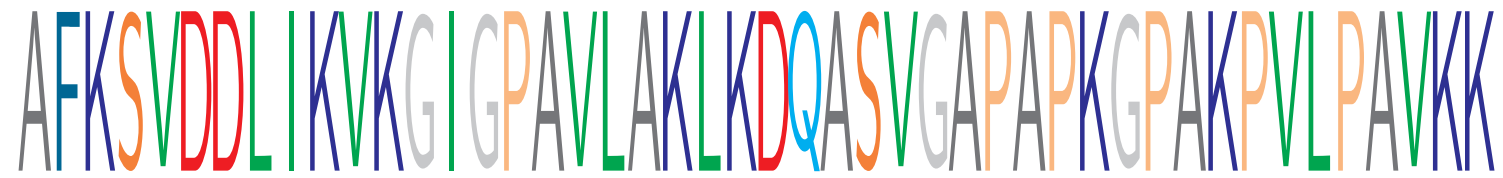
logo



NTDB id 1127 NGFG RS10560 WP 003703428.1	MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQNG	85
NTDB id 1126 NGFG RS09375 WP 003703428.1	MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQNG	85
NTDB id 1125 NGFG RS07215 WP 003703428.1	MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQNG	85
NTDB id 1108 NGFG RS06525 WP 003703428.1	MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSLAAVNINAASQQELEALPGIGPAKAKAIAEYRAQNG	85
consensus	!!!	

logo



NTDB id 1127 NGFG RS10560 WP 003703428.1	AFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK	131
NTDB id 1126 NGFG RS09375 WP 003703428.1	AFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK	131
NTDB id 1125 NGFG RS07215 WP 003703428.1	AFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK	131
NTDB id 1108 NGFG RS06525 WP 003703428.1	AFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK	131
consensus	!!	

X non conserved

X similar

X $\geq$ 50% conserved