logo

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NTDB id 1169 A1552VC RS11075 WP 000648511.1     .....MKATQTLPLKNYRWKGINSNG.KKVSGQMLAISEIEVRDKLKDQHIQIKKLKKGSVSLLAR.....   60
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NTDB id 1112 NGFG RS09215 WP 003689811.1        .....MAKNGGFSLFAKKEKRFIFEGRHSASDKLVNGEVSAFTEEEARKKLAKRGIRPLQITRVK.....TS....   62
NTDB id 369158 FIV34 RS05990 WP 139980636.1     MATATATKNARAVGAARAAVNQLTMYDWTALDKRG.KRMKGEMQAKNASLVKAELRRQGMNPQQVRERAKPLFGA.....   74
NTDB id 1252 GCO85 RS07730 WP 011213805.1       .....MDKNSPPLLTFHYQGINKAG.QKMEGDIQARSLAIAKADLRKQGIVTNKVVKKRKPLFDR.....   59
NTDB id 1198 PSJM300 03950 AFN76868.1           .....MAQKAIKNSVFTWEGLDRQG.AKIKGELSGVSPALVKAQLRKQGINPQKVRKKSISLFS.....   58
NTDB id 1016 ACIAD RS01680 WP 004920476.1       .....MAAKKTQVMPVFAYEGVDRKG.IKLKGELPARNMALAKVTLRKQGITIRTIREKRKNIFEGL....   61
NTDB id 1059 ABD1 RS01610 WP 000279215.1        .....MAVKKAQMMPTFAYEGVDRKG.VKIKGELPAKNMALAKVTLRKQGVTVRNIREKRKNILEGL....   61
consensus                                                * *  * ****** * *** *** * *  ***  !***** *  ****** *
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NTDB id 1112 NGFG RS09215 WP 003689811.1        SKRKITQEDITVFTRQLSTMIKAGLPLMQAFEIVARGHGNPSMTEMLMEIRGQVEQGSSLSRAFSNHPKYFDRFYCNLVA   142
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NTDB id 1059 ABD1 RS01610 WP 000279215.1        FKKKVTTLDITIFTRQLATMMKAGVPLVQGFEIVAEGLENPAMREVVLGIKGEVEGGSTFASALRKYPQHFDNLFCSLVE   141
consensus                                          ****  !*****!!**!!* *!****!********  ** ****** ** ****!**** **** * *!* ****!**
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NTDB id 1169 A1552VC RS11075 WP 000648511.1     TGEMSGNLPEVFERLATYREKSEQLRAKVIKALIYPSMVVLVALGVSYLMLTMVIPEFESMFKGFGAELPWFTQQVLKLS   220
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consensus                                       *!! *! !*******!***!*** ** !***!* !!* !***!********** **! ! **** ****!!**!**** *!
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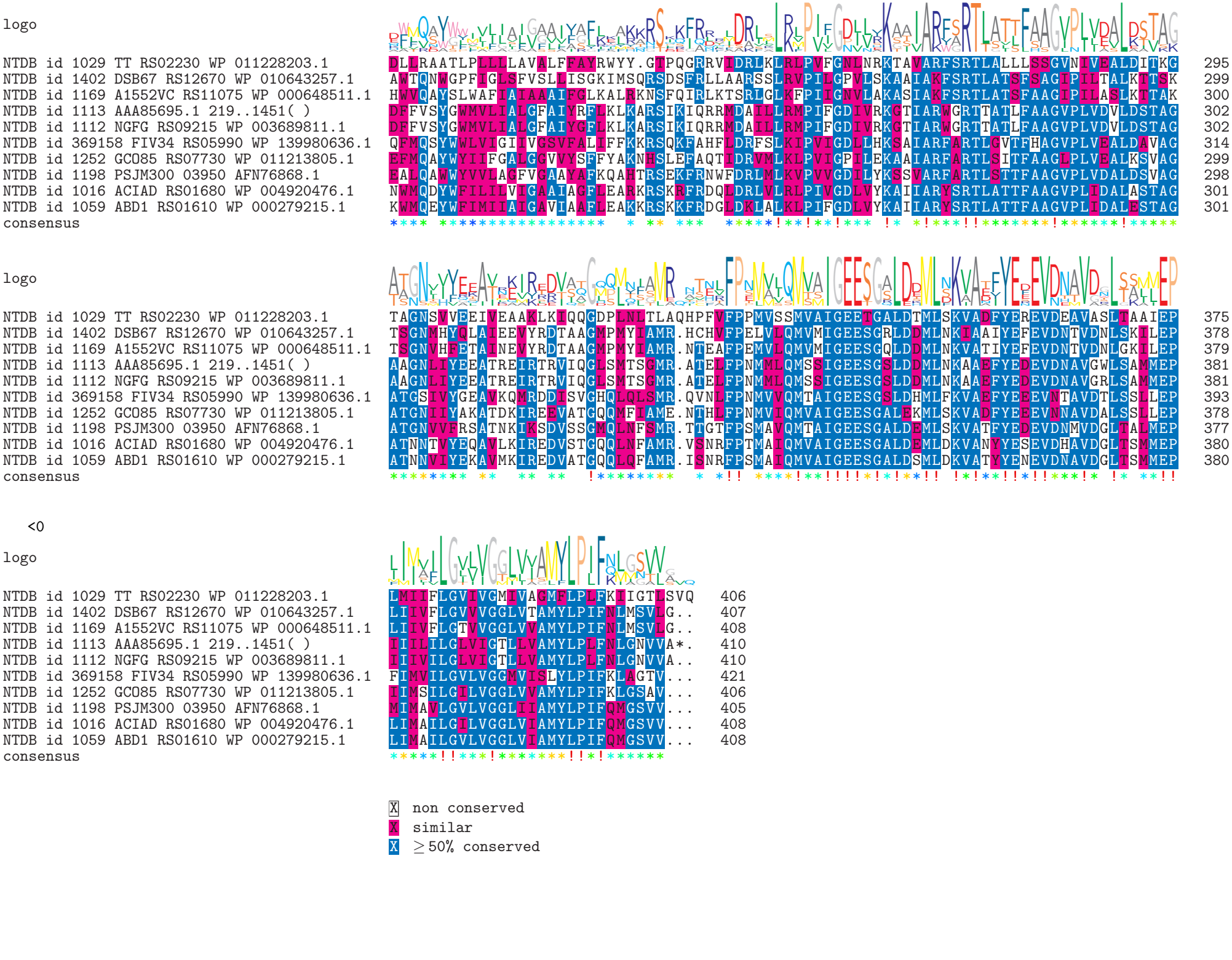
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consensus
logo
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consensus
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consensus
X non conserved
X similar
X ≥ 50% conserved