```
logo
NTDB id 1039427 ACC040 RS12500 WP 372401671.1  MKQIRKTWPLKDQAQLLKRLGEMTEGGYTLLDGIRLMELQMNKRQAADLTDAITHLREGAPFYQVLKGLSFHKEAVGICY   80
NTDB id 97 BSU 24720 NP 390352.1               ......................MTAGGYTLLDGLRLMELQMNKRQAADLTDSVTCLREGAPFYQVLKSLSFHKEAVGICY   58
consensus                                      **********************!! !!!!!!!!*!!!!!!!!!!!!!!!!! *! !!!!!!!!!!!! !!!!!!!!!!!!

logo
NTDB id 1039427 ACC040 RS12500 WP 372401671.1  FAETHGELPDSMIQSGELLERKIAQADQLKRVLRYPLFLIFTVAVMFYMLQAIIIPQFSGIYQSMNMETTRSTDMLFAFF  160
NTDB id 97 BSU 24720 NP 390352.1               FAETHGELPASMIQSGELLERKIAQADQLKRVLRYPLFLIFTVAVMFYMLQSIIIPQFSGIYQSMNMETSRSTDMLFAFF  138
consensus                                      !!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!*!!!!!!!!!!

logo
NTDB id 1039427 ACC040 RS12500 WP 372401671.1  QHIDLVIILLALFTGGIGLYYWLVFKKKSPARQMLICVRIPLVGNLVRLFNSYFFSLQLSSLLQSGLSIYDSLNAFKHQT  240
NTDB id 97 BSU 24720 NP 390352.1               QHIDLVIILLVLFTAGIGIYYWLVFKKKSPARQMLICIRIPLVGKLVKLFNSYFFSLQLSSLLKSGLSIYDSLNAFKHQT  218
consensus                                      !!!!!!!!!! !!!*!!!*!!!!!!!!!!!!!!!!!!*!!!!!! !!*!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!

logo
NTDB id 1039427 ACC040 RS12500 WP 372401671.1  FLPFYRCEAEQVIERLKAGESIETAICESPFYETDFSKVISHGQLSGRLDRELFTYSQFMLQRLEHKAQKWTGILQPMIY  320
NTDB id 97 BSU 24720 NP 390352.1               FLPFYRCEAEQLIERLKAGESIESAICGSLFYETDLSKVISHGQLSGRLDRELFTYSQFILQRLEHKAQKWTGILQPMIY  298
consensus                                      !!!!!!!!!!!*!!!!!!!!!!!*!!! !*!!!!! !!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1039427 ACC040 RS12500 WP 372401671.1  GFVAAMILIVYLSMLLPMYQMMNQM  345
NTDB id 97 BSU 24720 NP 390352.1               GFVAAMILLVYLSMLVPMYQMMNQM  323
consensus                                      !!!!!!!!*!!!!!!*!!!!!!!!!
```

X non conserved
X similar
X ≥50% conserved