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NTDB id 1113 AAA85695.1 219..1451( )           MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSAFTEEEARKKLAKRGIRPLQITRVKT......SSKRKITQEDITV   74
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NTDB id 1252 GCO85 RS07730 WP 011213805.1      ......MDKNSPPLLTFHYQGINKAGQ.KMEGDIQARSLAIAKADLRKQGIVTNKVVKKRKPLFDR..KNKKITQADITV   71
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consensus                                                * *  * **!*  *  ** !** * * **** !***** ****** *      **** !!**
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NTDB id 1070342 ACH784 RS14715 WP 404391440.1  FTRQIATMLTAGVPLIQSLEMISSGAKNKSLSKLIGSIADEVKAGQPLAKSLRAYPRYFDELYCDLIHSGEQSGALDKIF   156
NTDB id 1198 PSJM300 03950 AFN76868.1          FTRQMATMMKAGVPLLQSFDIIGEGFDNPNMRKLVDDLKQEVAAGNSFASSLRKKPQYFDDLYCNLVDSGEQSGSLETLL   150
consensus                                      **!!**!!* *!*!**!********  ** ****** ** ****!******** * *!* *** !***!!**! !* **
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NTDB id 1113 AAA85695.1 219..1451( )           DKLAIYKEKTQAIRKKVKTALTYPVSVIAVAIGLVFVMMIFVLPAFKEVYANMGAELPALTQTVMDMSDFFVSYGWMVLI   234
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NTDB id 1252 GCO85 RS07730 WP 011213805.1      DKVATYKEKIEIIKKKIKKALTYPIAVMVVALLVTAGLLIYVVPQFESLFKGFGADLPAMTRAVITMSEFMQAYWYIIFG   231
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NTDB id 1059 ABD1 RS01610 WP 000279215.1       DRVAIYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFQDLFASFGADLPAFTQMVVNMSKWMQEYWFIMII   233
NTDB id 1070342 ACH784 RS14715 WP 404391440.1  DRVAVYKEKAEALKSKIKKAMFYPIAVVVVALIVTSILLIFVVPQFQDIFSSFGAELPAFTLFVIGISEFMQKSWWIFLL   236
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consensus                                      ***!*!*!!** ** !***!* !!* !**!!**** **** !*! ! ***  *!!*!!**!* !* *!**** *******
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NTDB id 1016 ACIAD RS01680 WP 004920476.1      VIGAAIAGFLEARKRSKRFRDQLDRLVLRLPIVGDLVYKAIIARYSRTLATTFAAGVPLIDALASTAGATNNTVYEQAVL   313
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consensus                                      ********** ** *! *** ****!**!**!*** !* *!***!!****!*!!!!****!*********!**** !*
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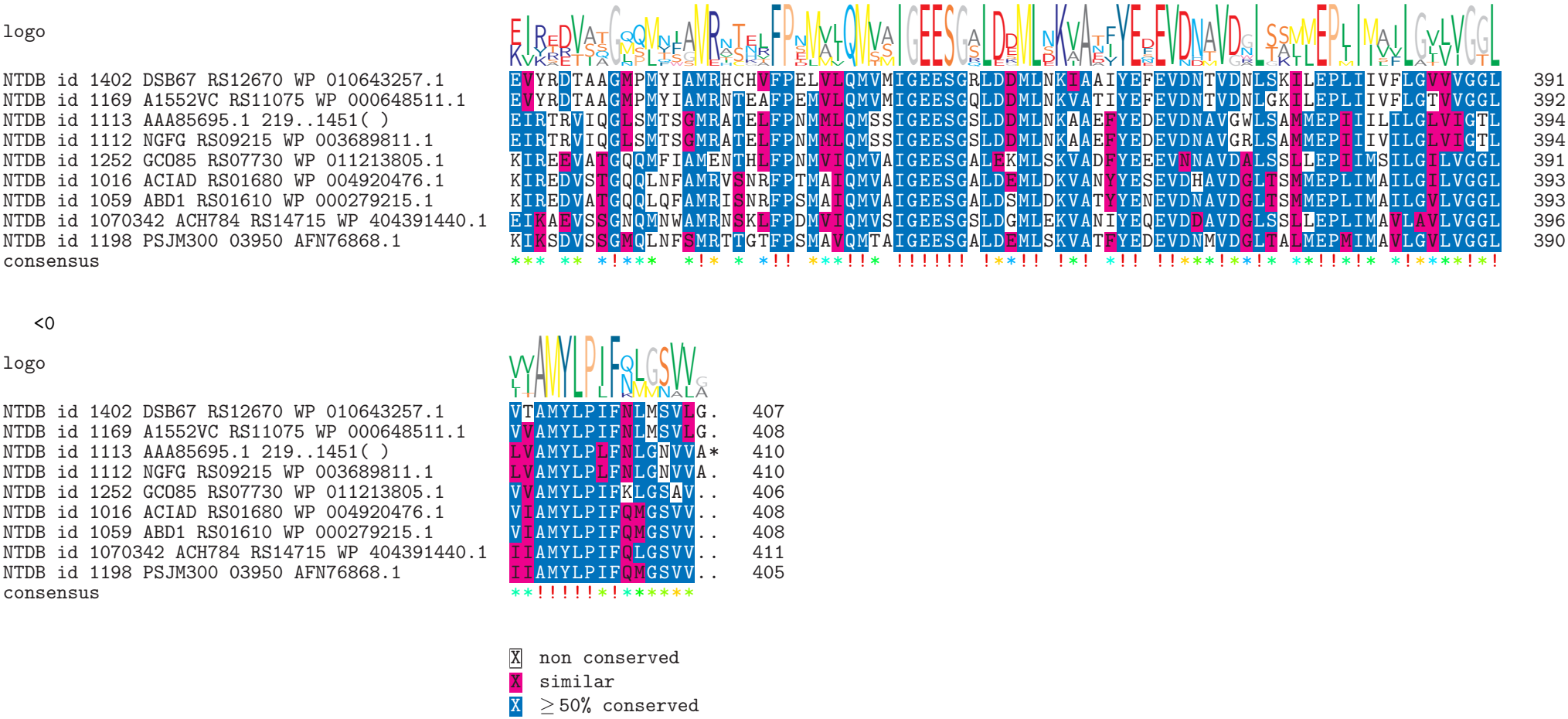
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consensus
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consensus
X non conserved
X similar
X ≥ 50% conserved