```
logo
NTDB id 1092447 ACME8E RS00145 WP 165079631.1  ..MKRKLLSFIAL..VLLAVGAYWVEGDNLFTKITGENPPSSAQVTLQFPSGRYPETAQHIKEAIQAGKSPVCTIDREGA   76
NTDB id 110 BSU 03430 NP 388225.3              MNITTDIIKTILLVIVIIAAAAVGLIKGDFFSADQKTSQTKEYDETMAFPSDRYPETAKHIKDAINEGHSEVCTIDRDGA   80
consensus                                      ***    **   !  !**!**! *!   *      !*       **     !* !!! !!!!!! !!!*!!* !*!*!!!!!!*!!

logo
NTDB id 1092447 ACME8E RS00145 WP 165079631.1  EQNRKHSLAGVPTRKGFDRDEWPMAMCSEGGKGANVKYIAPKDNRGAGSWVSHQLDQYEDGTRVKFVIK   145
NTDB id 110 BSU 03430 NP 388225.3              EERREQSLKDVPSKKGYDRDEWPMAMCKEGGEGASVEYISPADNRGAGSWVGHRLTDYPDGTKVLFTIQ   149
consensus                                      !   ! *!!   !!**!!*!!!!!!!!!! !!! !! ! !! ! !!!!!!!!! ! !   !*!!!*! ! !
```

X non conserved

X similar

X $\geq$ 50% conserved