```
logo
NTDB id 79297 HRS36 RS08525 WP 173236978.1   MTESAQLVNCTFPSEALLYNAYMPFVKGGGLFFRTKYVYPLGKELQLSLKIISEIEPYLIDVKVVWVTPKGAQGNKPPGIGVQFI   85
NTDB id 1275 LPP RS06810 WP 025519545.1      MTEKPQLINCSFPNEASLYMAYMPFVKGGGLFIRTQHVYSLGQSLELSVKLLNENEPYLVTGKIVWITPKGAQGNKPPGVGIQFT   85
NTDB id 1261 GCO85 RS07110 WP 025519545.1    MTEKPQLINCSFPNEASLYMAYMPFVKGGGLFIRTQHVYSLGQSLELSVKLLNENEPYLVTGKIVWITPKGAQGNKPPGVGIQFT   85
consensus                                    !!!**!!*!!*!!*!!*!!*!!!!!!!!!!!!!*!!**!!*!!**!*!!*!***!*!!!!***!*!!*!!!!!!!!!!!!!*!*!!*
```

```
logo
NTDB id 79297 HRS36 RS08525 WP 173236978.1   GENSRYLCNKIETHLAGMLKSTQPTDTI   113
NTDB id 1275 LPP RS06810 WP 025519545.1      GENSRYLCNKIETYLAGMLKSCQVTDTI   113
NTDB id 1261 GCO85 RS07110 WP 025519545.1    GENSRYLCNKIETYLAGMLKSCQVTDTI   113
consensus                                    !!!!!!!!!!!!!*!!!!!!!*!*!!!!
```


X non conserved

X similar

X $\geq$ 50% conserved