```
logo
NTDB id 1294 VP RS04495 WP 021449594.1      ................................MKWI.LTLCLVVFAPLSLAADTKADKYEGIEITVNINTASAEEIATMLKGIGE   52
NTDB id 1242 Cj0011c YP 002343483.1         ................................MKKL...LFLFFALTAF............LFGAVNINTATLKELK.SLNGIGE   37
NTDB id 26326 NEUT RS10540 WP 011635093.1   ................................MKKIFLILVIFFGLNLS............VLAGVDINTASQIDLE.SVKGLGP   40
NTDB id 1127 NGFG RS10560 WP 003703428.1    MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFS.............LAAVNINAASQQELE.ALPGIGP   71
NTDB id 1126 NGFG RS09375 WP 003703428.1    MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFS.............LAAVNINAASQQELE.ALPGIGP   71
NTDB id 1125 NGFG RS07215 WP 003703428.1    MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFS.............LAAVNINAASQQELE.ALPGIGP   71
NTDB id 1108 NGFG RS06525 WP 003703428.1    MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFS.............LAAVNINAASQQELE.ALPGIGP   71
consensus                                   ********************************!!***************             ***!*!!*!****** ***!*!*

logo
NTDB id 1294 VP RS04495 WP 021449594.1      KKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL...................   94
NTDB id 1242 Cj0011c YP 002343483.1         AKAKAILEYRKE.ANFTSIDDLKKVKGIGDKLFEKIKNDITIE..................   79
NTDB id 1127 NGFG RS10560 WP 003703428.1    AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
NTDB id 1126 NGFG RS09375 WP 003703428.1    AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
NTDB id 1125 NGFG RS07215 WP 003703428.1    AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
NTDB id 1108 NGFG RS06525 WP 003703428.1    AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
consensus                                   *!!**!**!!*****!****!!**!!!!!*****************************
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X non conserved

X similar

X ≥50% conserved