```
logo
NTDB id 1146 A1552VC RS00790 WP 001188316.1   MQRANYARTIYLLTTQPKALHPSIQAAIEQLNLPVPVIEPERLLREYQSDKHKILLLDHAENGLIRQQLGPLKLTSPYFE   80
NTDB id 91723 OCV52 RS01550 WP 137407230.1   MRKSRYARTLHFMCIDPSDTYLHVKEIEKHLSIILYKMTPGDLMLVDRKQSNRILLVDYKQVPELLATCPNLTVMWKSHE   80
NTDB id 1296 VP RS13295 WP 005480993.1   MRKSAYARKLFLISMEDD.AAQKVASLEKYIDMSIPVISTDALMEAKPEHRNKILLIDFSEHKSLVQSIKNLPLVWKNFE   79
NTDB id 1394 DSB67 RS13765 WP 005432827.1   MKKSAYARKLFLISMEED.AGKKVEALEKYTDIDIPVISTQALMEANPKHRNKILLIDFSEHKELVLSIKNLPLIWKNFE   79
consensus                                   !****!!! ******** ** ** ******* ********!*** ******!!!*!************!*******!

logo
NTDB id 1146 A1552VC RS00790 WP 001188316.1   TILFNVEKRLRTEDLLTFGNLKGLFYANEDTGFIAHGLGEIINGQNWLPRHVSSQLLHYYRYAFQSHQVQATVDLTAREI   160
NTDB id 91723 OCV52 RS01550 WP 137407230.1   IILFNVPQQLPTSELLIFGVLKGLFYNTDKKDKIAHGLQEIINGENWLPRKVTNQLLFYYRNMVNTNTTPTNVDLTIREI   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 1146 A1552VC RS00790 WP 001188316.1   QILRCLQTGASNMQIAENLFISEFTVKSHLYQIFKKLNVKNRVKAIAWVNQNLL.   214
NTDB id 91723 OCV52 RS01550 WP 137407230.1   QVIRCLQSGSSNTQIADDLFISEFTVKSHLYQIFRKLAVKNRVQAIAWANQNLLA   215
NTDB id 1296 VP RS13295 WP 005480993.1   QVLRCLQAGASNSQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS   214
NTDB id 1394 DSB67 RS13765 WP 005432827.1   QVLRCLQAGASNNQMAEELFVSEFTIKSHLYQIFKKLSVKNRVQAIAWADQNLMS   214
consensus                                   !**!!!!*!*!!*!*!**!!*!!!!*!!!!!!!!!*!!*!!!!!*!!!!* !!!**
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X non conserved
X similar
X $\geq$ 50% conserved