```
logo
```



```
NTDB id 56056 NC3 RS05155 WP 008348806.1  MSTKQPKKTLPPQHQNERPGLEYKMNPRPIFDREVQGKKLTGKTAIVTGGDSGIGRAVSVLFAKEGANVAIVYLNEHRDAEETKD  85
NTDB id 122 BSU 10400 NP 388921.1         .MANQKKKTLPPQHQNQQPGFEYLMDPRPVFDKPKKAKKLEGKTAIITGGDSGIGRAVSVLFAKEGANVVIVYLNEHQDAEETKQ  84
consensus                                 *    !*!!!!!!!!!!  !! !! ! !!!*!!**  *!!! !!!!!*!!!!!!!!!!!!!!!!!!!!!!  !!!!!!! !!!!!!
```

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logo
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```
NTDB id 56056 NC3 RS05155 WP 008348806.1  YIEKMGGRVILIAGDLGDEAFSHEVVKQTKDAFGSIDILVNNAGEQHPQKSIEQITSHQLLRTFQTNIFAMFYLTKAVLPHLKKG  170
NTDB id 122 BSU 10400 NP 388921.1         YVEKEGVKCLLIAGDVGDEAFCNDVVGQASQVFPSIDILVNNAAEQHVQPSIEKITSHQLIRTFQTNIFSMFYLTKAVLPHLKKG  169
consensus                                 !*!! ! ***!!!!!*!!!!!***!! !    !*!!!!!!!!*!!!*!*!!! !!!!!!*!!!!!!!! !!!!!!!!!!!!!!!
```

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logo
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```
NTDB id 56056 NC3 RS05155 WP 008348806.1  SSIINTASVTAYKGHETLIDYASTKGAVVTFTRSLSLSLIKQGIRVNGVAPGPIWTPLIPSTFTEKEVSEFGGDVPMERPGEPVE  255
NTDB id 122 BSU 10400 NP 388921.1         SSIINTASITAYKGNKTLIDYSATKGAIVTFTRSLSQSLVQQGIRVNAVAPGPIWTPLIPASFAAKDVEVFGSDVPMERPGQPVE  254
consensus                                 !!!!!!!!*!!!!!* !!!!!  !!!!*!!!!!!!! !!* !!!!!!*!!!!!!!!!!!! *!  !*!  !! !!!!!!!! !!!
```

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logo
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```
NTDB id 56056 NC3 RS05155 WP 008348806.1  LAPSYLFLASEDSSYMNGQMIHVNGGTILNG  286
NTDB id 122 BSU 10400 NP 388921.1         VAPSYLYLASDDSTYVTGQTIHVNGGTIVNG  285
consensus                                 *!!!!!*!!!*!!*!* !! !!!!!!!!*!!
```

X non conserved
X similar
X ≥ 50% conserved