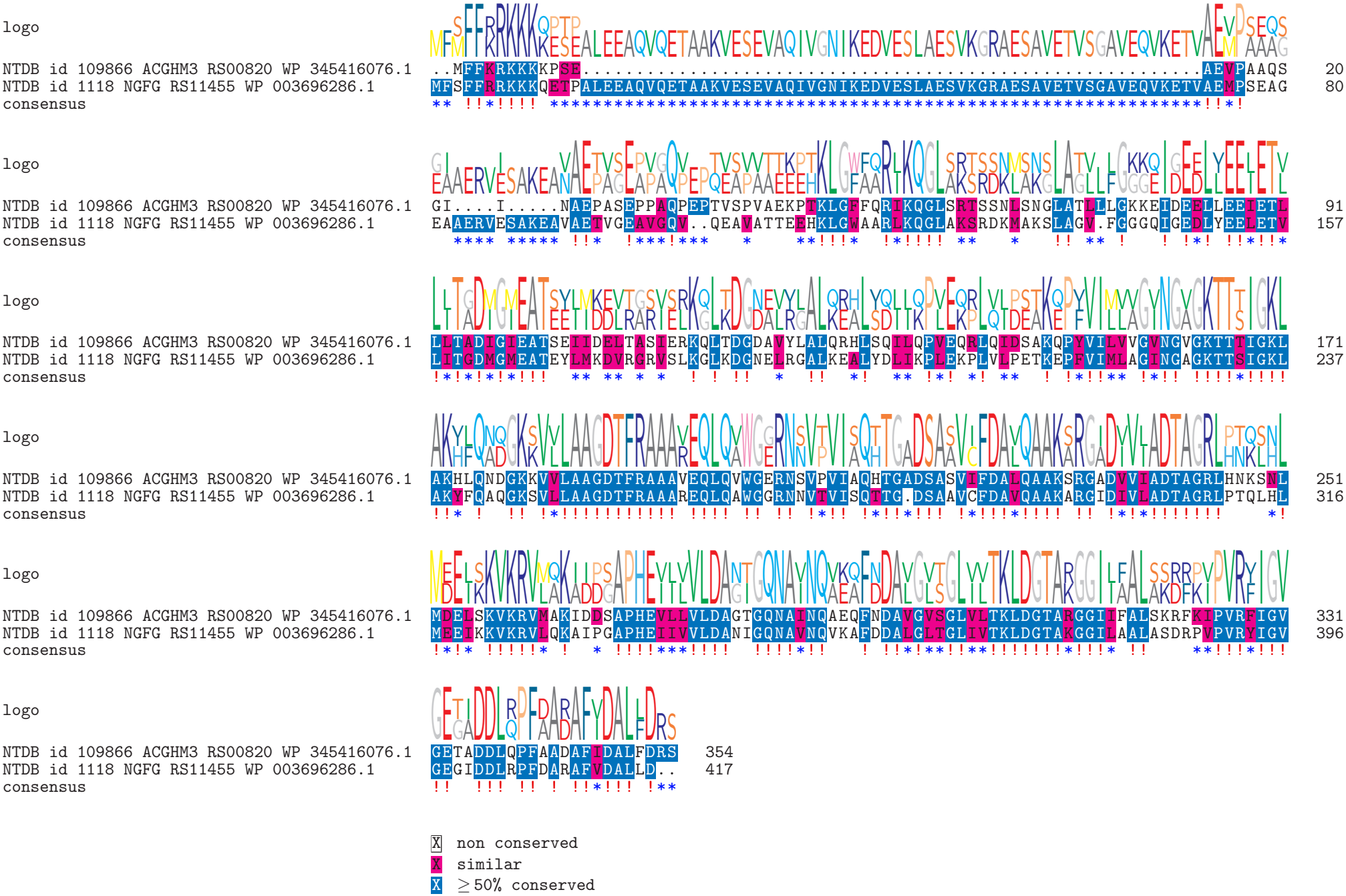
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NTDB id 109866 ACGHM3 RS00820 WP 345416076.1 ..MFFKRKKKKPSE.....AEVPAAQS 20
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consensus ** !!*!!!! ********** !!*!
logo
NTDB id 109866 ACGHM3 RS00820 WP 345416076.1 GI....I.....NAEPASEPPAQPEPTVSPVAEKPTKLGFFQRIKQGLSRTSSNLSNGLATLLLGKKEIDEELLEEIETL 91
NTDB id 1118 NGFG RS11455 WP 003696286.1 EAAERVESAKEAVAETVGEAVGQV..QEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGV.FGGGQIGEDLYEELETV 157
consensus **** ***** !!* !***!*** * **!!!* !*!!!! ** * !! ** ! ! !*! !!*!!*
logo
NTDB id 109866 ACGHM3 RS00820 WP 345416076.1 LLTADIGIEATSEIIDELTASIERKQLTDGDAVYLALQRHLSQILQPVEQRLQIDSAKQPYVILVVGVNGVGKTTTIGKL 171
NTDB id 1118 NGFG RS11455 WP 003696286.1 LITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKL 237
consensus !*!*!*!*!!! ** ** * * ! ! !! * !! *! ** !*! *! ** ! !*!!** !*!! !!!!*!!!!
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consensus !!* ! !! !*!!!!!!!!!!! !!!! !! !! !*!! !*!!*!!! !*!!!*!!!! !! !*!*!!!!!!! *!
logo
NTDB id 109866 ACGHM3 RS00820 WP 345416076.1 MDELSKVKRVMAKIDDSAPHEVLLVLDAGTGQNAINQAEQFNDAVGVSGLVLTKLDGTARGGIIFALSKRFKIPVRFIGV 331
NTDB id 1118 NGFG RS11455 WP 003696286.1 MEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGV 396
consensus !*!* !!!!!* ! * !!!!***!!!! !!!!*!! ! !!*!**!!**!!!!!!!*!!!* !! **!!!*!!!
logo
NTDB id 109866 ACGHM3 RS00820 WP 345416076.1 GETADDLQPFAADAFIDALFDRS 354
NTDB id 1118 NGFG RS11455 WP 003696286.1 GEGIDDLRPFDARAFVDALLD.. 417
consensus !! !!! !! ! !!*!!! !**
X non conserved
X similar
X ≥ 50% conserved