```
logo
NTDB id 396060 GFB47 RS01100 WP 153445876.1  MLHDINLLPWREEKREEYKQRFYGMVVLGVLLALALQWFGSLYIDHLKTQQQTRNQQLQTYVKELDKQLVSLKAIEAKHR   80
NTDB id 1405 DSB67 RS13995 WP 010643496.1    MLYHINLLAWRENQREEHRRRFIGLILLGAIAALGVQWGIGKYYQYQQDQQQQRLDYLAHYISQLDTRIEAMKIAEQEHS   80
NTDB id 1172 A1552VC RS12120 WP 000902745.1  MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
consensus                                    !!* *!!!*!!* *!!****!!*!***!!***!***!**** !*  *  *!! !* *!* **  !! **  *!**! *!*

logo
NTDB id 396060 GFB47 RS01100 WP 153445876.1  SILTRLRLVESLQNERNKTTDFMNLMPELVPEGVYIDKMKMNGREVEIAGISDSTARLATMLDSFERSAWLSDVEMHSII  160
NTDB id 1405 DSB67 RS13995 WP 010643496.1    KILERLKTVEGLQNGRNKTTEFMNLMPSVIPEGVYVDKIKMNDYEIEISGISDSTSRLATMLDNMERSAKLLDVDMHSIV  160
NTDB id 1172 A1552VC RS12120 WP 000902745.1  ALLTRLTVVEQLQQKRNKTTEFMNQMPNLIPEGVYVDKIKMNGHQIEITGISDSTARLATMLDNLEKSDKLTDVEMHEIV  160
consensus                                     *!*!!**!! !!* !!!!!*!!!*!! **!!!!!*!!*!!!* **!!*!!!!!!*!!!!!!!**!*!**!*!!*!!*!*

logo
NTDB id 396060 GFB47 RS01100 WP 153445876.1  SGKVLFGKKFQTFKVSFRFESKQPETAQA.KTKGAK  195
NTDB id 1405 DSB67 RS13995 WP 010643496.1    HGKARFGKEFQTFKVSFMFKDASI...KGGKQHG..  191
NTDB id 1172 A1552VC RS12120 WP 000902745.1  SGNKRFGKQFQSFKVSFQILTPASNPQAGGAHNG..  194
consensus                                    *!* *!!! !!*!!!!! * *        ***  !
```

X non conserved

X similar

X $\geq$ 50% conserved