```
logo
```

```
NTDB id 1242 Cj0011c YP 002343483.1              ..............................MKKLLFLFFAL..TAFLFGAVNINTATLKELKSLNGIGEAKAKAILEY   46
NTDB id 1099448 ACM67E RS03535 WP 416191716.1    ............................MHSTLKKIFFYGLATLAASVSSAAVNINTASESELTALPGVGPAKAKAIVAY   52
NTDB id 1127 NGFG RS10560 WP 003703428.1         MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEY   80
NTDB id 1126 NGFG RS09375 WP 003703428.1         MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEY   80
NTDB id 1125 NGFG RS07215 WP 003703428.1         MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEY   80
NTDB id 1108 NGFG RS06525 WP 003703428.1         MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEY   80
consensus                                        ********************************!!***************!!!!!*!***!!**!*!*!*!!!!!!**!
```

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logo
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```
NTDB id 1242 Cj0011c YP 002343483.1              RKE.ANFTSIDDLKKVKGIGDKLFEKIKNDITIE.................   79
NTDB id 1099448 ACM67E RS03535 WP 416191716.1    RKQHGNFKTVDELKNVKGIGEGIFSKLRNEATVAAPVKKAAQP....AVKK   99
NTDB id 1127 NGFG RS10560 WP 003703428.1         RAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
NTDB id 1126 NGFG RS09375 WP 003703428.1         RAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
NTDB id 1125 NGFG RS07215 WP 003703428.1         RAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
NTDB id 1108 NGFG RS06525 WP 003703428.1         RAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK  131
consensus                                        !*****!***!*!**!!!!!*****!************************
```

X non conserved
X similar
X ≥ 50% conserved