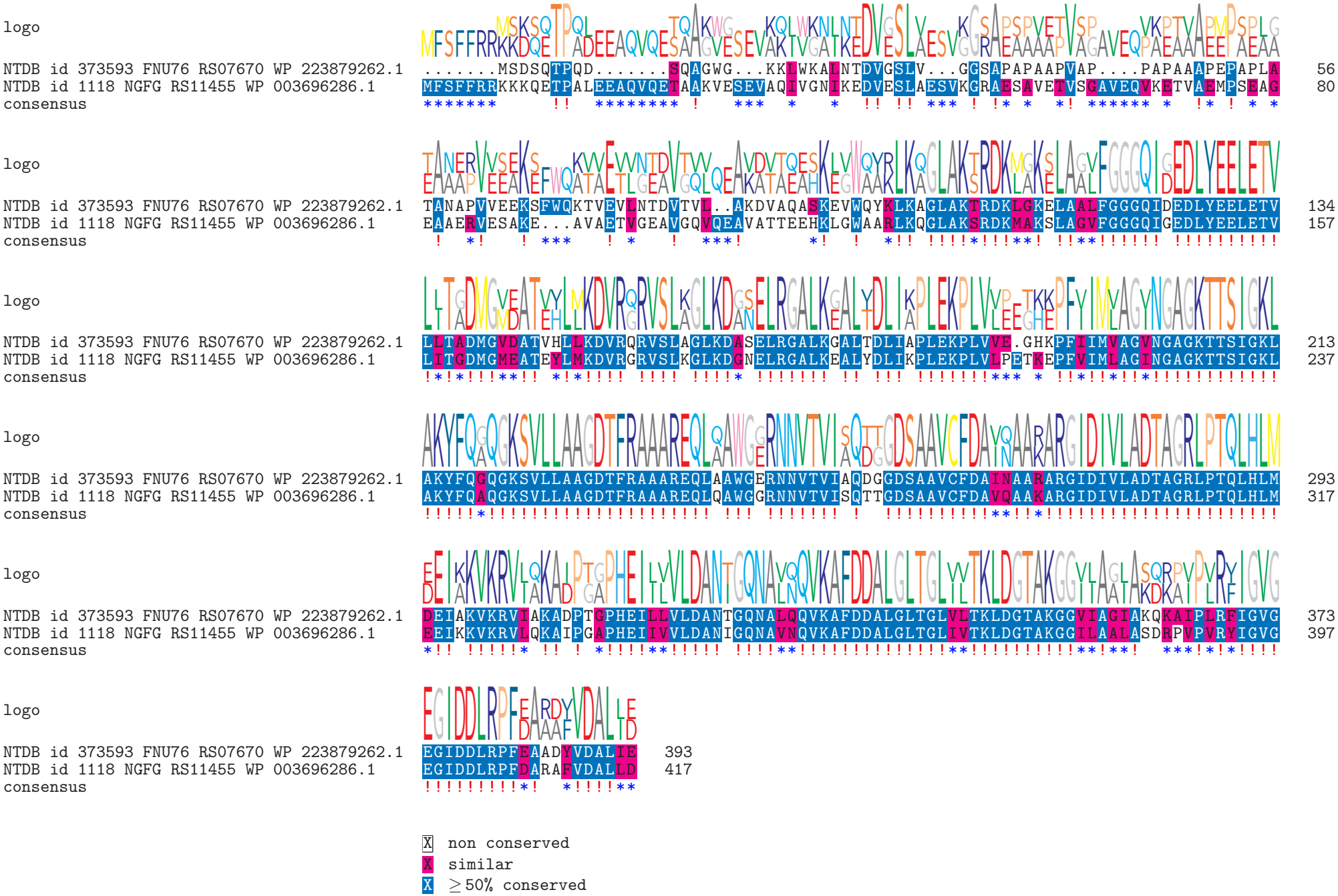
logo
NTDB id 373593 FNU76 RS07670 WP 223879262.1 .....MSDSQTPQD.....SQAGWG...KKLWKALNTDVGSLV...GGSAPAPAAPVAP....PAPAAAPEPAPLA 56
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus ******* !! ******** ! *** * * !! !! *** ! !* * *! ****** * !* ! * *
logo
NTDB id 373593 FNU76 RS07670 WP 223879262.1 TANAPVVEEKSFWQKTVEVLNTDVTVL..AKDVAQASKEVWQYKLKAGLAKTRDKLGKELAALFGGGQIDEDLYEELETV 134
NTDB id 1118 NGFG RS11455 WP 003696286.1 EAAERVESAKE...AVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETV 157
consensus ! *! ! *** ! * ! ***! *! ! *!! !!!!*!!!**! !!**!!!!!! !!!!!!!!!!
logo
NTDB id 373593 FNU76 RS07670 WP 223879262.1 LLTADMGVDATVHLLKDVRQRVSLAGLKDASELRGALKGALTDLIAPLEKPLVVE.GHKPFIIMVAGVNGAGKTTSIGKL 213
NTDB id 1118 NGFG RS11455 WP 003696286.1 LITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKL 237
consensus !*!*!!!**!! *!*!!!! !!!! !!!!* !!!!!!! !! !!! !!!!!!!*** * !!*!!*!!*!!!!!!!!!!!!
logo
NTDB id 373593 FNU76 RS07670 WP 223879262.1 AKYFQGQGKSVLLAAGDTFRAAAREQLAAWGERNNVTVIAQDGGDSAAVCFDAINAARARGIDIVLADTAGRLPTQLHLM 293
NTDB id 1118 NGFG RS11455 WP 003696286.1 AKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLM 317
consensus !!!!!*!!!!!!!!!!!!!!!!!!!!! !!! !!!!!!! ! !!!!!!!!!!**!!*!!!!!!!!!!!!!!!!!!!!
logo
NTDB id 373593 FNU76 RS07670 WP 223879262.1 DEIAKVKRVIAKADPTGPHEILLVLDANTGQNALQQVKAFDDALGLTGLVLTKLDGTAKGGVIAGIAKQKAIPLRFIGVG 373
NTDB id 1118 NGFG RS11455 WP 003696286.1 EEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVG 397
consensus *!! !!!!!* !! ! *!!!!**!!!!! !!!!**!!!!!!!!!!!!!!**!!!!!!!!!!**!**! ***!*!*!!!!
logo
NTDB id 373593 FNU76 RS07670 WP 223879262.1 EGIDDLRPFEAADYVDALIE 393
NTDB id 1118 NGFG RS11455 WP 003696286.1 EGIDDLRPFDARAFVDALLD 417
consensus !!!!!!!!!*! *!!!!**
X non conserved
X similar
X ≥ 50% conserved