logo
NTDB id 356537 E5Q06 RS09400 WP 010888498.1 MPVYEYRARDRGGQTLKSQMEADSEAQVRDALRARNLMILEIKAPKTGLNADIKIPGLTDRPPGLKQVAVFSKQLATLIN 80
NTDB id 1029 TT RS02230 WP 011228203.1 MPVYQYKARDRQGRLVEATIEAEDLRTAARLLRDRGLFVAEIKEPGKGLQAEVRIPALE.RGPGLKDLAIFSRQLATMLG 79
consensus
logo
NTDB id 356537 E5Q06 RS09400 WP 010888498.1 AGVPLVQSLAILQRQLDNKTFEAVIKKVRGEVESGQPLSEALQQHPKIFGRLFINLVRAGETSGTLDTVLERIADFQEKQ 160
NTDB id 1029 TT RS02230 WP 011228203.1 AGLTLLQALAILERQTENRKFREILKQVRTDVEGGMAFSEALSKH.KIFSRLYVNLVRAGETSGGLDLILDRLASFLEKE 158
consensus
logo
NTDB id 356537 E5Q06 RS09400 WP 010888498.1 LALQGKIKSALTYPTVVLVFALGITYFLLTTVVPQFAGILTQLNAPLPFITRMLMAVSNFLQHSGLLLLAIIAVIVLAYR 240
NTDB id 1029 TT RS02230 WP 011228203.1 LELRGKIRSAMTYPVIVFVFAVGVAYFLLTGIVPQFAQILTDLGSELPLLTRFLIAVSDLLRAATLPLLLLAVALFFAYR 238
consensus
logo
NTDB id 356537 E5Q06 RS09400 WP 010888498.1 WYYRTPQGRHAIDDFKLRLPVFGNLIQKSAISSFARTFGLLVSSGVNIIESLEITKGTADNAIVEETIENAKNVVISGDQ 320
NTDB id 1029 TT RS02230 WP 011228203.1 WYYGTPQGRRVIDRLKLRLPVFGNLNRKTAVARFSRTLALLLSSGVNIVEALDITKGTAGNSVVEEIVEAAKLKIQQGDP 318
consensus
logo
NTDB id 356537 E5Q06 RS09400 WP 010888498.1 MSSSLAT.SKVFPPMVVSMVAIGEETGSLDNMLNKVADFYDREVDEAVESLTAAIEPMMIVFLGGIVGLIVAGMFLPMFS 399
NTDB id 1029 TT RS02230 WP 011228203.1 LNLTLAQHPFVFPPMVSSMVAIGEETGALDTMLSKVADFYEREVDEAVASLTAAIEPLMIIFLGVIVGMIVAGMFLPLFK 398
consensus
logo
NTDB id 356537 E5Q06 RS09400 WP 010888498.1 IIGTLSQ. 406
NTDB id 1029 TT RS02230 WP 011228203.1 IIGTLSVQ 406
consensus
X non conserved
X similar
X ≥ 50% conserved