logo

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NTDB id 511 SMSK321 RS10730 WP 000668322.1              ..............................MKFGKRHY..RPQVDQMDCGVASLAMVFGYYGSYYSLA   36
NTDB id 247 KZH43 RS00220 WP 000668290.1                ..............................MKFGKRHY..RPQVDQMDCGVASLAMVFGYYGSYYFLA   36
NTDB id 206 SPD RS00235 WP 000668290.1                  ..............................MKFGKRHY..RPQVDQMDCGVASLAMVFGYYGSYYFLA   36
NTDB id 171 SPR RS00230 WP 000668290.1                  ..............................MKFGKRHY..RPQVDQMDCGVASLAMVFGYYGSYYFLA   36
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NTDB id 1116478 EQB47 RS11985 WP 167750760.1            ..................................................................      0
NTDB id 358 SMU RS01425 WP 002263523.1                  MKQVIYVVLIVIAVNILLEIIKRVTKRGGTVSSSNPLPDGQSKLFWRRHYKLVPQIDTRDCGPAVLASVAKHYGSNYSIA   80
consensus                                                                              ********  ************************ **
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NTDB id 511 SMSK321 RS10730 WP 000668322.1              HLRELAKTTMDGTTALGLVKVAEEIGFETRAIKADMTLFDLPDLTFPFVAHVLKEGKLLHYYVVTGQDKGSIHIADPDPG   116
NTDB id 247 KZH43 RS00220 WP 000668290.1                HLRELAKTTMDGTTALGLVKVAEEIGFETRAIKADMTLFDLPDLTFPFVAHVLKEGKLLHYYVVTGQDKDSIHIADPDPG   116
NTDB id 206 SPD RS00235 WP 000668290.1                  HLRELAKTTMDGTTALGLVKVAEEIGFETRAIKADMTLFDLPDLTFPFVAHVLKEGKLLHYYVVTGQDKDSIHIADPDPG   116
NTDB id 171 SPR RS00230 WP 000668290.1                  HLRELAKTTMDGTTALGLVKVAEEIGFETRAIKADMTLFDLPDLTFPFVAHVLKEGKLLHYYVVTGQDKDSIHIADPDPG   116
NTDB id 131 SP RS00255 WP 000668284.1                   HLRELAKTTMDGTTALGLVKVAEEIGFETRAIKADMTLFDLPDLTFPFVAHVLKEGKLLHYYVVTGQDKDSIHIADPDPG   116
NTDB id 1116478 EQB47 RS11985 WP 167750760.1            ..................................................................      0
NTDB id 358 SMU RS01425 WP 002263523.1                  YLRELSKTNKQGTTALGIVEAAKKLGFETRSIKADMTLFDYNDLTYPFIVHVIKGKRLQHYYVVYGSQNNQLIIGDPDPS   160
consensus                                               **********************************************************************
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NTDB id 511 SMSK321 RS10730 WP 000668322.1              VKLTKLPRERFAEEWTGVTLFMAPSPDYKPHKDQKNGLLSFIPILVKQRGLIANIVLATLLVTVINIVGSYYLQSIIDTY   196
NTDB id 247 KZH43 RS00220 WP 000668290.1                VKLTKLPRERFEEEWTGVTLFMAPSPDYKPHKEQKNGLLSFIPILVKQRGLIANIVLATLLVTVINIVGSYYLQSIIDTY   196
NTDB id 206 SPD RS00235 WP 000668290.1                  VKLTKLPRERFEEEWTGVTLFMAPSPDYKPHKEQKNGLLSFIPILVKQRGLIANIVLATLLVTVINIVGSYYLQSIIDTY   196
NTDB id 171 SPR RS00230 WP 000668290.1                  VKLTKLPRERFEEEWTGVTLFMAPSPDYKPHKEQKNGLLSFIPILVKQRGLIANIVLATLLVTVINIVGSYYLQSIIDTY   196
NTDB id 131 SP RS00255 WP 000668284.1                   VKLTKLPRERFEEEWTGVTLFMAPSPDYKPHKEQKNGLLSFIPILVKQRGLIANIVLATLLVTVINIVGSYYLQSIIDTY   196
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consensus                                               **********************************************************!!*!*!*!!!!!!*!*!*!
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NTDB id 247 KZH43 RS00220 WP 000668290.1                VPDQMRSTLGIISIGLVIVYILQQILSYAQEYLLLVLGQRLSIDVILSYIKHVFHLPMSFFATRRTGEIVSRFTDANSII   276
NTDB id 206 SPD RS00235 WP 000668290.1                  VPDQMRSTLGIISIGLVIVYILQQILSYAQEYLLLVLGQRLSIDVILSYIKHVFHLPMSFFATRRTGEIVSRFTDANSII   276
NTDB id 171 SPR RS00230 WP 000668290.1                  VPDQMRSTLGIISIGLVIVYILQQILSYAQEYLLLVLGQRLSIDVILSYIKHVFHLPMSFFATRRTGEIVSRFTDANSII   276
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consensus                                               *!*****!!!*!**!!*!*!!*!!*******!!!*!!**!!*!!!!!!!!*!*!*!!!!!!!!!!!!!!*!!!!!!!*!!
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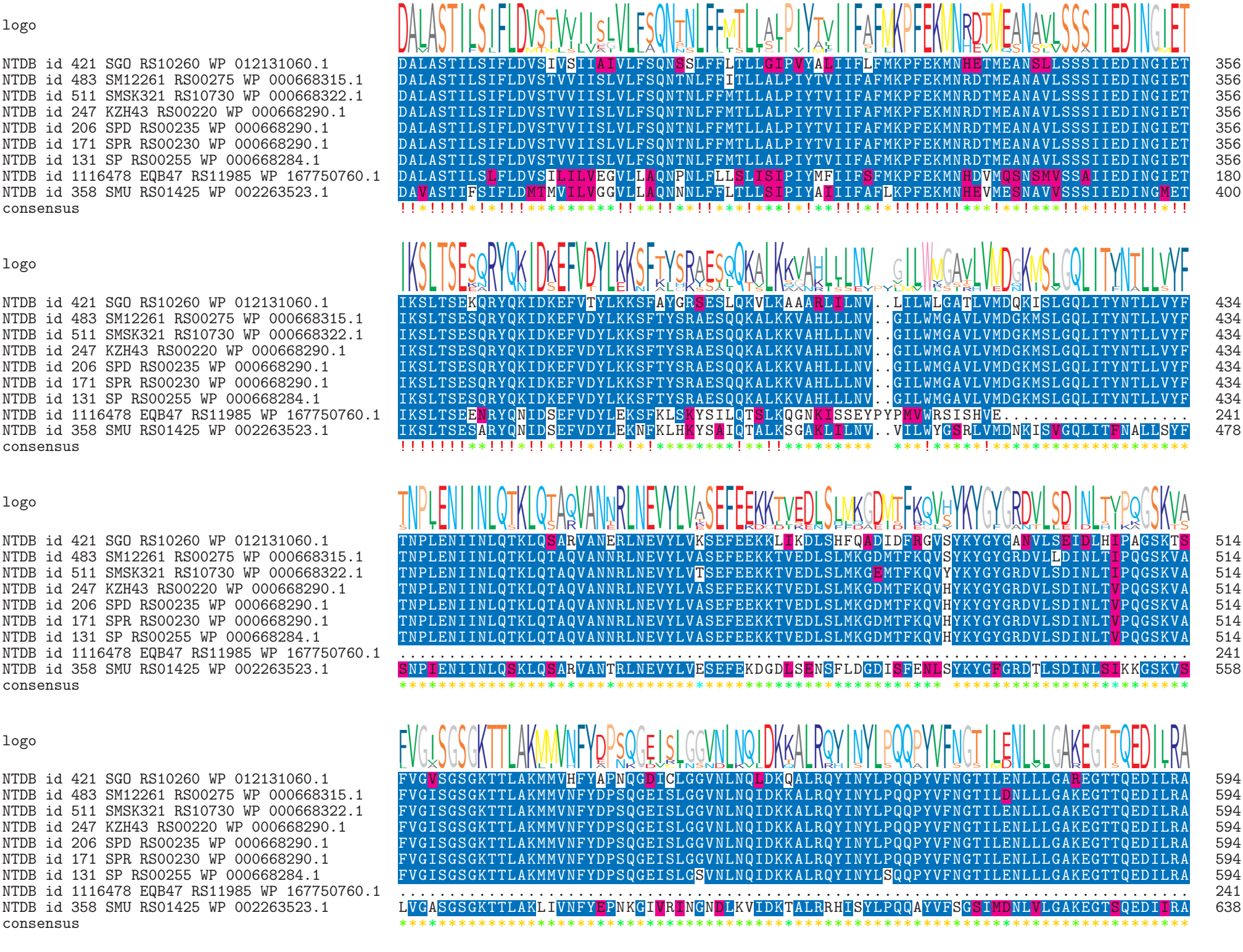
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NTDB id 131 SP RS00255 WP 000668284.1 DALASTILSIFLDVSTVVIISLVLFSQNTNLFFMTLLALPIYTVIIFAFMKPFEKMNRDTMEANAVLSSSIIEDINGIET 356
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consensus
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consensus
logo
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consensus
logo
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consensus

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logo                                                
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