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logo

NTDB id 1242 Cj0011c YP 002343483.1 ..............................MKKLLFLFF...ALTAFLFGAVNINTATLKELKSLNGIGEAKAKAILEYRKE.  49
NTDB id 85650 KJS40 RS11210 WP 213537807.1 ..............................MKKLLFVIIALFTFTCAAYAAVNINTATQAELETLQGIGPAKAKAIIDHRKKN  53
NTDB id 1127 NGFG RS10560 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKK.MFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQN  84
NTDB id 1126 NGFG RS09375 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKK.MFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQN  84
NTDB id 1125 NGFG RS07215 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKK.MFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQN  84
NTDB id 1108 NGFG RS06525 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKK.MFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQN  84
consensus                           ********************************!!! *!**********!!!!!*!**!!**!*!!!*!!!!!!!***!***

logo

NTDB id 1242 Cj0011c YP 002343483.1 ANFTSIDDLKKVKGIGDKLFEKIKNDITIE.................  79
NTDB id 85650 KJS40 RS11210 WP 213537807.1 GSFKSADDLEKVNGIGPATLKRLRKDIAVGGATTVKKENKPVPK...  97
NTDB id 1127 NGFG RS10560 WP 003703428.1 GAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1126 NGFG RS09375 WP 003703428.1 GAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1125 NGFG RS07215 WP 003703428.1 GAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1108 NGFG RS06525 WP 003703428.1 GAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
consensus                           **!*!*!!!*!!*!!!*****************************
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X non conserved
X similar
X ≥ 50% conserved