```
logo
NTDB id 174145 B34 RS00830 WP 080623899.1  MNVSSELLILLRLRGTLSPSLLTKWWKQDPSLSLTGEKNHLLTKLSLKRSDLCSIRKLAEQELSNVKRLIRSYEAAGVRMTAISS   85
NTDB id 114 BSU 16110 NP 389493.1          MDQAAVCLTICRINQLLSPSLLLKWWKADPSMSLTSPVLQTVTRDQIKA...AALKNEIEQFYPKLPRVLAAYREQGINTIPISS   82
consensus                                  !     *! **!*   !!!!!! !!!! !!!*!!! *  * *!*  *! **** **   !!  * **!**  !   !*   *!!!

logo
NTDB id 174145 B34 RS00830 WP 080623899.1  PDYPASLKTIHDPPPVLFLKGNKKLLHEQRLIGIVGTRCPSLYGKRAAVQLVRELCKNSWTIVSGLAKGIDGLAHQESIRSKGRT   170
NTDB id 114 BSU 16110 NP 389493.1          KQYPFWLKSIYDPPAVLFAKGDMTLLSKGRKIGIVGTRNPTAYGKQVVNHLTKEICRKGWVIVSGLASGIDGMSHAASIKAKGRT   167
consensus                                  * !!  !!*!*!!!*!!! !!   !!*  ! !!!!!!!*!* !!!    *! *!*!*  ! !!!!!! !!!!* !  !!* !!!!

logo
NTDB id 174145 B34 RS00830 WP 080623899.1  IGVIAGGFNSIYPREHRQLAGQMAESHLLVSEHPPHIKPQKWHFPMRNRLISGLTEGIVVVQGKEKSGSLITAYQALEQGREVFA   255
NTDB id 114 BSU 16110 NP 389493.1          IGVIAGGFQHIYPRENLQLADHMAKHHILLSEHPPETKPQKWHFPMRNRIISGLSEGVIVVQGKEKSGSLITAYQALEQGREVFA   252
consensus                                  !!!!!!!!**!!!!!* !!! *!! *!*!*!!!!!* !!!!!!!!!!!!*!!!!*!!**!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 174145 B34 RS00830 WP 080623899.1  VPGPIFDANSFGPSRLIQEGAKLVLTIEDILNELPPSNAQYTEPV   300
NTDB id 114 BSU 16110 NP 389493.1          VPGSLFDPYAGGPIKLIQQGAKAIWSAEDIFEELPERNVQYTEPF   297
consensus                                  !!!**!!*   !! *!!! !!! * * !!!  !!!* ! !!!!!
```

X non conserved

X similar

X ≥ 50% conserved