logo

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consensus
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consensus
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NTDB id 1110 NGFG RS09235 WP 003689817.1        ................................................................................     0
NTDB id 1251 GCO85 RS07725 WP 011946523.1       ................................................................................     0
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consensus
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NTDB id 1293 VP RS12240 WP 005479695.1          ..............MHSNLSTILRQKGLLTFSQEESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQ    66
NTDB id 1401 DSB67 RS12665 WP 010643259.1       ..............MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLSSGFFQSHELTEHLSSVFGLSCTSLNQ    66
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NTDB id 1110 NGFG RS09235 WP 003689817.1        ..............MSVGLLRILVQNQVVTVEQAEHYY.NESQAGKEVLPMLFSDGVISPKSLAALIARVFSYSILDLRH    65
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consensus                                             *       * ***  ***  ****  * * ** * *  *  *** ** ** ***  *** ** ***** ***
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NTDB id 1293 VP RS12240 WP 005479695.1          YEYASLCQQLGLRELITRHNALPLHRTPSTLLLAVADPTNQQAEDDFRFATG.....LQVELVLADFRELSTAIRRLYGR   141
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NTDB id 1018 ACIAD RS01685 WP 004920473.1       FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSK.....LKVEPIIVEHDKLERLLSEHFVE   149
consensus                                       ** * *   ** **** ** **!* **** * *****!*** **********     * *******   *  ********
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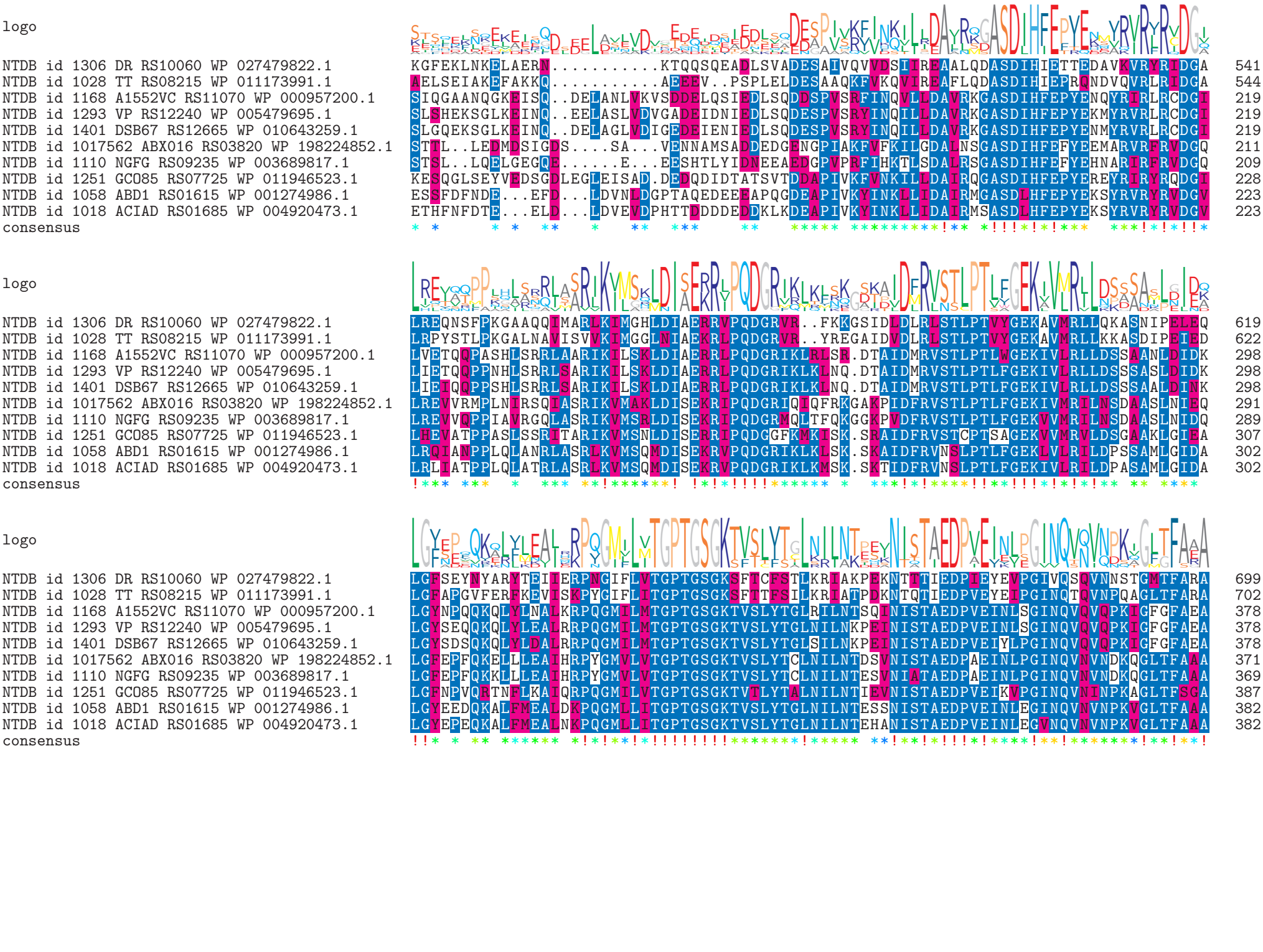
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consensus
logo
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NTDB id 1058 ABD1 RS01615 WP 001274986.1 LRQIANPPLQLANRLASRLKVMSQMDISEKRVPQDGRIKLKLSK.SKAIDFRVNSLPTLFGEKLVLRILDPSSAMLGIDA 302
NTDB id 1018 ACIAD RS01685 WP 004920473.1 LRLIATPPLQLATRLASRLKVMSQMDISEKRVPQDGRIKLKMSK.SKTIDFRVNSLPTLFGEKIVLRILDPASAMLGIDA 302
consensus
logo
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consensus

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consensus                                     !**!!!!!!!***!!!!!!**!!*!!**!**!!!*!**!!!!!*!*****!* *!!***!!**********!!!*!**!*
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consensus                                      !* *   *    *  ***** **      ******!! *!   !!*!!****!** ***** **** *** ***   **
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NTDB id 1018 ACIAD RS01685 WP 004920473.1     TLGFNNLRRSGLKKVMQGVTSLQEINRVTSE   570
consensus                                       !* *** *!**!***!*!***!***** *
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X non conserved
X similar
X ≥ 50% conserved