```
logo
NTDB id 41095 SGPB RS09855 WP 013852323.1  MKKYEQIFKDLEHDITSGIYQAGDYLPTENELTQIYHMSRDTIRKALSLLAEAGLIQKIHGSGSQVIKHEQIDFPVSQLTSYKEL   85
NTDB id 440 SMU RS09330 WP 002273717.1     MKKYEIIFKKLEEDILKGHYQMGDYLPPEMELSQTYASSRDTVRKALQLLTKAGFIKTVQGRGSQIIKRERINFPVSQLTSYQEL   85
consensus                                  !!!!! !!! !!*!!  !*!! !!!!!*! !!*! !* !!!!*!!!! !!  !! !  **! !!!*!!*! ! !!!!!!!!! !!

logo
NTDB id 41095 SGPB RS09855 WP 013852323.1  VDAHHINSKTNVIAIDKLIVDEKLATLTGFNKNNLVWRIIRQRVVDGTASVLDIDYLDKKIVPNMTREIAEQSIYDYLENQLHLI  170
NTDB id 440 SMU RS09330 WP 002273717.1     VKQLQMNVKTNVIAIDKLIVDEKLTKLTGFENKGLVWRITRQRVIDGVASILDTDYLDKALIPHMTREIAEHSIYDYLENQLKLD  170
consensus                                  !  ***! !!!!!!!!!!!!!!!!  !!!!   !!!!! !!!!*!! !!*!! !!!!! **!*!!!!!!!*!!!!!!!!!!*!

logo
NTDB id 41095 SGPB RS09855 WP 013852323.1  IAYAQKEITIDQITNRDKLLLDIGAEHHVVSVKSKVYLANKHQFQFTESRHKLEKFRFVDFATRKPQ  237
NTDB id 440 SMU RS09330 WP 002273717.1     IAYAQKIITIDQVSQKDKILLDLDSENHVVSVKSKVYLSNQQQFQFTESRHKLEKFRFVDFARRHRD  237
consensus                                  !!!!!! !!!!!****!!*!!!*  !*!!!!!!!!!!! ! *!!!!!!!!!!!!!!!!!!!! !**
```

X non conserved

X similar

X ≥ 50% conserved