```
NTDB id 1146 A1552VC RS00790 WP 001188316.1  MQRANYARTIYLLTTQPKALHPSIQAAIEQLNLPVPVIEPERLLREYQSDKHKILLLDHAENGLIRQQLGPLKLTSPYFE   80
NTDB id 1394 DSB67 RS13765 WP 005432827.1    MKKSAYARKLFLISMEEDAG.KKVEALEKYTDIDIPVISTQALMEANPKHRNKILLIDFSEHKELVLSIKNLPLIWKNFE   79
NTDB id 27110 VEA RS03955 WP 006742331.1     MRKSAYARKLFLISMEDNAQ.KKVAALEKYIDMSIPVISTEALMEAKPQHRNKILLIDFSEHKSFVQSIKNLPLVWKNFE   79
NTDB id 1296 VP RS13295 WP 005480993.1       MRKSAYARKLFLISMEDDAA.QKVASLEKYIDMSIPVISTDALMEAKPEHRNKILLIDFSEHKSLVQSIKNLPLVWKNFE   79
consensus                                    !****!!!***!******!* **************!!!****!***** ***!!!!*!**!**********!*!****!!
```

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NTDB id 1146 A1552VC RS00790 WP 001188316.1  TILFNVEKRLRTEDLLTFGNLKGLFYANEDTGFIAHGLGEIINGQNWLPRHVSSQLLHYYRYAFQSHQVQATVDLTAREI  160
NTDB id 1394 DSB67 RS13765 WP 005432827.1    TVVFNVPKRLTTDELLAFGQLKGIFYEESSLEQIGEGLKEIINGQNWLPRNVTSQLLHYYRNVISTHTAPATVDLTIREL  159
NTDB id 27110 VEA RS03955 WP 006742331.1     TVVFNVPKRLTTDELLSFGQLKGLFYSEDKLEQIGEGLKGIVNGQNWLPRNVTSQLLHYYRNVINTHTAPATVDLTIREL  159
NTDB id 1296 VP RS13295 WP 005480993.1       TVVFNVPKRLTTDELLAFGQLKGLFYSEDSLEQVGEGLKGIVNGQNWLPRNVTSQLLHYYRNVINTHTAPATVDLTIREL  159
consensus                                    !**!!!*!!!*!**!!*!!*!!!*!!***********!!* !*!!!!!!!!!*!*!!!!!!!!*****!***!!!!!!*!!*
```

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NTDB id 1146 A1552VC RS00790 WP 001188316.1  QILRCLQTGASNMQIAENLFISEFTVKSHLYQIFKKLNVKNRVKAIAWVNQNLL.   214
NTDB id 1394 DSB67 RS13765 WP 005432827.1    QVLRCLQAGASNNQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS   214
NTDB id 27110 VEA RS03955 WP 006742331.1     QVLRCLQAGASNSQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS   214
NTDB id 1296 VP RS13295 WP 005480993.1       QVLRCLQAGASNSQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS   214
consensus                                    !*!!!!!*!!!!*!*!!*!!*!!!!*!!!!!!!!!!!!!*!!!!!*!!!!**!!!**
```

X non conserved
X similar
X ≥ 50% conserved