```
logo
NTDB id 1063032 ACG4I4 RS12260 WP 001188316.1  MQRANYARTIYLLTTQPKALHPSIQAAIEQLNLPVPVIEPERLLREYQSDKHKILLLDHAENGLIRQQLGPLKLTSPYFE   80
NTDB id 1146 A1552VC RS00790 WP 001188316.1    MQRANYARTIYLLTTQPKALHPSIQAAIEQLNLPVPVIEPERLLREYQSDKHKILLLDHAENGLIRQQLGPLKLTSPYFE   80
NTDB id 1296 VP RS13295 WP 005480993.1         MRKSAYARKLFLISMEDDAAQ.KVASLEKYIDMSIPVISTDALMEAKPEHRNKILLIDFSEHKSLVQSIKNLPLVWKNFE   79
NTDB id 1394 DSB67 RS13765 WP 005432827.1      MKKSAYARKLFLISMEEDAGK.KVEALEKYTDIDIPVISTQALMEANPKHRNKILLIDFSEHKELVLSIKNLPLIWKNFE   79
consensus                                      !**  !!! **!**  * !*** ***    * ***!!! ** !*  ******!!!!*!* !* ** * * *!*!* * !!

logo
NTDB id 1063032 ACG4I4 RS12260 WP 001188316.1  TILFNVEKRLRTEDLLTFGNLKGLFYANEDTGFIAHGLGEIINGQNWLPRHVSSQLLHYYRYAFQSHQVQATVDLTAREI  160
NTDB id 1146 A1552VC RS00790 WP 001188316.1    TILFNVEKRLRTEDLLTFGNLKGLFYANEDTGFIAHGLGEIINGQNWLPRHVSSQLLHYYRYAFQSHQVQATVDLTAREI  160
NTDB id 1296 VP RS13295 WP 005480993.1         TVVFNVPKRLTTDELLAFGQLKGLFYSEDSLEQVGEGLKGIVNGQNWLPRNVTSQLLHYYRNVINTHTAPATVDLTIREL  159
NTDB id 1394 DSB67 RS13765 WP 005432827.1      TVVFNVPKRLTTDELLAFGQLKGIFYEESSLEQIGEGLKEIINGQNWLPRNVTSQLLHYYRNVISTHTAPATVDLTIREL  159
consensus                                      !**!!!*!!!  !**!! !!*!!!*!!* *    ***!! *!*!!!!!!!!!*!*!!!!!!!!   **!  *!!!!!! !!*

logo
NTDB id 1063032 ACG4I4 RS12260 WP 001188316.1  QILRCLQTGASNMQIAENLFISEFTVKSHLYQIFKKLNVKNRVKAIAWVNQNLL.  214
NTDB id 1146 A1552VC RS00790 WP 001188316.1    QILRCLQTGASNMQIAENLFISEFTVKSHLYQIFKKLNVKNRVKAIAWVNQNLL.  214
NTDB id 1296 VP RS13295 WP 005480993.1         QVLRCLQAGASNSQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS  214
NTDB id 1394 DSB67 RS13765 WP 005432827.1      QVLRCLQAGASNNQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS  214
consensus                                      !*!!!!! !!!!*!*!! !!*!!!!*!!!!!!!!!!!! !!!!! !!!!  !!!**
```

X non conserved

X similar

X $\geq$ 50% conserved