```
logo
NTDB id 1077 ABD1 RS06150 WP 000203218.1     MTSNFSQL....STDALSPGQLPESVKALLKRAHINRYPKRTTIVDAGTESKSLYLILKGSVSIILREDDEREIVVAYLN   76
NTDB id 1145 A1552VC RS12025 WP 000242749.1  ..............MVLGKPQTDPTLEWFLSHCHIHKYPSKSTLIHAGEKAETLYYIVKGSVAVLIKDEEGKEMILSYLN   66
NTDB id 42223 PARA RS00335 WP 014064026.1    MSEDVELLEE..QKLQDDDQQRDPALDWFLTHCHLHKYPAKSTLIHAGEDANILYFLIKGTVMVSSKDDEGKEMILSYLG   78
NTDB id 1409 HI 0957 AAC22618.1              MSNELIEIDEVVTSSQEEATQRDPVLDWFLTHCHLHKYPAKSTLIHAGEDATTLYYVIKGSVMVSSKDDEGKEMILTYLG   80
consensus                                    ** *** ***  ******  !*****!***!***!!***!***!!*** *!!***!!*!**********!****!!

logo
NTDB id 1077 ABD1 RS06150 WP 000203218.1     PGDFFGEMGLFEPNPQRTAEVRTRDVCEIAEISYDNFHELSKQYPDLSYAVFAQLVRRLKNTTRKMTDLAFIDVSGRIAR   156
NTDB id 1145 A1552VC RS12025 WP 000242749.1  QGDFIGELGLFEEGQERTAWVRAKTPCEVAEISFKKFRQLIQVNPDILMRLSGQMARRLQVTSQKVGDLAFLDVTGRIAQ   146
NTDB id 42223 PARA RS00335 WP 014064026.1    AGQFFGEAGLFDEGSKRSAWVKTKTPCEIAEISYKKYRQLIQINPEILMFLTSQLSKRLQNTSRQVTNLAFLDVAGRIAQ   158
NTDB id 1409 HI 0957 AAC22618.1              AGQFFGEAGLFDEGSKRSAWVKTKTTCEIAEISYKKYRQLIQANPEILMFLTAQLARRLQNTSRQVTNLAFLDVAGRIAQ   160
consensus                                    *! !*!!*!!!*****!*!*!*****!!*!!!!******!****!********!***!!**!** ** !!!*!!*!!!!*

logo
NTDB id 1077 ABD1 RS06150 WP 000203218.1     CLIDLSSQPEAMILPNGRQIRITRQEIGRIVGCSREMVGRVLKTLEDQGMIQTDGKAILIFDTSLEETPVTDEDYDEEE   235
NTDB id 1145 A1552VC RS12025 WP 000242749.1  TLLNLARQPDAMTHPDGMQIKITRQEIGQIVGCSRETVGRILKMLEEQNLISAHGKTIVVYGTR...............   210
NTDB id 42223 PARA RS00335 WP 014064026.1    ALMHLAKQPEAMTHPDGMQIKITRQEIGQMVGCSRETVGRIIKMLEDQNLIHAHGKTIVVYGTR...............   222
NTDB id 1409 HI 0957 AAC22618.1              TLMNLAKQPEAMTHPDGMQIKITRQEIGQMVGCSRETVGRIIKMLEDQNLIHAHGKTIVVYGAR...............   224
consensus                                    *!**!**!!*!!**!*!*!!*!!!!!!!**!!!!!!*!!!**!*!!*!**!***!!*!******
```

X non conserved

X similar

X $\geq$ 50% conserved