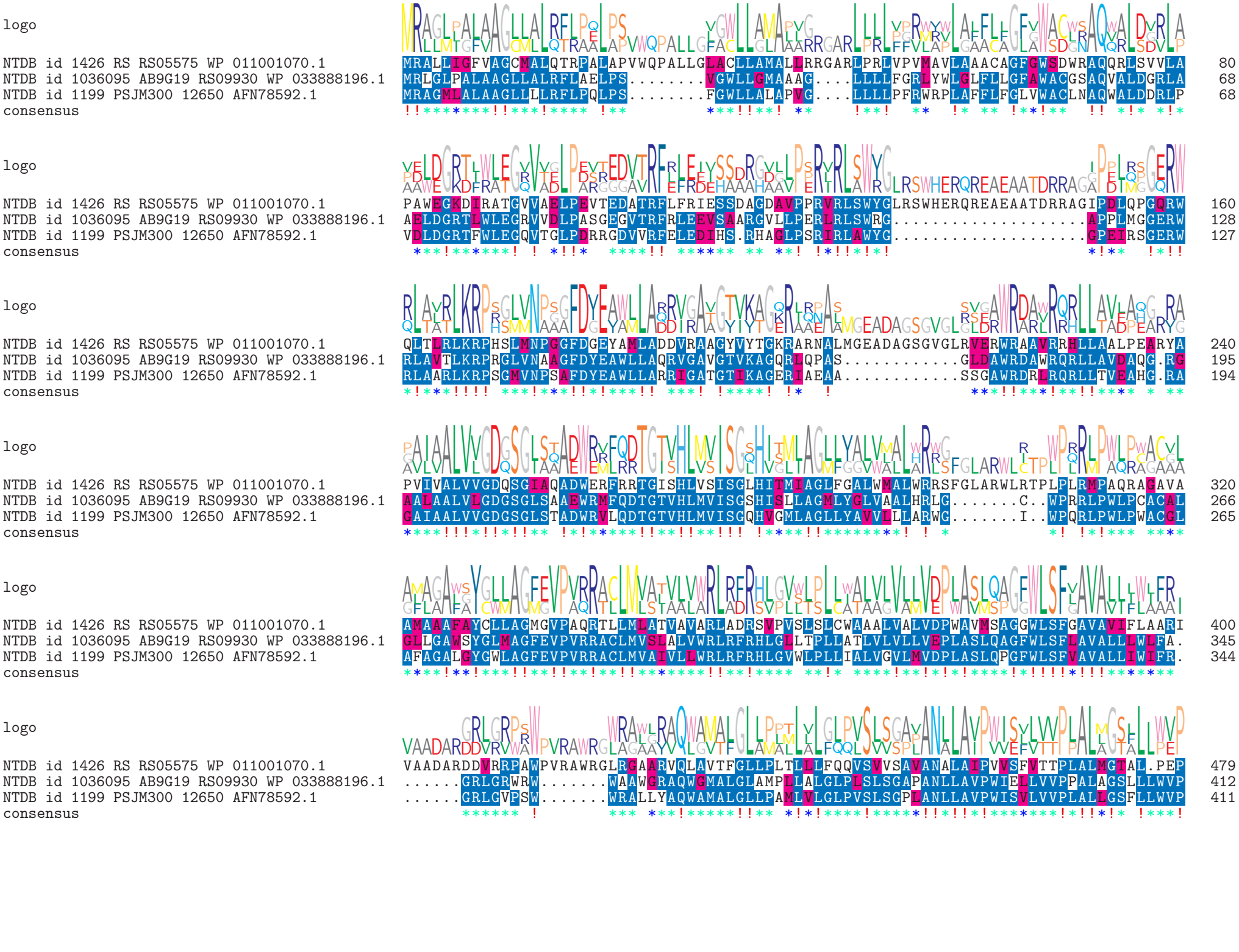
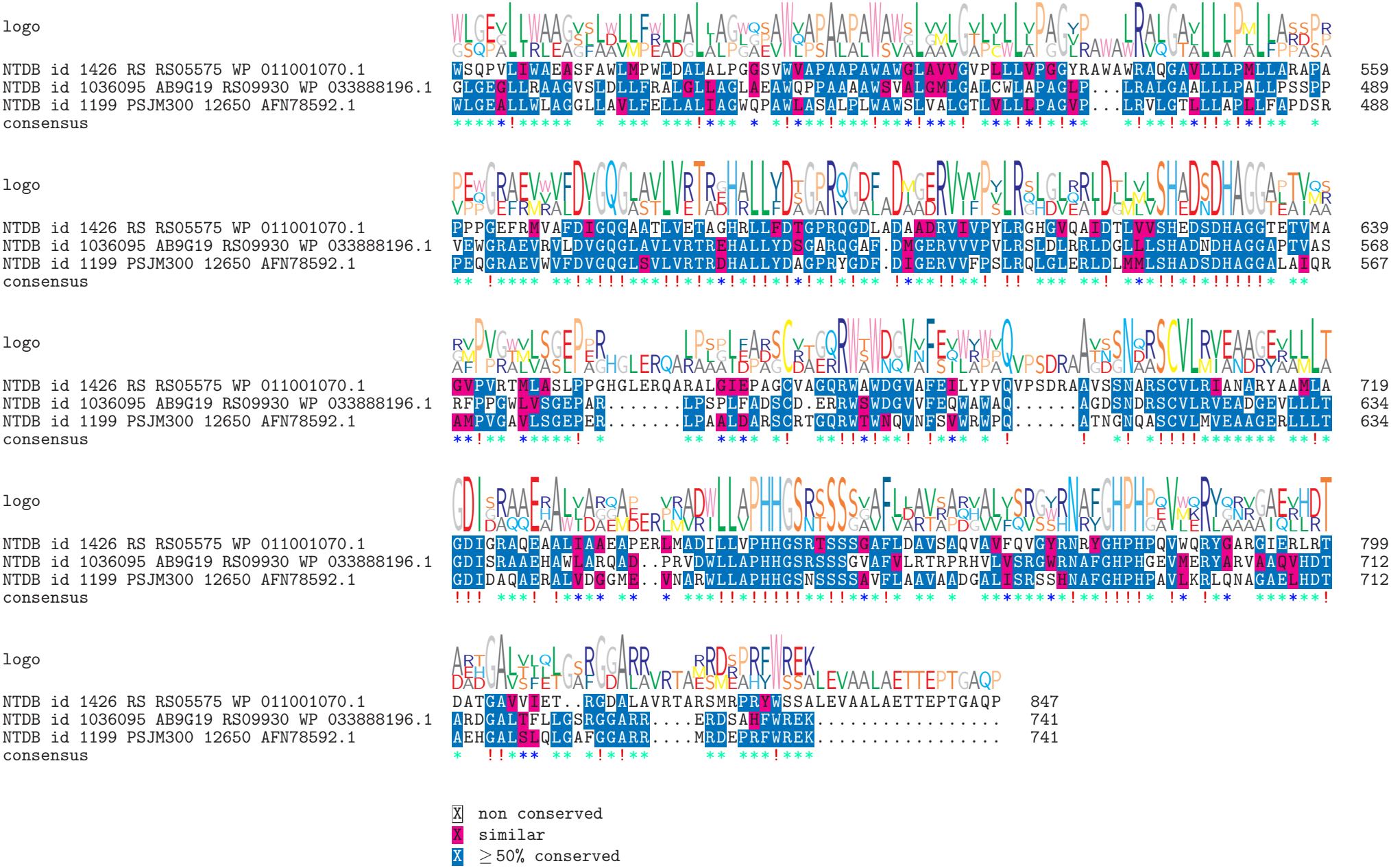

logo
NTDB id 1426 RS RS05575 WP 011001070.1 MRALLIGFVAGCMALQTRPALAPVWQPALLGLACLLAMALLRRGARLPRLVPVMAVLAAACAGFGWSDWRAQQRLSVVLA 80
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 MRLGLPALAAGLLALRFLAELPS.....VGWLLGMAAAG....LLLLFGRLYWLGLFLLGFAWACGSAQVALDGRLA 68
NTDB id 1199 PSJM300 12650 AFN78592.1 MRAGMLALAAGLLLLRFLPQLPS.....FGWLLALAPVG....LLLLPFRWRPLAFFLFGLVWACLNAQWALDDRLP 68
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 PAWEGKDIRATGVVAELPEVTEDATRFLFRIESSDAGDAVPPRVRLSWYGLRSWHERQREAEAATDRRAGIPDLQPGQRW 160
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 AELDGRTLWLEGRVVDLPASGEGVTRFRLEEVSAARGVLLPERLRLSWRG.....APPLMGGERW 128
NTDB id 1199 PSJM300 12650 AFN78592.1 VDLDGRTFWLEGQVTGLPDRRGDVVRFELEDIHS.RHAGLPSRIRLAWYG.....GPEIRSGERW 127
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 QLTLRLKRPHSLMNPGGFDGEYAMLADDVRAAGYVYTGKRARNALMGEADAGSGVGLRVERWRAAVRRHLLAALPEARYA 240
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 RLAVTLKRPRGLVNAAGFDYEAWLLAQRVGAVGTVKAGQRLQPAS.....GLDAWRDAWRQRLLAVDAQG.RG 195
NTDB id 1199 PSJM300 12650 AFN78592.1 RLAARLKRPSGMVNPSAFDYEAWLLARRIGATGTIKAGERIAEAA.....SSGAWRDRLRQRLLTVEAHG.RA 194
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 PVIVALVVGDQSGIAQADWERFRRTGISHLVSISGLHITMIAGLFGALWMALWRRSFGLARWLRTPLPLRMPAQRAGAVA 320
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 AALAALVLGDGSGLSAAEWRMFQDTGTVHLMVISGSHISLLAGMLYGLVAALHRLG.....C..WPRRLPWLPCACAL 266
NTDB id 1199 PSJM300 12650 AFN78592.1 GAIAALVVGDGSGLSTADWRVLQDTGTVHLMVISGQHVGMLAGLLYAVVLLLARWG.....I..WPQRLPWLPWACGL 265
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 AMAAAFAYCLLAGMGVPAQRTLLMLATVAVARLADRSVPVSLSLCWAAALVALVDPWAVMSAGGWLSFGAVAVIFLAARI 400
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 GLLGAWSYGLMAGFEVPVRRACLMVSLALVWRLRFRHLGLLTPLLATLVLVLLVEPLASLQAGFWLSFLAVALLLWLFA. 345
NTDB id 1199 PSJM300 12650 AFN78592.1 AFAGALGYGWLAGFEVPVRRACLMVAIVLLWRLRFRHLGVWLPLLIALVGVLMVDPLASLQPGFWLSFVAVALLIWIFR. 344
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 VAADARDDVRRPAWPVRAWRGLRGAARVQLAVTFGLLPLTLLLFQQVSVVSAVANALAIPVVSFVTTPLALMGTAL.PEP 479
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 ......GRLGRWRW.....WAAWGRAQWGMALGLAMPLLALGLPLSLSGAPANLLAVPWIELLVVPPALAGSLLLWVP 412
NTDB id 1199 PSJM300 12650 AFN78592.1 ......GRLGVPSW.....WRALLYAQWAMALGLLPAMLVLGLPVSLSGPLANLLAVPWISVLVVPLALLGSFLLWVP 411
consensus


logo
NTDB id 1426 RS RS05575 WP 011001070.1 559
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 489
NTDB id 1199 PSJM300 12650 AFN78592.1 488
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 639
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 568
NTDB id 1199 PSJM300 12650 AFN78592.1 567
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 719
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 634
NTDB id 1199 PSJM300 12650 AFN78592.1 634
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 799
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 712
NTDB id 1199 PSJM300 12650 AFN78592.1 712
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 847
NTDB id 1036095 AB9G19 RS09930 WP 033888196.1 741
NTDB id 1199 PSJM300 12650 AFN78592.1 741
consensus
X non conserved
X similar
X ≥ 50% conserved