```
logo
```

```
NTDB id 304750 DV129 RS01695 WP 114865959.1  MSEIGFKYSILASGSSGNSFYLETSKNKLLVDAGLSGKKITSLLAEINRKPEDLDAILITHEHSDHIHGVGVLARKYGMD  80
NTDB id 383 SMU RS06875 WP 002262928.1       MTETGFRYSILASGSSGNSFYLETPQKRILIDAGLSGKKISQLMAEIDRKVDDVDAILVTHEHKDHIHGVGVLARKYGLD  80
consensus                                    !*!  !!*!!!!!!!!!!!!!!!!!!*  **!*!!!!!!!!!* !*!!!  !!**!*!!!!*!!!!  !!!!!!!!!!!!!!*!
```

```
logo
```

```
NTDB id 304750 DV129 RS01695 WP 114865959.1  LYANEKTWQAMENSKYLGKVDSSQKHIFEMGKTKTFGDIDIESFGVSHDAVAPQFYRFMKDDKSFVLLTDTGYVSDRMAG  160
NTDB id 383 SMU RS06875 WP 002262928.1       VYANAATWQAMDG..MIGKIDVSQKHIFDLGKTITFGDIDIESFGVSHDAAQPQFYRLMKDGKSFVILTDTGYVSDRMAD  158
consensus                                    *!!!  !!!!!* ** *!!*! !!!!!!**!!! !!!!!!!!!!!!!!!!  !!!!! !!! !!!!*!!!!!!!!!!!!
```

```
logo
```

```
NTDB id 304750 DV129 RS01695 WP 114865959.1  IVENADGYLIEANHDVEILRSGSYAWRLKQRILSDLGHLSNEDGAEAMIRTLGNRTKKIYLGHLSKENNIKELAHMTMVN  240
NTDB id 383 SMU RS06875 WP 002262928.1       LIENADGYLIESNHDIEILRSGIYPWSLKQRILSDKGHLSNDDGAETMVRTLGNKTKKIYLGHLSKENNVKELAHMTMKN  238
consensus                                    **!!!!!!!!! !!!*!!!!!! !*! !!!!!!!! !!!!!*!!!! !*!!!!!*!!!!!!!!!!!!!!!*!!!!!!!! !
```

```
logo
```

```
NTDB id 304750 DV129 RS01695 WP 114865959.1  QLAQADLGVGVDFKVYDTSPDTPTPLTEI  269
NTDB id 383 SMU RS06875 WP 002262928.1       ALMQADLAVDHDFKIYDTSPDTALPLAKI  267
consensus                                     ! !!!!*! *!!!*!!!!!!!* !!  !
```

X non conserved
X similar
X ≥ 50% conserved