logo
NTDB id 1252 GC085 RS07730 WP 011213805.1 MDKNSPPLLTFHYQGINKAGQKMEGDIQARSLAIAKADLRKQGIVTNKVVKKRKPLFDR.....KNKKITQADITV 71
NTDB id 329715 EHF33 RS02885 WP 124867717.1 .....MPVFEYRVRDRSGKILKSQTEADTIAQVRDSLRAKGLLIVDIKSPKGGMQTDLNDLKIPFLSDRPPGLKPIAI 73
NTDB id 1029 TT RS02230 WP 011228203.1 .....MPVYQYKARDRQGRLVEATIEAEDLRTAARLLRDRGLFVAEIKEPGKGLQAEVR...IPA.LERGPGLKDLAI 69
consensus
logo
NTDB id 1252 GC085 RS07730 WP 011213805.1 FSRQLATMIESGIPLVQAFDIVAKGQSNKRLKDLIETIKHDIETGLTLAESLIKHPLYFNELFCNLVDAGEKSGSLDIML 151
NTDB id 329715 EHF33 RS02885 WP 124867717.1 FSKQLATLINAGVPLVQSLTILQRQIEHKGLQNVVRKVRNDVEAGTPLSEAIAKFPKVFNRLYINLVRAGETSGTLDSVL 153
NTDB id 1029 TT RS02230 WP 011228203.1 FSRQLATMLGAGLTLLQALAILERQTENRKFREILKQVRTDVEGGMAFSEALSK.HKIFSRLYVNLVRAGETSGGLDLIL 148
consensus
logo
NTDB id 1252 GC085 RS07730 WP 011213805.1 DKVATYKEKIEIIKKKIKKALTYPIAVMVVALLVTAGLLIYVVPQFESLFKGFGADLPAMTRAVITMSEFMQAYWYIIFG 231
NTDB id 329715 EHF33 RS02885 WP 124867717.1 ERIAEFQEKDLATRGKIKSAMTYPVIVLVFALLITYFLLTTIVPQFGGMLTQLGAELPLLTRVLMAMSDFLKTKTIVIVL 233
NTDB id 1029 TT RS02230 WP 011228203.1 DRLASFLEKELELRGKIRSAMTYPVIVFVFAVGVAYFLLTGIVPQFAQILTDLGSELPLLTRFLIAVSDLLRAATLPLLL 228
consensus
logo
NTDB id 1252 GC085 RS07730 WP 011213805.1 ALGGV..VYSFFYAKNHSLEFAQTIDRVMLKLPVIGPILEKAAIARFARTLSITFAAGLPLVEALKSVAGATGNIIYAKA 309
NTDB id 329715 EHF33 RS02885 WP 124867717.1 IVAALVFAYRYYYSTKK...GRRVVDDFKLKIPVFGTLIQRSAIASFARTFGLLLINSGVNIIESLEITKGTANNAIVEDT 310
NTDB id 1029 TT RS02230 WP 011228203.1 LAVALFFAYRWYYGTPQ...GRRVIDRLKLRLPVFGNLNRKTAVARFSRTLALLLSSGVNIVEALDITKGTAGNSVVEEI 305
consensus
logo
NTDB id 1252 GC085 RS07730 WP 011213805.1 TDKIREEVATGQQMFIAMEN.THLFPNMVIQMVAIGEESGALEKMLSKVADFYEEEVNNAVDALSSLLEPIIMSILGILV 388
NTDB id 329715 EHF33 RS02885 WP 124867717.1 IENAKNVVMVGEQMSGSLAASP.VFPPMVVSMVAIGEETGALDTMLAKVADFYDREVEEAIEGLTAAIEPIMIVFLGAIV 389
NTDB id 1029 TT RS02230 WP 011228203.1 VEAAKLKIQQGDPLNLTLAQHPFVFPPMVSSMVAIGEETGALDTMLSKVADFYEREVDEAVASLTAAIEPLMIIFLGVIV 385
consensus
logo
NTDB id 1252 GC085 RS07730 WP 011213805.1 GGLVVAMYLPIFKLGSAV... 406
NTDB id 329715 EHF33 RS02885 WP 124867717.1 GLIVAGMFLPMFSIISALSK. 409
NTDB id 1029 TT RS02230 WP 011228203.1 GMIVAGMFLPLFKIIGTLSVQ 406
consensus
X non conserved
X similar
X ≥ 50% conserved