logo

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NTDB id 21583 NMB RS00320 WP 009348628.1  MLCPEKMSVMAGRHPYGVRSGLRRNGLKLWDIHFRMTRFIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE   85
NTDB id 1127 NGFG RS10560 WP 003703428.1   .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE   62
NTDB id 1126 NGFG RS09375 WP 003703428.1   .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE   62
NTDB id 1125 NGFG RS07215 WP 003703428.1   .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE   62
NTDB id 1108 NGFG RS06525 WP 003703428.1   .....................MSRGGAT..PYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQE   62
consensus                                                      **!*!**  ***!***!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
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logo

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NTDB id 21583 NMB RS00320 WP 009348628.1  LEALPGIG..............................PAVLAKLKDQASVGAPAPKGPAKPVLPADKK   124
NTDB id 1127 NGFG RS10560 WP 003703428.1   LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1   LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1   LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1   LEALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
consensus                                 !!!!!!!!******************************!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!
```

X non conserved
X similar
X $\geq$50% conserved