```
logo
```

```
NTDB id 48028 NMBM04240196 RS14940 WP 162467419.1  MLCPEKMSVMAGRHPYGVRSGLRRNGLKLWDIHFRMTRFIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAA   75
NTDB id 1127 NGFG RS10560 WP 003703428.1           .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAA   52
NTDB id 1126 NGFG RS09375 WP 003703428.1           .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAA   52
NTDB id 1125 NGFG RS07215 WP 003703428.1           .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAA   52
NTDB id 1108 NGFG RS06525 WP 003703428.1           .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAA   52
consensus                                                               **!*!**  ***!***!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

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logo
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```
NTDB id 48028 NMBM04240196 RS14940 WP 162467419.1  VNINAASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLTKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLP   150
NTDB id 1127 NGFG RS10560 WP 003703428.1           VNINAASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLP   127
NTDB id 1126 NGFG RS09375 WP 003703428.1           VNINAASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLP   127
NTDB id 1125 NGFG RS07215 WP 003703428.1           VNINAASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLP   127
NTDB id 1108 NGFG RS06525 WP 003703428.1           VNINAASQQELEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLP   127
consensus                                          !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

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logo
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```
NTDB id 48028 NMBM04240196 RS14940 WP 162467419.1  ADKK   154
NTDB id 1127 NGFG RS10560 WP 003703428.1           AVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1           AVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1           AVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1           AVKK   131
consensus                                          !*!!
```

X non conserved
X similar
X ≥ 50% conserved