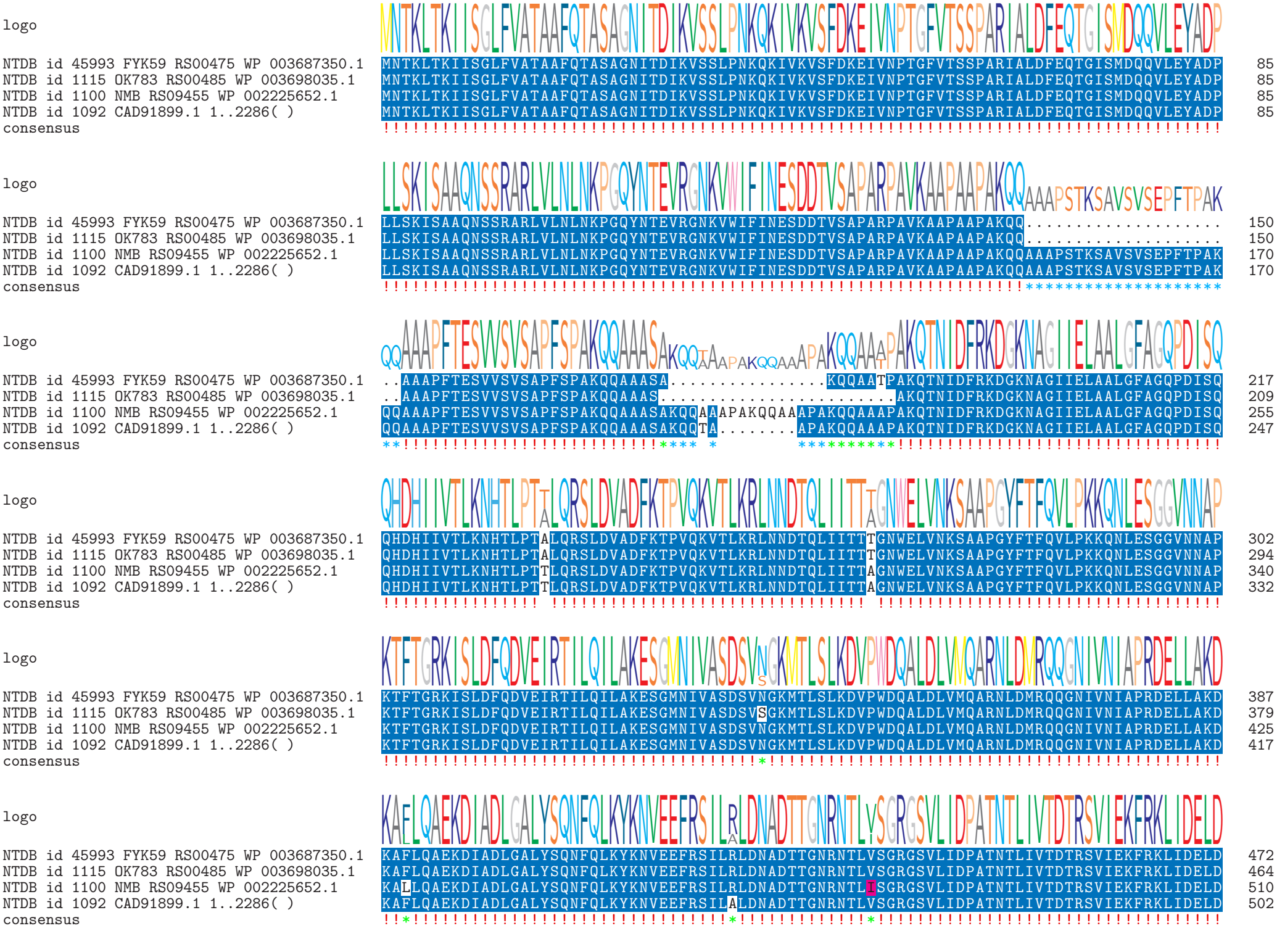
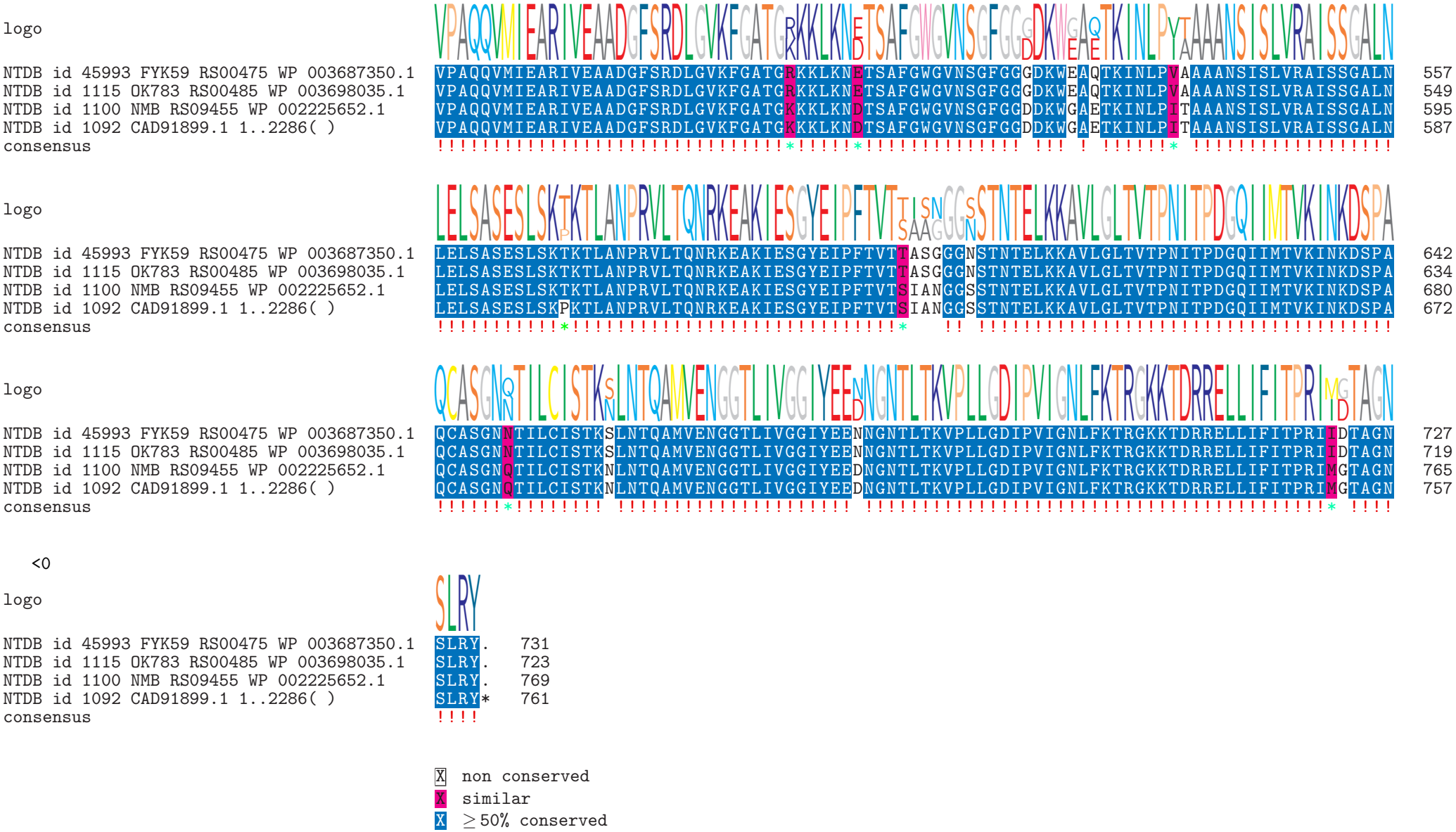

logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 MNTKLTKIISGLFVATAAFQTASAGNITDIKVSSLPNKQKIVKVSFDKEIVNPTGFVTSSPARIALDFEQTGISMDQQVLEYADP 85
NTDB id 1115 OK783 RS00485 WP 003698035.1 MNTKLTKIISGLFVATAAFQTASAGNITDIKVSSLPNKQKIVKVSFDKEIVNPTGFVTSSPARIALDFEQTGISMDQQVLEYADP 85
NTDB id 1100 NMB RS09455 WP 002225652.1 MNTKLTKIISGLFVATAAFQTASAGNITDIKVSSLPNKQKIVKVSFDKEIVNPTGFVTSSPARIALDFEQTGISMDQQVLEYADP 85
NTDB id 1092 CAD91899.1 1..2286( ) MNTKLTKIISGLFVATAAFQTASAGNITDIKVSSLPNKQKIVKVSFDKEIVNPTGFVTSSPARIALDFEQTGISMDQQVLEYADP 85
consensus
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 LLSKISAAQNSSRARLVLNLNKPGQYNTEVRGNKVWIFINESDDTVSAPARPAVKAAPAAPAKQQ..... 150
NTDB id 1115 OK783 RS00485 WP 003698035.1 LLSKISAAQNSSRARLVLNLNKPGQYNTEVRGNKVWIFINESDDTVSAPARPAVKAAPAAPAKQQ..... 150
NTDB id 1100 NMB RS09455 WP 002225652.1 LLSKISAAQNSSRARLVLNLNKPGQYNTEVRGNKVWIFINESDDTVSAPARPAVKAAPAAPAKQQAAAPSTKSAVSVSEPFTPAK 170
NTDB id 1092 CAD91899.1 1..2286( ) LLSKISAAQNSSRARLVLNLNKPGQYNTEVRGNKVWIFINESDDTVSAPARPAVKAAPAAPAKQQAAAPSTKSAVSVSEPFTPAK 170
consensus
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 ..AAAPFTESVVSVSAPFSPAKQQAAASA.....KQQAATPAKQTNIDFRKDGKNAGIIELAALGFAGQPDISQ 217
NTDB id 1115 OK783 RS00485 WP 003698035.1 ..AAAPFTESVVSVSAPFSPAKQQAAAS.....AKQTNIDFRKDGKNAGIIELAALGFAGQPDISQ 209
NTDB id 1100 NMB RS09455 WP 002225652.1 QQAAAPFTESVVSVSAPFSPAKQQAAASAKQQAAPAKQQAAAPAKQQAAAPAKQTNIDFRKDGKNAGIIELAALGFAGQPDISQ 255
NTDB id 1092 CAD91899.1 1..2286( ) QQAAAPFTESVVSVSAPFSPAKQQAAASAKQQTA.....APAKQQAAAPAKQTNIDFRKDGKNAGIIELAALGFAGQPDISQ 247
consensus
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 QHDHIIVTLKNHTLPTALQRSLDVADFKTPVQKVTLKRLNNDTQLIITTTGNWELVNKSAAPGYFTFQVLPKKQNLESGGVNNAP 302
NTDB id 1115 OK783 RS00485 WP 003698035.1 QHDHIIVTLKNHTLPTALQRSLDVADFKTPVQKVTLKRLNNDTQLIITTTGNWELVNKSAAPGYFTFQVLPKKQNLESGGVNNAP 294
NTDB id 1100 NMB RS09455 WP 002225652.1 QHDHIIVTLKNHTLPTTLQRSLDVADFKTPVQKVTLKRLNNDTQLIITTAGNWELVNKSAAPGYFTFQVLPKKQNLESGGVNNAP 340
NTDB id 1092 CAD91899.1 1..2286( ) QHDHIIVTLKNHTLPTTLQRSLDVADFKTPVQKVTLKRLNNDTQLIITTAGNWELVNKSAAPGYFTFQVLPKKQNLESGGVNNAP 332
consensus
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 KTFTGRKISLDFQDVEIRTILQILAKESGMNIVASDSVNGKMTLSLKDVPWDQALDLVMQARNLDMRQQGNIVNIAPRDELLAKD 387
NTDB id 1115 OK783 RS00485 WP 003698035.1 KTFTGRKISLDFQDVEIRTILQILAKESGMNIVASDSVSGKMTLSLKDVPWDQALDLVMQARNLDMRQQGNIVNIAPRDELLAKD 379
NTDB id 1100 NMB RS09455 WP 002225652.1 KTFTGRKISLDFQDVEIRTILQILAKESGMNIVASDSVNGKMTLSLKDVPWDQALDLVMQARNLDMRQQGNIVNIAPRDELLAKD 425
NTDB id 1092 CAD91899.1 1..2286( ) KTFTGRKISLDFQDVEIRTILQILAKESGMNIVASDSVNGKMTLSLKDVPWDQALDLVMQARNLDMRQQGNIVNIAPRDELLAKD 417
consensus
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 KAFLQAEKDIADLGALYSQNFQLKYKNVEEFRSILRLDNADTTGNRNTLVSGRGSVLIDPATNTLIVTDTRSVIEKFRKLIDELD 472
NTDB id 1115 OK783 RS00485 WP 003698035.1 KAFLQAEKDIADLGALYSQNFQLKYKNVEEFRSILRLDNADTTGNRNTLVSGRGSVLIDPATNTLIVTDTRSVIEKFRKLIDELD 464
NTDB id 1100 NMB RS09455 WP 002225652.1 KALLQAEKDIADLGALYSQNFQLKYKNVEEFRSILRLDNADTTGNRNTLISGRGSVLIDPATNTLIVTDTRSVIEKFRKLIDELD 510
NTDB id 1092 CAD91899.1 1..2286( ) KAFLQAEKDIADLGALYSQNFQLKYKNVEEFRSILALDNADTTGNRNTLVSGRGSVLIDPATNTLIVTDTRSVIEKFRKLIDELD 502
consensus


logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 VPAQQVMIEARIVEAADGFSRDLGVKFGATGRKKLKNETSAFGWGVNSGFGGGDKWEAQTKINLPVAAAANSISLVRAISSGALN 557
NTDB id 1115 OK783 RS00485 WP 003698035.1 VPAQQVMIEARIVEAADGFSRDLGVKFGATGRKKLKNETSAFGWGVNSGFGGGDKWEAQTKINLPVAAAANSISLVRAISSGALN 549
NTDB id 1100 NMB RS09455 WP 002225652.1 VPAQQVMIEARIVEAADGFSRDLGVKFGATGKKKLKNDTSAFGWGVNSGFGGDDKWGAETKINLPITAAANSISLVRAISSGALN 595
NTDB id 1092 CAD91899.1 1..2286( ) VPAQQVMIEARIVEAADGFSRDLGVKFGATGKKKLKNDTSAFGWGVNSGFGGDDKWGAETKINLPITAAANSISLVRAISSGALN 587
consensus
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 LELSASESLSKTKTLANPRVLTQNRKEAKIESGYEIPFTVTTASGGGNSTNTELKKAVLGLTVTPNITPDGQIIMTVKINKDSPA 642
NTDB id 1115 OK783 RS00485 WP 003698035.1 LELSASESLSKTKTLANPRVLTQNRKEAKIESGYEIPFTVTTASGGGNSTNTELKKAVLGLTVTPNITPDGQIIMTVKINKDSPA 634
NTDB id 1100 NMB RS09455 WP 002225652.1 LELSASESLSKTKTLANPRVLTQNRKEAKIESGYEIPFTVTSIANGGSSTNTELKKAVLGLTVTPNITPDGQIIMTVKINKDSPA 680
NTDB id 1092 CAD91899.1 1..2286( ) LELSASESLSKPKTLANPRVLTQNRKEAKIESGYEIPFTVTSIANGGSSTNTELKKAVLGLTVTPNITPDGQIIMTVKINKDSPA 672
consensus
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 QCASGNNTILCISTKSLNTQAMVENGGTLIVGGIYEENNGNTLTKVPLLGDIPVIGNLFKTRGKKTDRRELLIFITPRIIDTAGN 727
NTDB id 1115 OK783 RS00485 WP 003698035.1 QCASGNNTILCISTKSLNTQAMVENGGTLIVGGIYEENNGNTLTKVPLLGDIPVIGNLFKTRGKKTDRRELLIFITPRIIDTAGN 719
NTDB id 1100 NMB RS09455 WP 002225652.1 QCASGNQTILCISTKNLNTQAMVENGGTLIVGGIYEEDNGNTLTKVPLLGDIPVIGNLFKTRGKKTDRRELLIFITPRIMGTAGN 765
NTDB id 1092 CAD91899.1 1..2286( ) QCASGNQTILCISTKNLNTQAMVENGGTLIVGGIYEEDNGNTLTKVPLLGDIPVIGNLFKTRGKKTDRRELLIFITPRIMGTAGN 757
consensus
<0
logo
NTDB id 45993 FYK59 RS00475 WP 003687350.1 SLRY. 731
NTDB id 1115 OK783 RS00485 WP 003698035.1 SLRY. 723
NTDB id 1100 NMB RS09455 WP 002225652.1 SLRY. 769
NTDB id 1092 CAD91899.1 1..2286( ) SLRY* 761
consensus
X non conserved
X similar
X ≥ 50% conserved