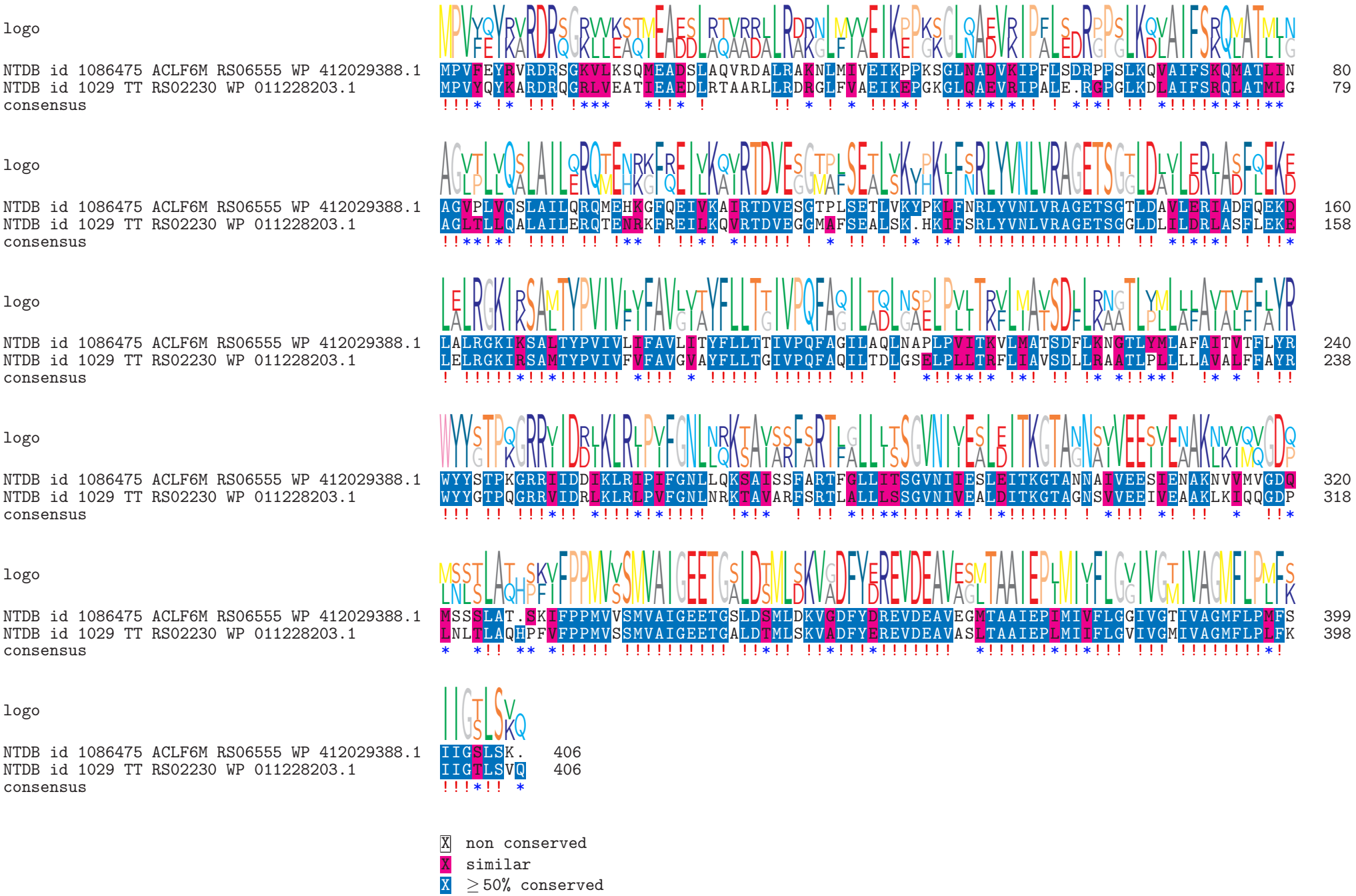
logo
NTDB id 1086475 ACLF6M RS06555 WP 412029388.1 MPVFEYRVRDRSGKVLKSQMEADSLAQVRDALRAKNLMIVEIKPPKSGLNADVKIPFLSDRPPSLKQVAIFSKQMATLIN 80
NTDB id 1029 TT RS02230 WP 011228203.1 MPVYQYKARDRQGRLVEATIEAEDLRTAARLLRDRGLFVAEIKEPGKGLQAEVRIPALE.RGPGLKDLAIFSRQLATMLG 79
consensus
logo
NTDB id 1086475 ACLF6M RS06555 WP 412029388.1 AGVPLVQSLAILQRQMEHKGFQEIVKAIRTDVESGTPLSETLVKYPKLFNRLYVNLVRAGETSGTLDAVLERIADFQEKD 160
NTDB id 1029 TT RS02230 WP 011228203.1 AGLTLLQALAILERQTENRKFREILKQVRTDVEGGMAFSEALSK.HKIFSRLYVNLVRAGETSGGLDLILDRLASFLEKE 158
consensus
logo
NTDB id 1086475 ACLF6M RS06555 WP 412029388.1 LALRGKIKSALTYPVIVLIFAVLITYFLLTTIVPQFAGILAQLNAPLPVITKVLMATSDFLKNGTLYMLAFAITVTFLYR 240
NTDB id 1029 TT RS02230 WP 011228203.1 LELRGKIRSAMTYPVIVFVFAVGVAYFLLTGIVPQFAQILTDLGSELPLLTRFLIAVSDLLRAATLPLLLLAVALFFAYR 238
consensus
logo
NTDB id 1086475 ACLF6M RS06555 WP 412029388.1 WYYSTPKGRRIIDDIKLRIPIFGNLLQKSAISSFARTFGLLITSGVNIIESLEITKGTANNAIVEESIENAKNVVMVGDQ 320
NTDB id 1029 TT RS02230 WP 011228203.1 WYYGTPQGRRVIDRLKLRLPVFGNLNRKTAVARFSRTLALLLSSGVNIVEALDITKGTAGNSVVEEIVEAAKLKIQQGDP 318
consensus
logo
NTDB id 1086475 ACLF6M RS06555 WP 412029388.1 MSSSLAT.SKIFPPMVVSMVAIGEETGSLDSMLDKVCDFYDREVDEAVEGMTAAIEPIMIVFLGGIVGTIVAGMFLPMFS 399
NTDB id 1029 TT RS02230 WP 011228203.1 LNLTLAQHPFVFPPMVSSMVAIGEETGALDTMLSKVADFYEREVDEAVASLTAAIEPLMIIFLGVIVGMIVAGMFLPLFK 398
consensus
logo
NTDB id 1086475 ACLF6M RS06555 WP 412029388.1 IIGSLSK. 406
NTDB id 1029 TT RS02230 WP 011228203.1 IIGTLSVQ 406
consensus
X non conserved
X similar
X ≥ 50% conserved