```
logo
NTDB id 54863 OC457 RS06500 WP 080174725.1  MAKLHPLSFVIGSIFA.ANSYAMTPQPDPDNIGGYVIERSEIITAENIKTMDPMYDVWAQALVTASNSIVESILPGLSTNPDNVK   84
NTDB id 1299 VP RS02975 WP 005483021.1      MFALKHLPIALSCVLAAQANASMNIQPDPQNPNGYVIARGDVQAMEQSKTSDPMYAIWSQALETRPNSVVEAIVPGAATNPENVK   85
consensus                                   !  !* !*  * ** !*     ! *!!!! !* !!!! ! **   !* !! !!!! *! !!! ! *!!*!! !*!!  !!!*!!!

logo
NTDB id 54863 OC457 RS06500 WP 080174725.1  RVEHVFPETEWNYLTQMAAPEYTYTRFLRAIGKFPAFCGDYTDDRDADAICKKSIVTAFAHFAQETGGHISIENVSDNPLGLEEW  169
NTDB id 1299 VP RS02975 WP 005483021.1      RVERVFPESEWDFLTQMAAPEYTYTRFLRAIGKFPAFCGEYTDGRDSDAICKKSIVTAFAHFAQETGGHISIDNVSDNPLGLEEW  170
consensus                                   !!!*!!!!*!! *!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!! !! !!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!

logo
NTDB id 54863 OC457 RS06500 WP 080174725.1  QQALVHVREMGWSEGQPGYTTGCGQNDWQNKRWPCAEGQGYFGRGAKQLSYHFNYGAFSEVMFDGDATVLLNNPGLVADSWLNLA  254
NTDB id 1299 VP RS02975 WP 005483021.1      QQALVHVREMGWSEGQEGYTTGCGQNDWQNKRWPCEPGQGYFGRGAKQLSYHFNYGAFSEVMFDGDASVLLKNPGLVADSWLNLA  255
consensus                                   !!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!! *!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!! !!!!!!!!!!!!!

logo
NTDB id 54863 OC457 RS06500 WP 080174725.1  SAIWFFLTPQAPKPAMLHVIDRTWVPSQRETDAGIGYGFGTTINVINGGIECGEQNKDKGQPVNRIRYWEGLVENYQITLKSDET  339
NTDB id 1299 VP RS02975 WP 005483021.1      SAIWFFLTPQAPKPAMLHVIDRTWVPSQRELDAGIGYGFGTTINVINGGIECGEQNKDKGQPVNRIRYWEGLAAHYQIPTEADEK  340
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X $\geq$ 50% conserved