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NTDB id 1127 NGFG RS10560 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFS.....................................  45
NTDB id 1126 NGFG RS09375 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFS.....................................  45
NTDB id 1125 NGFG RS07215 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFS.....................................  45
NTDB id 1108 NGFG RS06525 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFS.....................................  45
NTDB id 1440 BW25113 RS02290 WP 000680312.1   ........................MKHGIKALLITLSLACAGMSHSALAAASVAKPTAVETKAEAPAAQ.SKAAVPAKA  54
NTDB id 1459 ASU2 RS10590 WP 015674357.1      ........................MKTPSK...LGLSL.LIGMLSTGVLAAEKTKPSVTEPTPVVQNAQATSPEHSHAE  51
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    ........................MQIKTKIVTLFLSL.CLPTL...................PLLANAEETAPA..AQV  34
NTDB id 1150 A1552VC RS08580 WP 001166105.1   ........................MQIKTKIVTLFLSL.CLPTL...................PLLANAEETAPA..AQV  34
NTDB id 1294 VP RS04495 WP 021449594.1        ..............................MKWILTL.CLVVF...................APLSLAA..DTK..ADK  25
NTDB id 1395 DSB67 RS04445 WP 010649155.1     ..............................MKWMLTL.CLLIL...................APMSWAE.TKTK..ADK  26
consensus                                                                   ** * ** ***! *** *                 * * *     *
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NTDB id 1127 NGFG RS10560 WP 003703428.1      CAFSLAAVNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKP  124
NTDB id 1126 NGFG RS09375 WP 003703428.1      CAFSLAAVNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKP  124
NTDB id 1125 NGFG RS07215 WP 003703428.1      CAFSLAAVNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKP  124
NTDB id 1108 NGFG RS06525 WP 003703428.1      CAFSLAAVNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKP  124
NTDB id 1440 BW25113 RS02290 WP 000680312.1   SDEEGTRVSINNASAEELARAMNGVGLKKAQAIVSYREEYGPFKTVEDLKQVPGMGNSLVERNLAVLTL...........  123
NTDB id 1459 ASU2 RS10590 WP 015674357.1      MTNNANLVNINIATAAEIQDKLVGIGAKKAQAIVEYRTKNGAFSSLEQLTDVSGIGKATLDKNRDRILLQ..........  121
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    EEGIVITVNINTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...........  103
NTDB id 1150 A1552VC RS08580 WP 001166105.1   EEGIVITVNINTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...........  103
NTDB id 1294 VP RS04495 WP 021449594.1        YEGIEITVNINTASAEEIATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL...........   94
NTDB id 1395 DSB67 RS04445 WP 010649155.1     YEGIEITVNVNSATAQEIATLLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL...........   95
consensus                                      *  *  !**! !***!***** !*! *!!**!**!!  *!*!*****!**!*!*! *** ********
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NTDB id 1127 NGFG RS10560 WP 003703428.1      VLPAVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1      VLPAVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1      VLPAVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1      VLPAVKK   131
NTDB id 1440 BW25113 RS02290 WP 000680312.1   .......   123
NTDB id 1459 ASU2 RS10590 WP 015674357.1      .......   121
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    .......   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1   .......   103
NTDB id 1294 VP RS04495 WP 021449594.1        .......    94
NTDB id 1395 DSB67 RS04445 WP 010649155.1     .......    95
consensus
```

X non conserved
X similar
X ≥ 50% conserved