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NTDB id 1113 AAA85695.1 219..1451( )          MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSAFTEEEARKKLAKRGIRPLQITRVKTS.....SKRKITQEDITVF   75
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consensus                                                * *  * **!* * * *** !** * ******  !***** *  *** * **       **** !****
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consensus                                     *!!***!***!*!**!********** ** ****** ** ****!***  **** * *!* ****!***!! *! !*****
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NTDB id 1151576 MKZ32 RS01240 WP 239795601.1  RLATYQEKTLAIKSKIKGALMYPVSIIAIAFIITAVIMIFVIPAFKQLFSSFGADLPAPTLIMMSISDFFVAYWWAIFSI   230
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NTDB id 1016 ACIAD RS01680 WP 004920476.1     RVAVYKEKSEVLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFKDLFTSFGADLPAFTQLVVNMSNWMQDYWFILILV   234
NTDB id 1059 ABD1 RS01610 WP 000279215.1      RVAIYKEKSELLKQKIKKAMKYPATVIVVAIVVTIILMVKVVPVFQDLFASFGADLPAFTQMVVNMSKWMQEYWFIMIIA   234
consensus                                     **!*!*!!** ** !***!* !!******!**** **** !*! ! ***  *!!*!!**!**** *!**** ********
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NTDB id 1113 AAA85695.1 219..1451( )          LGFAIYRFLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLDSTAGAAGNLIYEEATRE   315
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NTDB id 1252 GCO85 RS07730 WP 011213805.1     LGGVVYSFFYAKNHSLEFAQTIDRVMLKLPVIGPILEKAAIARFARTLSITFAAGLPLVEALKSVAGATGNIIYAKATDK   312
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consensus                                     ********   * ** ***  *****!**!**!*** !***!***!!****!*!!*!****! *******!****  !*
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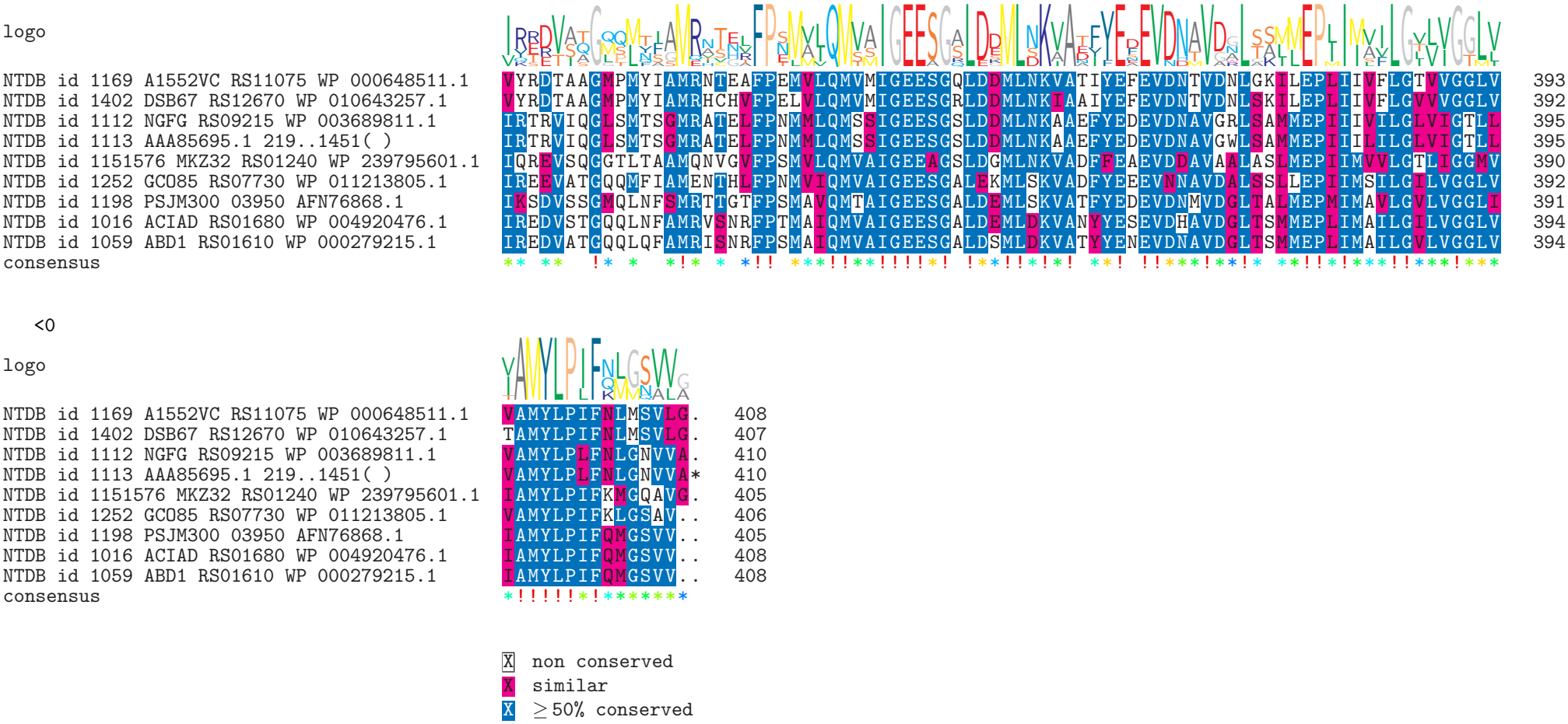
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NTDB id 1112 NGFG RS09215 WP 003689811.1 IRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYEDEVDNAVGRLSAMMEPIIIVILGLVIGTLL 395
NTDB id 1113 AAA85695.1 219..1451( ) IRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYEDEVDNAVGWLSAMMEPIIILILGLVIGTLL 395
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consensus
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consensus
X non conserved
X similar
X ≥ 50% conserved