```
logo

NTDB id 1362 HI 1008 AAC22669.1              ..MKTLFTSVVLCGALVVSSSFAEEKATXQTAQSVVTTQAEAQVAPAVVSDKLNINTATASEIQKSLTGIGAKKAEAIVQ   78
NTDB id 364326 FEM41 RS11020 WP 138096018.1  ..MKTLLSPLWLTLALLVNGSVAHAASEEVT......PQSPVPALITASESQISINTASAEELAEALNGIGLKKAQSIVR   72
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   MQIKTKIVTLFLSLCLPTLPLLANAEE..T.........APAAQVEEGIVITVNINTASAEELATLLKGIGLKKAQAIVD   69
NTDB id 1150 A1552VC RS08580 WP 001166105.1  MQIKTKIVTLFLSLCLPTLPLLANAEE..T.........APAAQVEEGIVITVNINTASAEELATLLKGIGLKKAQAIVD   69
NTDB id 1294 VP RS04495 WP 021449594.1       .......MKWILTLCLVVFAPLSLAAD.............TKADKYEGIEITVNINTASAEEIATMLKGIGEKKAQSIVD   60
consensus                                      *** *****!***!**   ** * *            **** * *** ****!!!!*!*!***!*!!!*!!!**!!*

logo

NTDB id 1362 HI 1008 AAC22669.1              YREKHGNFXNAEQLLEVQGIGKATLEKNRDRITF   112
NTDB id 364326 FEM41 RS11020 WP 138096018.1  YREENGQFASIDDLRSVPGVGNALVERNLSRLKL   106
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   YREANGPFTHIDDLTNVKGIGEATVRNNAARILL   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1  YREANGPFTHIDDLTNVKGIGEATVRNNAARILL   103
NTDB id 1294 VP RS04495 WP 021449594.1       YRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL    94
consensus                                    !!* *!*!  ***!**!*!*!*!****!  !***
```

X non conserved

X similar

X $\geq$ 50% conserved