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NTDB id 1169 A1552VC RS11075 WP 000648511.1     .........MKATQTLPLKNYRWKGINSNGKK.VSGQMLAISEIEVRDKLKDQHIQIKKLKKGSVSLLAR.....LTHRVK   66
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NTDB id 1113 AAA85695.1 219..1451( )            MA...KNGGFSLFAKKEKRFIFECGRHSASDKLVNGEVSAFTEEEARKKLAKRGIRPLQITRVKT.........SSKRKIT   68
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NTDB id 1016 ACIAD RS01680 WP 004920476.1       .........MAAKKTQVMPVFAYEGVDRKGIK.LKGELPARNMALAKVTLRKQGITIRTIREKRKNIFEG....LLKKKVK   67
NTDB id 1059 ABD1 RS01610 WP 000279215.1        .........MAVKKAQMMPTFAYEGVDRKGVK.IKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEG....LFKKKVT   67
consensus                                                       *  **  *  ****  *  *  *  *  ***  *  *    **   !*****  **  ***  *   *          *****
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NTDB id 1113 AAA85695.1 219..1451( )            QEDITVFTRQLSTMINAGLPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVAAGETGG   148
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NTDB id 1059 ABD1 RS01610 WP 000279215.1        TLDITIFTRQLATMMKAGVPLVQGFEIVAEGLENPAMREVVLGIKGEVEQGSTFASALRKYPQHFDNLFCSLVESGEQSG   147
consensus                                         !*****!!***!***!****!******  *****  ******  **  ****!***   ******  *!*  ****!***!!**!
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NTDB id 1169 A1552VC RS11075 WP 000648511.1     NLPEVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMLTMVIPEFESMFKGFGAELPWFTQQVLKLSHWVQAY   226
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NTDB id 35080 KKOR RS03555 WP 012800641.1       TLEQMLDRIATYKEKTEALKSKIKSAMMYPAAVITVSLAVTAILLIYVVPMFEDLFSNAGADLPALTAMVVSMSENMQAN   233
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NTDB id 1059 ABD1 RS01610 WP 000279215.1        ALETMLDRVAIYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFQDLFASFGADLPAFTQMVVNMSKWMQEY   227
consensus                                        !* *****!***!!** ** !***!* !!* !*************** **! ! ***  *!**!!**!**** *!******
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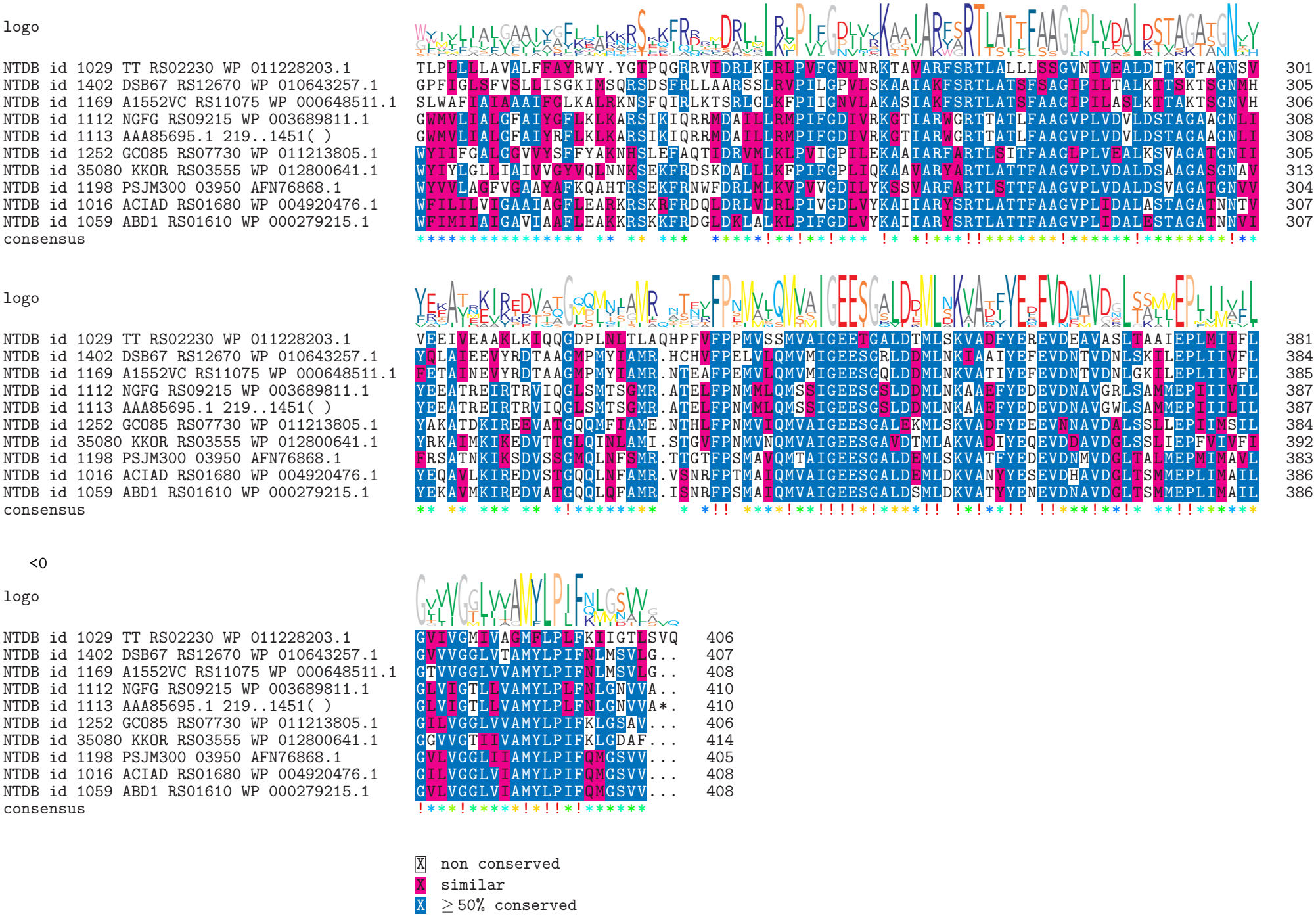
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consensus
logo
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consensus
X non conserved
X similar
X ≥50% conserved