```
logo

NTDB id 1093157 ACMGE3 RS07880 WP 414053607.1  MLNTYLNRFMQMFSKQFKKKDGFTLIEMLLVLLVISVLIIVIIPNIAKQSDNVQNTGCKAQVKMVQGQIEAFKLQNGKTP   80
NTDB id 32 MW RS07995 WP 000472256.1           ............MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP   68
NTDB id 16 SA RS07765 WP 000472256.1           ............MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP   68
consensus                                                  **!**!!***!!!!!!!!!!!*!!*!*!!!!!!!!!!****!*!!!*!!!!!!**!!!!**!*****!

logo

NTDB id 1093157 ACMGE3 RS07880 WP 414053607.1  ASINELVP.EFLKDTQVKCKNGDPIQIEDGEAVVQKA   116
NTDB id 32 MW RS07995 WP 000472256.1           SSIEDLIADGFIKEAQKTCKSGETITISNGEAVAN..   103
NTDB id 16 SA RS07765 WP 000472256.1           SSIEDLIADGFIKEAQKTCKSGETITISNGEAVAN..   103
consensus                                      *!!**!****!*!**!**!!*!**!*!**!!!!**
```

X non conserved

X similar

X $\geq$ 50% conserved