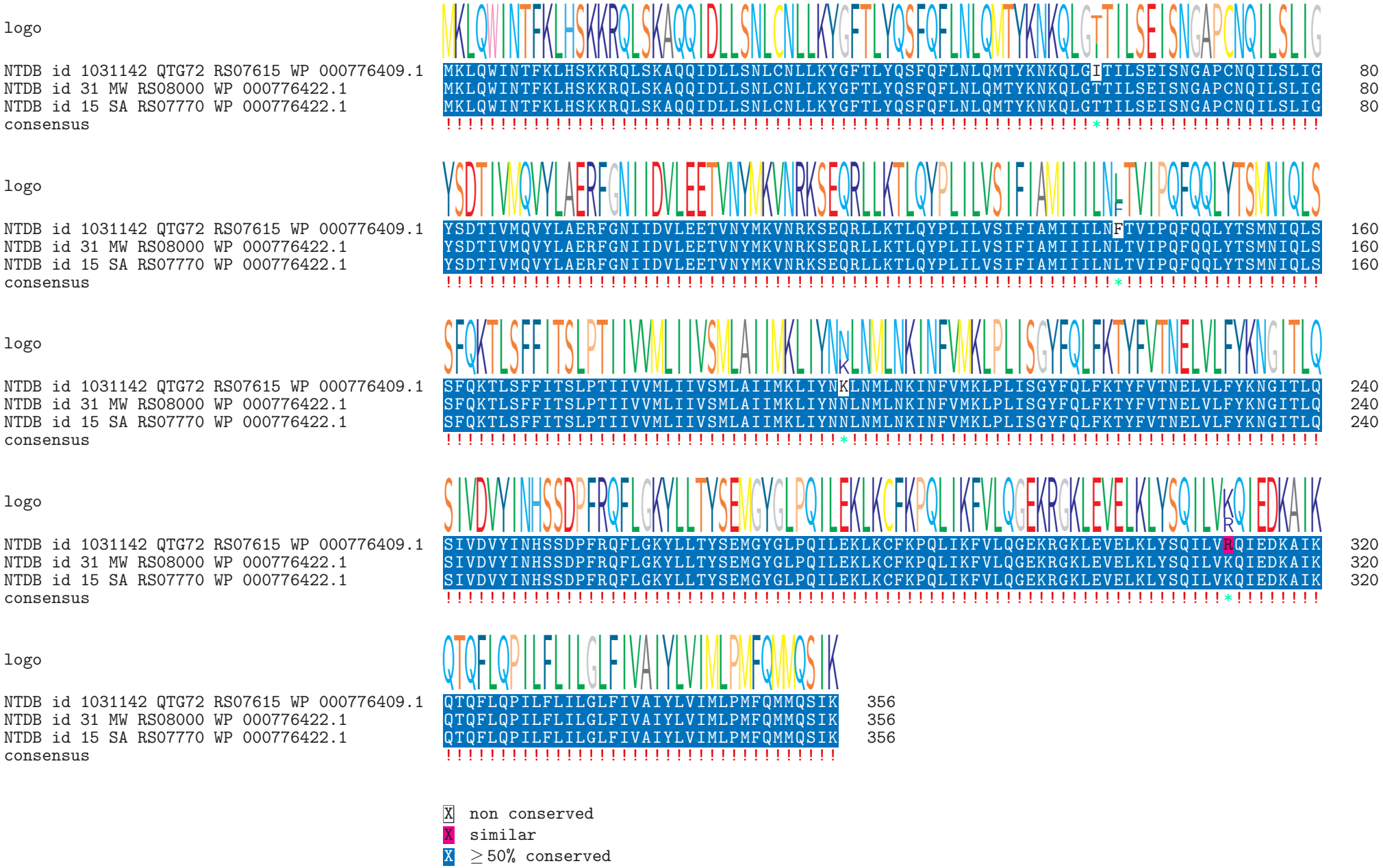
logo
NTDB id 1031142 QTG72 RS07615 WP 000776409.1 MKLQWINTFKLHSKKRQLSKAQQIDLLSNLCNLLKYGFTLYQSFQFLNLQMTYKNKQLGITILSEISNGAPCNQILSLIG 80
NTDB id 31 MW RS08000 WP 000776422.1 MKLQWINTFKLHSKKRQLSKAQQIDLLSNLCNLLKYGFTLYQSFQFLNLQMTYKNKQLGTTILSEISNGAPCNQILSLIG 80
NTDB id 15 SA RS07770 WP 000776422.1 MKLQWINTFKLHSKKRQLSKAQQIDLLSNLCNLLKYGFTLYQSFQFLNLQMTYKNKQLGTTILSEISNGAPCNQILSLIG 80
consensus
logo
NTDB id 1031142 QTG72 RS07615 WP 000776409.1 YSDTIVMQVYLAERFGNIIDVLEETVNYMKVNRKSEQRLLKTLQYPLILVSIFIAMIIILNFTVIPQFQQLYTSMNIQLS 160
NTDB id 31 MW RS08000 WP 000776422.1 YSDTIVMQVYLAERFGNIIDVLEETVNYMKVNRKSEQRLLKTLQYPLILVSIFIAMIIILNLTVIPQFQQLYTSMNIQLS 160
NTDB id 15 SA RS07770 WP 000776422.1 YSDTIVMQVYLAERFGNIIDVLEETVNYMKVNRKSEQRLLKTLQYPLILVSIFIAMIIILNLTVIPQFQQLYTSMNIQLS 160
consensus
logo
NTDB id 1031142 QTG72 RS07615 WP 000776409.1 SFQKTLSFFITSLPTIIVVMLIIVSMLAIIMKLIYNKLNMLNKINFVMKLPLISGYFQLFKTYFVTNELVLFYKNGITLQ 240
NTDB id 31 MW RS08000 WP 000776422.1 SFQKTLSFFITSLPTIIVVMLIIVSMLAIIMKLIYNNLNMLNKINFVMKLPLISGYFQLFKTYFVTNELVLFYKNGITLQ 240
NTDB id 15 SA RS07770 WP 000776422.1 SFQKTLSFFITSLPTIIVVMLIIVSMLAIIMKLIYNNLNMLNKINFVMKLPLISGYFQLFKTYFVTNELVLFYKNGITLQ 240
consensus
logo
NTDB id 1031142 QTG72 RS07615 WP 000776409.1 SIVDVYINHSSDPFRQFLGKYLLTYSEMGYGLPQILEKLKCFKPQLIKFVLQGEKRGKLEVELKLYSQILVRQIEDKAIK 320
NTDB id 31 MW RS08000 WP 000776422.1 SIVDVYINHSSDPFRQFLGKYLLTYSEMGYGLPQILEKLKCFKPQLIKFVLQGEKRGKLEVELKLYSQILVKQIEDKAIK 320
NTDB id 15 SA RS07770 WP 000776422.1 SIVDVYINHSSDPFRQFLGKYLLTYSEMGYGLPQILEKLKCFKPQLIKFVLQGEKRGKLEVELKLYSQILVKQIEDKAIK 320
consensus
logo
NTDB id 1031142 QTG72 RS07615 WP 000776409.1 QTQFLQPILFLILGLFIVAIYLVIMLPMFQMMQSIK 356
NTDB id 31 MW RS08000 WP 000776422.1 QTQFLQPILFLILGLFIVAIYLVIMLPMFQMMQSIK 356
NTDB id 15 SA RS07770 WP 000776422.1 QTQFLQPILFLILGLFIVAIYLVIMLPMFQMMQSIK 356
consensus
X non conserved
X similar
X ≥ 50% conserved