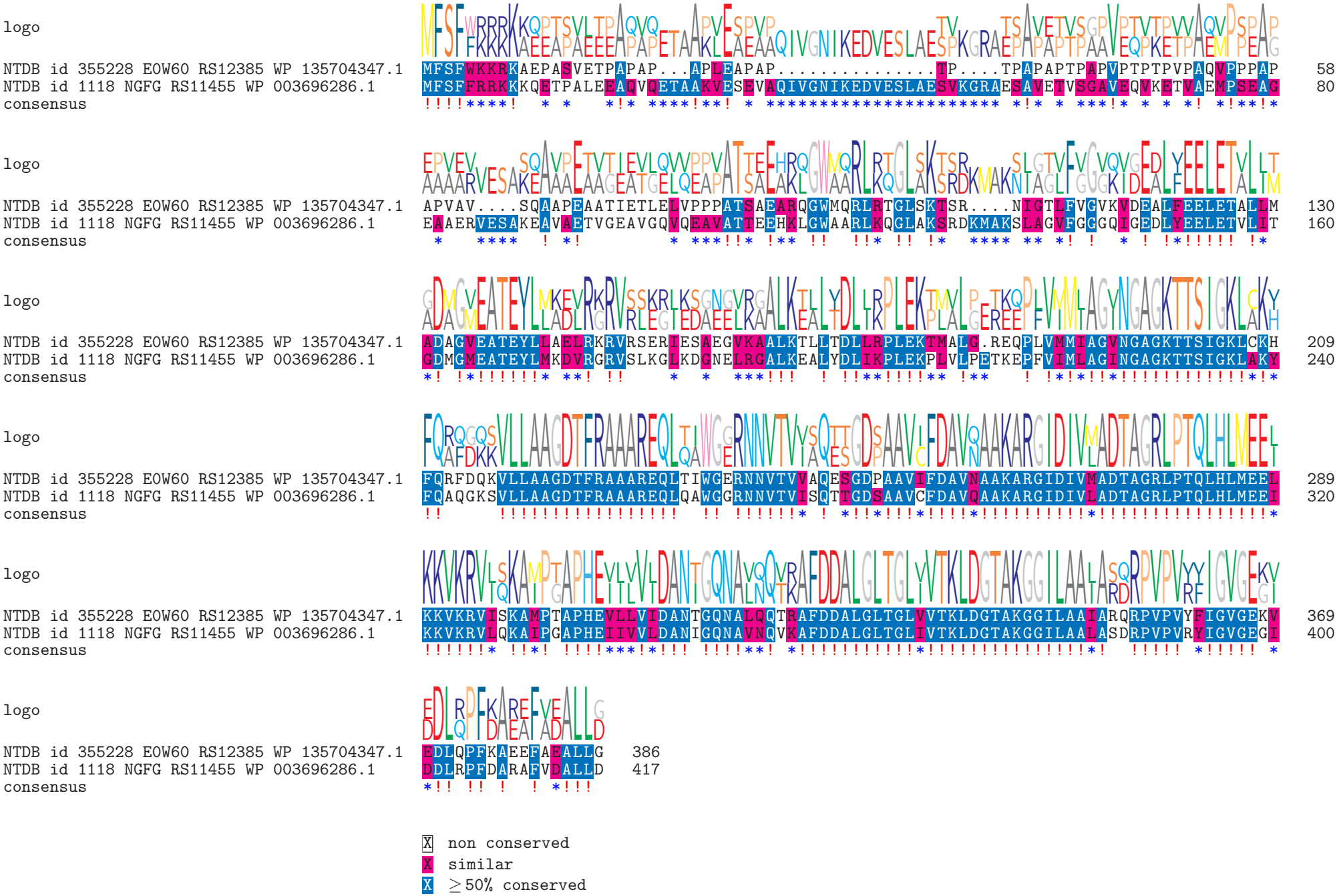
logo
NTDB id 355228 E0W60 RS12385 WP 135704347.1 MFSFWKKRKAEPASVETPAPAP...APLEAPAP.....TP....TPAPAPTPAPVPTPTPVPAQVPPPAP 58
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus
logo
NTDB id 355228 E0W60 RS12385 WP 135704347.1 APVAV....SQAAPEAATIETLELVPPPATSAEARQGWMQRLRTGLSKTSR....NICTLFVGVKVDEALREELETALLM 130
NTDB id 1118 NGFG RS11455 WP 003696286.1 EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT 160
consensus
logo
NTDB id 355228 E0W60 RS12385 WP 135704347.1 ADAGVEATEYLLAELRKRVRSERIESAEGVKAALKTLLTDLLRPLEKTMALG.REQPLVMMIAGVNGAGKTTSIGKLCKH 209
NTDB id 1118 NGFG RS11455 WP 003696286.1 GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY 240
consensus
logo
NTDB id 355228 E0W60 RS12385 WP 135704347.1 FQRFDQKVLLAAGDTFRAAAREQLTIWGERNNVTVVAQESGDPAAVIFDAVNAAKARGIDIVMADTAGRLPTQLHLMEEL 289
NTDB id 1118 NGFG RS11455 WP 003696286.1 FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI 320
consensus
logo
NTDB id 355228 E0W60 RS12385 WP 135704347.1 KKVKRVISKAMPTAPHEVLLVIDANTGQNALQQTRAFDDALGLTGLVVTKLDGTAKGGILAAIARQRPVPVYFIGVGEKV 369
NTDB id 1118 NGFG RS11455 WP 003696286.1 KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI 400
consensus
logo
NTDB id 355228 E0W60 RS12385 WP 135704347.1 EDLQPFKAEEFAEALLG 386
NTDB id 1118 NGFG RS11455 WP 003696286.1 DDLRPFDARAFVDALLD 417
consensus
X non conserved
X similar
X ≥ 50% conserved