```
logo

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
NTDB id 122899 NELON RS01865 WP 003775734.1  ................................MKKALFGLLSYAVTALALAAVNINTATVEELKALPGIGPAKAQAIVDY   48
NTDB id 1242 Cj0011c YP 002343483.1           ................................MKKLLF..LFFALTAFLFGAVNINTATLKELKSLNGIGEAKAKAILEY   46
consensus                                     ********************************!!!***********!****!!!!!*!***!!**!*!!!*!!!*!!**!

logo

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
NTDB id 122899 NELON RS01865 WP 003775734.1  RKENGNFKSVEDLKKVKGIGEGIFGKLKDEATVGTAAPAKTTKPAAEKPAAQPAAGKNTAK   109
NTDB id 1242 Cj0011c YP 002343483.1           RKE.ANFTSIDDLKKVKGIGDKLFEKIKNDITIE.........................   79
consensus                                     !*****!*!**!!*!!!!!!*****!*!*********************
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X non conserved

X similar

X $\geq$ 50% conserved