```
logo
```

```
NTDB id 1162905 ABNR97 RS03275 WP 010951046.1  MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSYIVLIAWSSYFTSRKLNDVANERLAIQQDLRNAATLIVRDARMAGSFG  80
NTDB id 1138 NGFG RS02430 WP 003694978.1       MKRKMLNVPKGGYDGMKGFTIVEFLVAGLLSVIVLIAVVSSYFTSRKLNDAANERLAEQQDLRNAATLIVRDARMAGSFG  80
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

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logo
```

```
NTDB id 1162905 ABNR97 RS03275 WP 010951046.1  CFNMSEHTKDDIVDSSNQTQSNLAKPGAKQENPLFSLKRSG..MDKQLIPVAESIDIKYPGFIQRLNALVFQYGIDDLDA  158
NTDB id 1138 NGFG RS02430 WP 003694978.1       CFNMSEHIGNDVVFNV............AQKNALFSLKRNSTNSTNKLIPITESPNINYQNFFQVSSALIFQYGIDDVDA  148
consensus                                      !!!!!!!   !*!   ************ ! !*!!!!!!  **    !!!* !!* ! !* ! !   !!*!!!!!!!*!!
```

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logo
```

```
NTDB id 1162905 ABNR97 RS03275 WP 010951046.1  SAETVVVSSCSKIAKPGKKISTLQEAKSALQITNDD.KQNGNITRQKHVVNAYAVGRFGNNEESLFRFQLDDKGKWGNPQ  237
NTDB id 1138 NGFG RS02430 WP 003694978.1       SADTTVVSSCAAISKPGKQIPTLENAKKELKIQNSDKEQNGNIARQRHVVNAYAVGKIAG.EEGLFRFQLDDKGKWGNPQ  227
consensus                                      !!*! !!!!!  ! !!!! !*!!  !!  ! ! ! !* !!!!! !!*!!!!!!!!!* * *!! !!!!!!!!!!!!!!!!
```

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logo
```

```
NTDB id 1162905 ABNR97 RS03275 WP 010951046.1  LLVKKVKRMDVRYIYVSGCPEDEDAGKEEKFRYTNKFDKSKNAVTPAGVEVLLDSGLNAKIAASSDNSIYAYRINATIRG  317
NTDB id 1138 NGFG RS02430 WP 003694978.1       LLAKKVKRMRVRYIYVSGCPEDEDAGKEEQFKYTDKFDS...SVTPAGVEVLLDSGSDAKIAASSDNIIYAYRINATIRG  304
consensus                                      !! !!!!!! !!!!!!!!!!!!!!!!!!!! !*!! !!! *** !!!!!!!!!!!!!!  !!!!!!!!! !!!!!!!!!!!!
```

```
logo
```

```
NTDB id 1162905 ABNR97 RS03275 WP 010951046.1  GNVCANRTL  326
NTDB id 1138 NGFG RS02430 WP 003694978.1       GNVCANRTL  313
consensus                                      !!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved