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NTDB id 1070916 ACIO07 RS06320 WP 410683430.1  ..........................................MAEYSLDE....DDIVYAVD.........GKCFEITPHLW   27
NTDB id 1355 HI 0298 AAC21962.1                ..........................................MTSYALLH....TQRVTAKN.........GEVFTISPDLW   27
NTDB id 1366 NTHI RS01950 WP 011271972.1       ..........................................MTSYALLH....TQRVTAQN.........GEIFTISPDLW   27
NTDB id 1168 A1552VC RS11070 WP 000957200.1    ........MLTNLVAILRQAELISATQEQAVVTQVSASGTSVPEALLELSIFHAQELTEQLSHIFGLPETDLSRYDYANL   72
NTDB id 1401 DSB67 RS12665 WP 010643259.1      ........MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLSSGFFQSHELTEHLSSVFGLSCTSLNQYEYASL   72
NTDB id 1293 VP RS12240 WP 005479695.1         ........MHSNLSTILRQKGLLTFSQEESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQYEYASL   72
NTDB id 1251 GCO85 RS07725 WP 011946523.1      MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAQNFGVPMLDINCIDVGTI   80
NTDB id 1018 ACIAD RS01685 WP 004920473.1      MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIVPYLIDNFSISPLTIAEIISLEFGEPLLDLGVFDPALF   80
consensus                                                 *     *                *   ***         *         **     **   **** *
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NTDB id 1070916 ACIO07 RS06320 WP 410683430.1  QQNKQQKDILIRYLAVPLQEDEHTLWLGIDDLTNLSACESFSFLTGKIVEPILLSEQDLKSLLQKLSPTLVEEPA.....  102
NTDB id 1355 HI 0298 AAC21962.1                ERNQQQQSLLLRYFALPLKEENNRLWLGVDSLSNLSACETIAFITGKPVEPILLESSQLKELLQQLTPNQMQVEE.....  102
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NTDB id 1401 DSB67 RS12665 WP 010643259.1      CQTLGLRDLITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATGLQVELVLADFRELTAAIRRLYGRSLGQEKSGLKE  152
NTDB id 1293 VP RS12240 WP 005479695.1         CQQLGLRELITRHNALPLHRTPSTLLLAVADPTNQQAEDDFRFATGLQVELVLADFRELSTAIRRLYGRSLSHEKSGLKE  152
NTDB id 1251 GCO85 RS07725 WP 011946523.1      PVNLVNEKLIKRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSEYVE  160
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consensus                                        *   * **  **  **!*       !**********  *  *  ! ***********    !   **   *      *
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NTDB id 1366 NTHI RS01950 WP 011271972.1       .......QVKFYQHQETHFE......QEDDEPVIRLLNQIFESALQKNASDIHLETLADQFQVRFRIDGVLQPQPLISKI  169
NTDB id 1168 A1552VC RS11070 WP 000957200.1    ISQDELANL...VKVSDDELQSIEDLSQDDSPVSRFINQVLLDAVRKGASDIHFEPYENQYRIRLRCDGILVETQQPASH  229
NTDB id 1401 DSB67 RS12665 WP 010643259.1      INQDELAGL...VDIGEDEIENIEDLSQDESPVSRYINQILLDAVRKGASDIHFEPYENMYRVRLRCDGILIEIQQPPSH  229
NTDB id 1293 VP RS12240 WP 005479695.1         INQEELASL...VDVGADEIDNIEDLSQDESPVSRYINQILLDAVRKGASDIHFEPYEKMYRVRLRCDGILIETQQPPNH  229
NTDB id 1251 GCO85 RS07725 WP 011946523.1      D.SGDLEGLEISAD.DEDQDIDTATSVTDDAPIVKFVNKILLDAIRQGASDIHFEPYEREYRIRYRQDGILHEVATPPAS  238
NTDB id 1018 ACIAD RS01685 WP 004920473.1      ..TEEL.DLDVEVDPHTTDDDDEDDKLKDEAPIVKYINKLLIDAIRMSASDLHFEPYEKSYRVRYRVDGVLRLIATPPLQ  233
consensus                                        **  *   *     *   *    ** !*****************  !!!*!*!***   ***!*!*!!*!      *
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NTDB id 1355 HI 0298 AAC21962.1                FANRIISRLKLLAKLDISENRLPQDGRFQFKTTFSDILDFRLSTLPTHWGEKIVLRAQQNKPVELSFSELGMTENQQQAF  249
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NTDB id 1401 DSB67 RS12665 WP 010643259.1      LSRRLSARIKILSKLDIAERRLPQDGRIKLKLNQDTAIDMRVSTLPTLFGEKIVLRLLDSSSAALDINKLGYSDSQKQLY  309
NTDB id 1293 VP RS12240 WP 005479695.1         LSRRLSARIKILSKLDIAERRLPQDGRIKLKLNQDTAIDMRVSTLPTLFGEKIVLRLLDSSSASLDIDKLGYSEQQKQLY  309
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NTDB id 1018 ACIAD RS01685 WP 004920473.1      LATRLASRLKVMSQMDISEKRVPQDGRIKLKMSKSKTIDFRVNSLPTLFGEKIVLRILDPASAMLGIDALGYEPEQKALF  313
consensus                                      ** !* *!*!*!*****!**!*!*!*!!!!*******  *   *!*!****!!**!!!*!*!***    *  ***   !!** *!**  *
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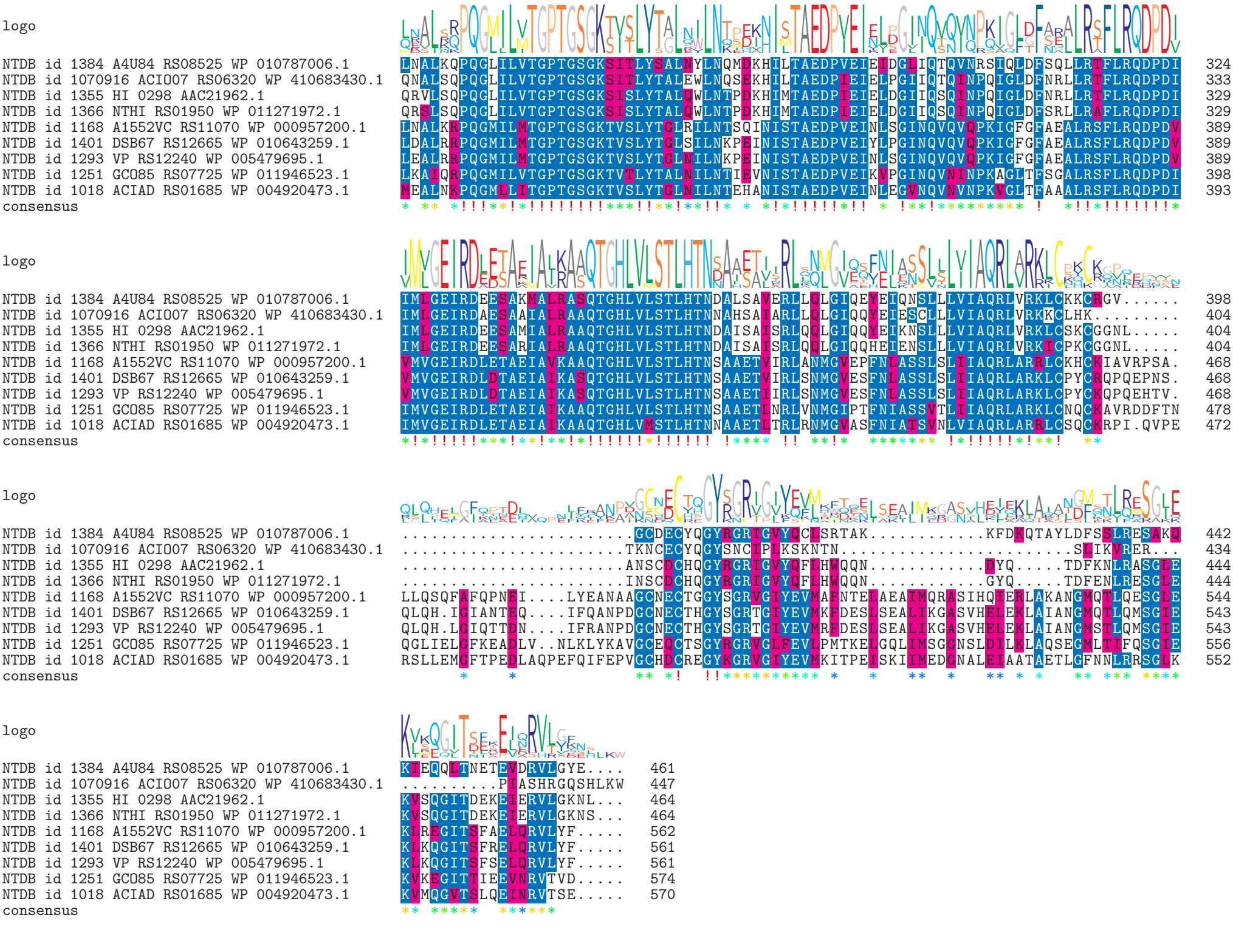
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NTDB id 1293 VP RS12240 WP 005479695.1 LEALRRPQGMILMTGPTGSGKTVSLYTGLNILNKPEINISTAEDPVEINLSGINQVQVQPKIGFGFAEALRSFLRQDPDV 389
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consensus
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consensus

X non conserved
X similar
X ≥ 50% conserved