```
logo
NTDB id 374 SMU RS06885 WP 002262930.1          ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK   69
NTDB id 112727 PFLUOLIPICF7 RS10060 WP 010207217.1 MEQHSNALRVMVIDDSKTIRRTAETLLKNVGCDVITAIDGFDALARIVDHHPHIIFVDIMMPRLDGYQTCALVKN   75
NTDB id 1047 HON27 RS03110 WP 000389061.1       MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKN   75
consensus                                       **   * ******!!*!*!*******! **!***!!!*!!**!!*** * *!*****!*!*!*!!!*********

logo
NTDB id 374 SMU RS06885 WP 002262930.1          ...NSHVPIIMLSAKDSEFDKVIGLEICADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGD  141
NTDB id 112727 PFLUOLIPICF7 RS10060 WP 010207217.1 NPAFKSIPVIMLSSKDGLFDKAKGRIVGVDQFLTKPFSKDELLGAIKAYVPGFAAVEQAH...............  135
NTDB id 1047 HON27 RS03110 WP 000389061.1       SQNYQNIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.......................  127
consensus                                          *  *!*!!!!*!!**!!***!***! !***!!!!!**!!!*******     ** *

logo
NTDB id 374 SMU RS06885 WP 002262930.1          LQILPDAFVAKKRGTEVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPS  216
NTDB id 112727 PFLUOLIPICF7 RS10060 WP 010207217.1 ..........................................................................  135
NTDB id 1047 HON27 RS03110 WP 000389061.1       ..........................................................................  127
consensus

logo
NTDB id 374 SMU RS06885 WP 002262930.1          RPEYILTRRGVGYYMKSYD   235
NTDB id 112727 PFLUOLIPICF7 RS10060 WP 010207217.1 ...................   135
NTDB id 1047 HON27 RS03110 WP 000389061.1       ...................   127
consensus
```

X non conserved
X similar
X ≥50% conserved