logo

```
NTDB id 50096 HT078 RS06110 WP 123771562.1  MRKQNTLTGIPTSDGQRGSALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGEFQVLDLEYAADSKVT   85
NTDB id 1139 NGFG RS02435 WP 003687918.1    MRKQNTLTGIPTSDGQRGSALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGEFQVLDLEYTADSKVT   85
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!
```

logo

```
NTDB id 50096 HT078 RS06110 WP 123771562.1  FSENCEKGLCTAVNVRTNNNGSEEAFGNIVVQGKPIVEAVKRSCPAKSGKNSTGLCIDNEGMEYKKGTAGVSKMPRYIIEYLGVK   170
NTDB id 1139 NGFG RS02435 WP 003687918.1    FSENCEKGLCTAVNVRTNNNGNEEVFGNIVVQGTPTVEAVKRSCPAKSGKNSTGLCIDNQGVEYEKGTGNVSKMPRYIIEYLGEK   170
consensus                                   !!!!!!!!!!!!!!!!!!!!! !! !!!!!!!! ! !!!!!!!!!!!!!!!!!!!!!!! !*!! !!!* !!!!!!!!!!!!! !
```

logo

```
NTDB id 50096 HT078 RS06110 WP 123771562.1  NGQNVYRVTAKAWGKNANTVVVLQSYVGNNDEQ   203
NTDB id 1139 NGFG RS02435 WP 003687918.1    NNQNIYRVTAKAWGKNANTVVVLQSYVGNNDEQ   203
consensus                                   ! !!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved