logo
NTDB id 285101 DA379 RS14550 WP 003151388.1 ....MAEGASEFSEVRDFFYGRHLLRSEIPFSDHCIKRFIEKEYITAEPSIIRRKNRYVCQRCGQSDQACFAAFWAPSAK 76
NTDB id 108 BSU 35470 NP 391427.1 MNVPVEKNSSFSKELQQTLRSRHLLRTELSFSDEMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHS..SG 78
consensus
NTDB id 285101 DA379 RS14550 WP 003151388.1 RQITYCRACVMMGRADELTLLYSWNQAPENRWKPVKLSWEGTLTDGQKQAAAALTDAIKERQELLVWAVCGSGKTEMLFP 156
NTDB id 108 BSU 35470 NP 391427.1 KNKLYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSSGQQKAANVLIEAISKKEELLIWAVCGAGKTEMLFP 158
consensus
NTDB id 285101 DA379 RS14550 WP 003151388.1 GIEFALNQGLRVCVATPRTDVVLELLPRLKKAFEKVDVSALYGGSEDKGRLTPLMISTAHQLMRYRDIFDVMIIDEVDAF 236
NTDB id 108 BSU 35470 NP 391427.1 GIESALNQGLRVCIATPRTDVVLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAF 238
consensus
NTDB id 285101 DA379 RS14550 WP 003151388.1 PFSADETLRFAVDKARKKNSALVYVTATPSDTLKKNAEAGLLKSVRIPARYHRKPLPEPRFLWCGNWKKKLQKGKLPRSV 316
NTDB id 108 BSU 35470 NP 391427.1 PYSADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRIPARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAV 318
consensus
NTDB id 285101 DA379 RS14550 WP 003151388.1 TDWVRQKLQSQLPVFLFVPSVHVLKKTTDYFQKLNIRAEGVHAEDTFRKDKVKRFRDGRLDLLVTTTILERGVTVPKVQT 396
NTDB id 108 BSU 35470 NP 391427.1 KRWIEFHVKEGRPVFLFVPSVSILEKAAACFKGVHCRTASVHAEDKHRKEKVQQFRDGQLDLLITTTILERGVTVPKVQT 398
consensus
NTDB id 285101 DA379 RS14550 WP 003151388.1 CVLGAEAPIFTESALVQIAGRTGRHYKHFSGDVVLFHFGITNGMKKAKKHIEHMNKLAQKSKLLD 461
NTDB id 108 BSU 35470 NP 391427.1 GVLGAESSIFTESALVQIAGRTGRHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD 463
consensus
X non conserved
X similar
X ≥ 50% conserved