!!

logo

NTDB id 1080788 ACK2WG RS12280 WP 014114401.1   QHIDLVIILLALFAGGIGLYYWLVFKKKSPARQMLICVRIPLVGNLVRLFNSYFFSLQLSSLLQSGLSIYDSLNAFKHQT   240
NTDB id 97 BSU 24720 NP 390352.1                QHIDLVIILLVLFTAGIGIYYWLVFKKKSPARQMLICIRIPLVGKLVKLFNSYFFSLQLSSLLKSGLSIYDSLNAFKHQT   218
consensus                                       !!!!!!!!!! !! *!!!*!!!!!!!!!!!!!!!!!!*!!!!!! !!*!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!

logo

NTDB id 1080788 ACK2WG RS12280 WP 014114401.1   FLPFYRCEAEQVIERLKAGESIETAICESPFYETDFSKVISHGQLSGRLDRELFTYSQFILQRLEHKAQKWTGILQPMIY   320
NTDB id 97 BSU 24720 NP 390352.1                FLPFYRCEAEQLIERLKAGESIESAICGSLFYETDLSKVISHGQLSGRLDRELFTYSQFILQRLEHKAQKWTGILQPMIY   298
consensus                                       !!!!!!!!!!!*!!!!!!!!!!!*!!! !*!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1080788 ACK2WG RS12280 WP 014114401.1   GFVAAMILIVYLSMLLPMYQMMNQM   345
NTDB id 97 BSU 24720 NP 390352.1                GFVAAMILLVYLSMLVPMYQMMNQM   323
consensus                                       !!!!!!!!*!!!!!!*!!!!!!!!!
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X non conserved

X similar

X ≥50% conserved