logo

```
NTDB id 35968 B938 RS05360 WP 015239527.1  MANQKKKTLPPQHQNRQPGFEYVMDPRPVFDKPKAAKKLEGKTAVITGGDSGIGRAVSVLFAKEGANVVIVYFDEHQDAEETKQY  85
NTDB id 122 BSU 10400 NP 388921.1          MANQKKKTLPPQHQNQQPGFEYLMDPRPVFDKPKKAKKLEGKTAIITGGDSGIGRAVSVLFAKEGANVVIVYLNEHQDAEETKQY  85
consensus                                  !!!!!!!!!!!!!!! !!!!!!*!!!!!!!!!!! !!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!  !!!!!!!!!!!
```

logo

```
NTDB id 35968 B938 RS05360 WP 015239527.1  VEKEGAKCLLIAGDVGDEAFCNDVIRQAGQAFPSIDILVNNAGEQHVQPGIEKITSHQLIRTFQTNIFSMFYLTKAALPHLKKGS  170
NTDB id 122 BSU 10400 NP 388921.1          VEKEGVKCLLIAGDVGDEAFCNDVVGQASQVFPSIDILVNNAAEQHVQPSIEKITSHQLIRTFQTNIFSMFYLTKAVLPHLKKGS  170
consensus                                  !!!!! !!!!!!!!!!!!!!!!!!* !! ! !!!!!!!!!!!*!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!
```

logo

```
NTDB id 35968 B938 RS05360 WP 015239527.1  SIINTASITAYKGHKTLIDYSATKGAIVTFTRSLSQSLVTQGIRVNAVAPGPIWTPLIPASFSAKEVEVFGSDVPMQRPGEPVEV  255
NTDB id 122 BSU 10400 NP 388921.1          SIINTASITAYKGNKTLIDYSATKGAIVTFTRSLSQSLVQQGIRVNAVAPGPIWTPLIPASFAAKDVEVFGSDVPMERPGQPVEV  255
consensus                                  !!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!! !!*!!!!!!!!!! !!! !!!!!
```

logo

```
NTDB id 35968 B938 RS05360 WP 015239527.1  APSYLYLASDDSSYVTGQTIHVNGGTIVNG  285
NTDB id 122 BSU 10400 NP 388921.1          APSYLYLASDDSTYVTGQTIHVNGGTIVNG  285
consensus                                  !!!!!!!!!!!!!*!!!!!!!!!!!!!!!!
```

X non conserved

X similar

X ≥ 50% conserved