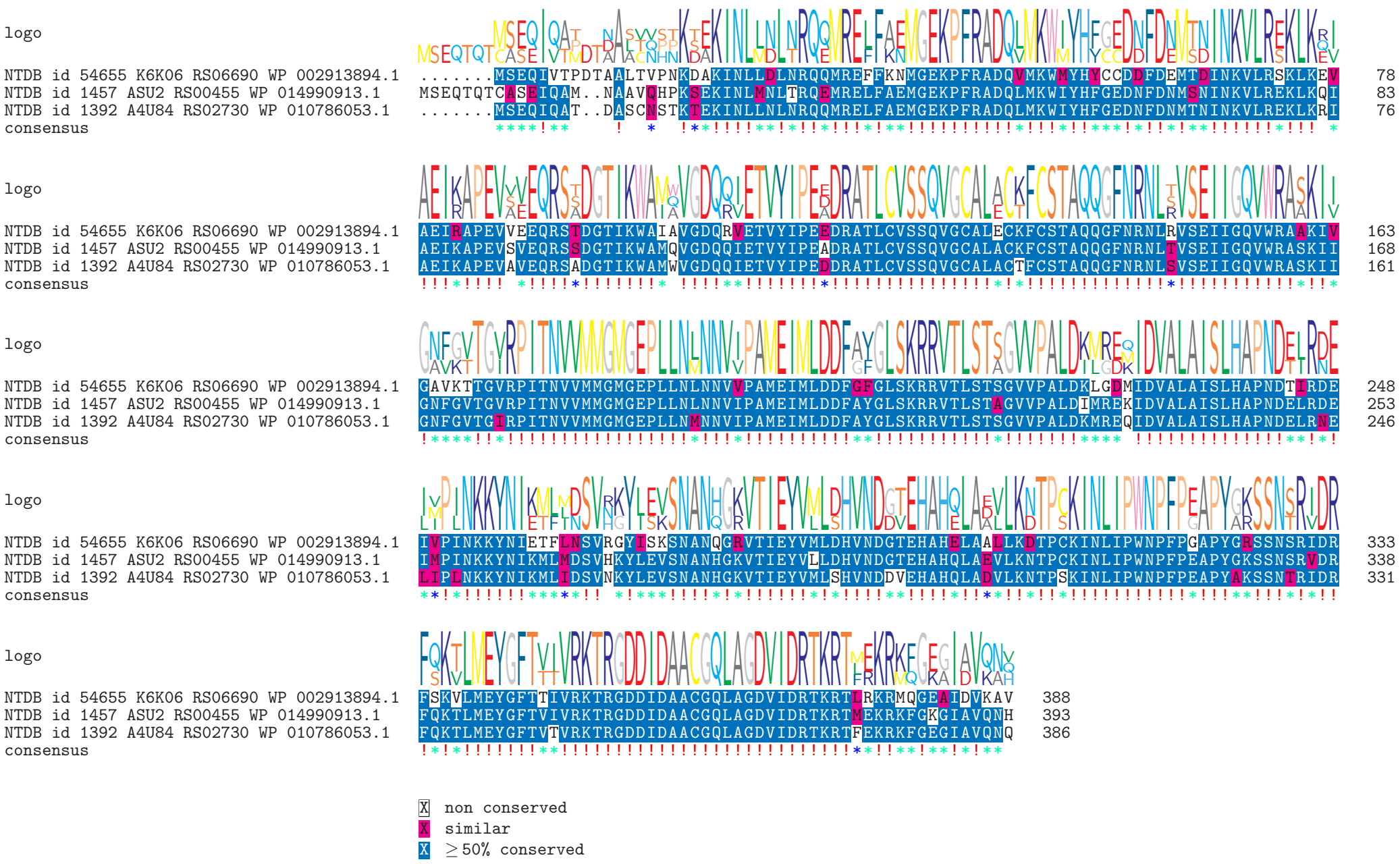

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NTDB id 1457 ASU2 RS00455 WP 014990913.1 MSEQTQTCASEIQAM..NAAVQHPKSEKINLMNLTRQEMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMSNINKVLREKLKQI 83
NTDB id 1392 A4U84 RS02730 WP 010786053.1 .....MSEQIQAT..DASCNSTKTEKINLLNLNRQQMRELFAEMGEKPFRADQLMKWIYHFGEDNFDNMTNINKVLREKLKRI 76
consensus ****!** ! * !*!!!!**!*!!*!!!*!**!!!!!!!!!!*!!!*!!***!*!!*!**!!!!!!*!!! *
logo
NTDB id 54655 K6K06 RS06690 WP 002913894.1 AEIRAPEVVEEQRSTDGTIKWAIAVGDQRVETVYIPEEDRATLCVSSQVGCALECKFCSTAQQGFNRNLRVSEIIGQVWRAAKIV 163
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consensus !!!*!!!! *!!!!*!!!!!!!* !!!!**!!!!!!!*!!!!!!!!!!!!!!!!*!*!!!!!!!!!!!!!*!!!!!!!!!!!*!!*
logo
NTDB id 54655 K6K06 RS06690 WP 002913894.1 GAVKTTGVRPITNVVMMGMGEPLLNLNNVVPAMEIMLDDFGFGLSKRRVTLSTSGVVPALDKLGDMIDVALAISLHAPNDTIRDE 248
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consensus !****!!*!!!!!!!!!!!!!!!!!!*!!!*!!!!!!!!!!**!!!!!!!!!!!!*!!!!!!!****** !!!!!!!!!!!!!!**!*!
logo
NTDB id 54655 K6K06 RS06690 WP 002913894.1 IVPINKKYNIETFLNSVRGYISKSNANQGRVTIEYVMLDHVNDGTEHAHELAALLKDTPCKINLIPWNPFPGAPYGRSSNSRIDR 333
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consensus **!*!!!!!!****!! *!***!!!!*!*!!!!!!*!*!!!!**!!!!*!!*!!*!*!!!!!!!!!!!*!!!**!!!*!*!!
logo
NTDB id 54655 K6K06 RS06690 WP 002913894.1 FSKVLMEYGFTTIVRKTRGDDIDAACGQLAGDVIDRTKRTLRKRMQGEAIDVKAV 388
NTDB id 1457 ASU2 RS00455 WP 014990913.1 FQKTLMEYGFTVIVRKTRGDDIDAACGQLAGDVIDRTKRTMEKRKFGKGIAVQNH 393
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consensus !*!*!!!!!!!**!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!**!**!*!**
X non conserved
X similar
X ≥ 50% conserved