```
logo
NTDB id 1127 NGFG RS10560 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLA..................AVNINAASQQEL   63
NTDB id 1126 NGFG RS09375 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLA..................AVNINAASQQEL   63
NTDB id 1125 NGFG RS07215 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLA..................AVNINAASQQEL   63
NTDB id 1108 NGFG RS06525 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLA..................AVNINAASQQEL   63
NTDB id 1272 LPP RS04360 WP 010946546.1      ............................MK.ARLIAILLSFIV..I.PFSSQAENEPQKTPAKQVLVKGKINLNTADVYSL   49
NTDB id 1249 GC085 RS04230 WP 010946546.1    ............................MK.ARLIAILLSFIV..I.PFSSQAENEPQKTPAKQVLVKGKINLNTADVYSL   49
NTDB id 23880 PBPR RS13395 WP 011219290.1    ..............................M.NTIIKMALFSVLLLLAPL......SQAA..NEYEGIEIKINNNANAEEL   43
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   ..........................MQIK.TKIVTLFLSLCLPTLPLLANAEETAPAA..QVEEGIVITVNINTASAEEL   52
NTDB id 1150 A1552VC RS08580 WP 001166105.1  ..........................MQIK.TKIVTLFLSLCLPTLPLLANAEETAPAA..QVEEGIVITVNINTASAEEL   52
NTDB id 1395 DSB67 RS04445 WP 010649155.1    ..................................MKWMLTLCLLILAPMSWAE.TKTKA..DKYEGIEITVNVNSATAQEI   44
NTDB id 1294 VP RS04495 WP 021449594.1       ..................................MKWILTLCLVVFAPLSLAA..DTKA..DKYEGIEITVNINTASAEEI   43
consensus                                                                   **   *********** *  ** *              *   *!*!*!*  **

logo
NTDB id 1127 NGFG RS10560 WP 003703428.1     E.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1     E.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1     E.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1     E.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1272 LPP RS04360 WP 010946546.1      TGSFKGIGKKRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN.............   105
NTDB id 1249 GC085 RS04230 WP 010946546.1    TGSFKGIGKKRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN.............   105
NTDB id 23880 PBPR RS13395 WP 011219290.1    DKLLIGIGPEKAESIIIYRKEKGDFQTADDLINVKGIGLATVDKNRERIEL...................    94
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   ATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...................   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1  ATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...................   103
NTDB id 1395 DSB67 RS04445 WP 010649155.1    ATLLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL...................    95
NTDB id 1294 VP RS04495 WP 021449594.1       ATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL...................    94
consensus                                       * !!! **! *!**!! ****!*****!  !!!!! * * ****** *
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X non conserved

X similar

X ≥ 50% conserved