```
logo

NTDB id 1044759 ACCB35 RS06160 WP 374056220.1  MGKISKITKQVKNDERYNLFIDGKYAFSVDEAVLAKFQLRKGLEIDELEISELQYEDDVKKAFNKAIQYLAYRMRTEKEI   80
NTDB id 116 BSU 08520 NP 388733.1              MPFITKISTQKKNTERFNIFLDDKYAFSVDADVLVKFELKKGKELDDLDIIEIQYGDEVKKGFNRALDFLSYRMRSTKEV   80
consensus                                      !* !*!!* ! !! !!*!*!*! !!!!!!!  !! !! !*!! !*!*!*! !*!! !*!!!*!!*!* *! !!!!* !!*

logo

NTDB id 1044759 ACCB35 RS06160 WP 374056220.1  RDHLQEGDPEDGIIQEVIHKLNDIKYINDLEFAHAFVGTHMNTSDKGPTWIRGELRRKGVDDPYIEEALAVFSVEKQVDK  160
NTDB id 116 BSU 08520 NP 388733.1              EDHLKKKETSPPVIAEVIHRLNDYKYLNDQEFAAAYVSTHKKTNGKGPDVLFRELRAKGIDDDTIKEALSSFSFEDQTRE  160
consensus                                       !!!   ** ***! !!!!*!!! !!*!! !!!*!*! !!  !  !!!  *  !!! !!*!!* ! !!!  !! ! !

logo

NTDB id 1044759 ACCB35 RS06160 WP 374056220.1  AVGLTEKLLQKYRKDSKMIARQKIEQNLMRKGYPGSVMKIVWETVESDRDDDEKWDAVYHQGLKAHRKLSSKYDGYEYVQ  240
NTDB id 116 BSU 08520 NP 388733.1              AVKHVEKLLKKDKKLSTKELKQRAQLQLQRKGFSFDVISAAMDQIEYENDEDTEKEALRLHAEKAFRKYR.YDGSYESAM  239
consensus                                      !! * !!!! ! *! !    *!*   *! !!!**  !*    * *! * !*!   *!* *** !!*!!  *    !!

logo

NTDB id 1044759 ACCB35 RS06160 WP 374056220.1  KMKQTLYRKGFSMEDIERLLEELKENEE  268
NTDB id 116 BSU 08520 NP 388733.1              KVKQFLFRKGFSLDLIEQLLQEEEY...  264
consensus                                      !*!! !*!!!!!** !! !! !   ***
```

X non conserved

X similar

X ≥ 50% conserved