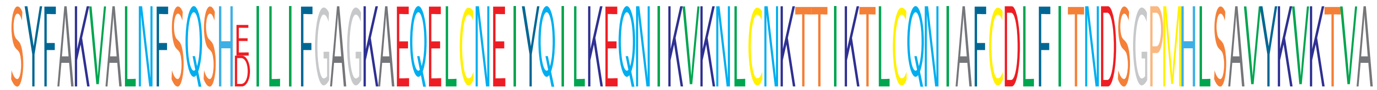
```
logo
```

```
NTDB id 84956 RB95 RS05495 WP 002933188.1  MKIFIHLPTWLGDAVMASPALYTIKEHFKNAQFILYGSLVSTALFKEFPNSKIIIENKQSRYKQALSLHKELGKIDLSFAFRSAF  85
NTDB id 1241 Cj1148 YP 002344539.1         MKIFIHLPTWLGDTVMASPALYTIKEHFKDAQFILYGSFVSTALFKEFPNSKIIIENKLSRYKQALSLRKELGKIDLSFAFRSAF  85
consensus                                  !!!!!!!!!!!!! !!!!!!!!!!!!!!! !!!!!!!! !!!!!!!!!!!!!!!!!!! !!!!!!!!!*!!!!!!!!!!!!!!!!
```

```
logo
```

```
NTDB id 84956 RB95 RS05495 WP 002933188.1  SSKIILHILKTKQRYFFDKNKHKEEHQVLKYLYFIENSLSIKAHSKDLKLPFKLKFQNPLVLKNGKKILGLNPGASFGSAKRWDA  170
NTDB id 1241 Cj1148 YP 002344539.1         SSKIILHILKTKQRYFFDKNKYKEEHQVLKYLYFIENSLSIKAHFKDLKLPFKLKFQNPLILRNGKKILGLNPGASFGSAKRWDA  170
consensus                                  !!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!*!*!!!!!!!!!!!!!!!!!!!!!!
```

```
logo
```


```
NTDB id 84956 RB95 RS05495 WP 002933188.1  SYFAKVALNFSQSHEILIFGAGKAEQELCNEIYQILKEQNIKVKNLCNKTTIKTLCQNIAFCDLFITNDSGPMHLSAVYKVKTVA  255
NTDB id 1241 Cj1148 YP 002344539.1         SYFAKVALNFSQSHDILIFGAGKAEQELCNEIYQILKEQNIKVKNLCNKTTIKTLCQNIAFCDLFITNDSGPMHLSAVYKVKTVA  255
consensus                                  !!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

```
logo
```

```
NTDB id 84956 RB95 RS05495 WP 002933188.1  IFGPTKFTQTSPWQNENATLVHLDLACMPCMQKTCPLKHHKCMKDLKPERVIEEIKKLSTP...   316
NTDB id 1241 Cj1148 YP 002344539.1         IFGPTKFTQTSPWQNENAKLVHLDLACMPCMQKTCPLKHHKCMKDLKPEKVIEQAKNLLKNSHL   319
consensus                                  !!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!  ! !  ****
```

X non conserved
X similar
X $\geq$ 50% conserved