```
logo
NTDB id 106292 AABA14 RS09165 WP 000760673.1  MKKYQQLFKQIQETIQNETYAVGDFLPSEHDLMEQYQVSRDTVRKALSLLQEEGLIKKIRGQGSQVVKEETVNFPVSNLT   80
NTDB id 440 SMU RS09330 WP 002273717.1        MKKYEIIFKKLEEDILKGHYQMGDYLPPEMELSQTYASSRDTVRKALQLLTKAGFIKTVQGRGSQIIKRERINFPVSQLT   80
consensus                                     !!!!  *!! * ! !   *! *!!*!!*!**!   !  !!!!!!!!! !!   ! !! * ! !!!**! ! *!!!!!*!!

logo
NTDB id 106292 AABA14 RS09165 WP 000760673.1  SYQELVKELGLRSKTNVVSLDKIIIDKKSSLITGFPEFRMVWKVVRQRVVDDLVSVLDTDYLDMELIPNLTRQIAEQSIY  160
NTDB id 440 SMU RS09330 WP 002273717.1        SYQELVKQLQMNVKTNVIAIDKLIVDEKLTKLTGFENKGLVWRITRQRVIDGVASILDTDYLDKALIPHMTREIAEHSIY  160
consensus                                     !!!!!!! ! *  !!!!* *!!*!*! ! * *!!!*   *!!** !!!!*! * !*!!!!!!!  !!!**!! !!!*!!!

logo
NTDB id 106292 AABA14 RS09165 WP 000760673.1  SYIENGLKLLIDYAQKEITIDHSSDRDKILMDIGKDPYVVSIKSKVYLQDGRQFQFTESRHKLEKFRFVDFAKRKK.   236
NTDB id 440 SMU RS09330 WP 002273717.1        DYLENQLKLDIAYAQKIITIDQVSQKDKILLDLDSENHVVSVKSKVYLSNQQQFQFTESRHKLEKFRFVDFARRHRD   237
consensus                                      !*!! !!! ! !!!! !!!!* ! *!!!!*!*  ***!!!*!!!!!!    !!!!!!!!!!!!!!!!!!!!*!***
```

X non conserved

X similar

X $\geq$ 50% conserved