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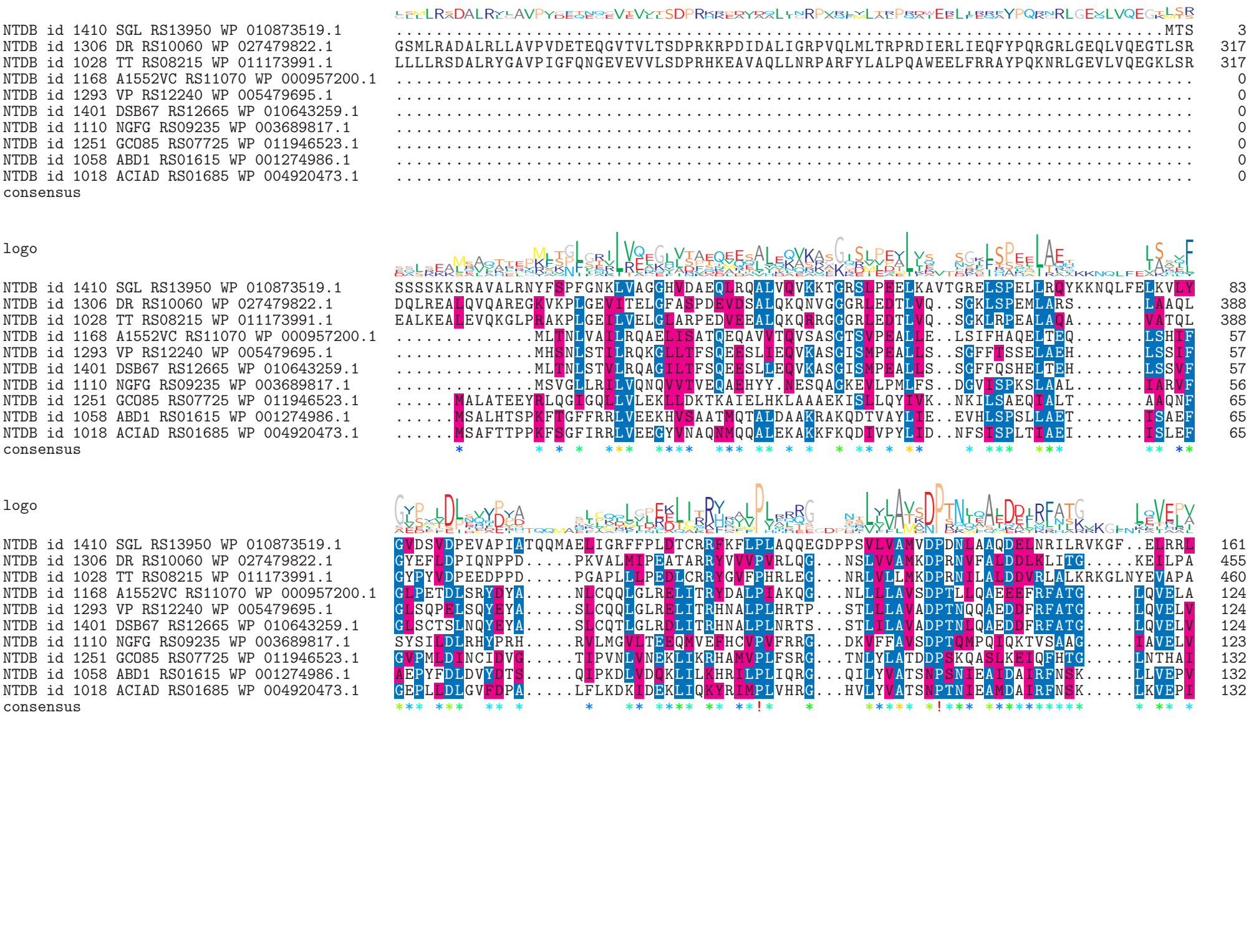
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NTDB id 1018 ACIAD RS01685 WP 004920473.1 GEPLLDLGVFDPA.....LFLKDKIDEKLIQKYRIMPLVHRG...HVLYVATSNPTNIEAMDAIRFNSK.....LKVEPI 132
consensus

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NTDB id 1028 TT RS08215 WP 011173991.1       VATEAAITKLIERFYGKAELSEIAKEF.........AK.............KQAEEEV..PSPLELDESAAQKFVKQVI  515
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NTDB id 1251 GCO85 RS07725 WP 011946523.1    VVETDKLSALIDNLLTAKESQGLSEYVEDSG.DLEGLEIS...........AD.DEDQDIDTATSVTDDAPIVKFVNKIL  199
NTDB id 1058 ABD1 RS01615 WP 001274986.1     IVEHHKLEKVLGQHFAEESSFDFNDE...EF.D...LDVN...........LDGPTAQEDEEEAPQGDEAPIVKYINKLL  194
NTDB id 1018 ACIAD RS01685 WP 004920473.1     IVEHDKLERLLSEHFVEETHFNFDTE...EL.D...LDVE...........VDPHTTDDDDEDDKLKDEAPIVKYINKLL  194
consensus                                    ***   *  ********    *   *   *    * **              **  ***   **     ************
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NTDB id 1293 VP RS12240 WP 005479695.1       MRVSTLPTLFGEKIVLRLLDSSSASLDIDKLGYSEQQKQLYLEALRRPQGMILMTGPTGSGKTVSLYTGLNILNKPEINI  348
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consensus                                    *!*!!!*!****!**!*******!**!*  *!**!*!!!!!***!!!*!!**!!**!**!**!!!*!**!!!!!*!*****
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logo



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consensus                                    !*  *!!***!**************!!!*!**!* !*  *    **   *  ***** **              ******!
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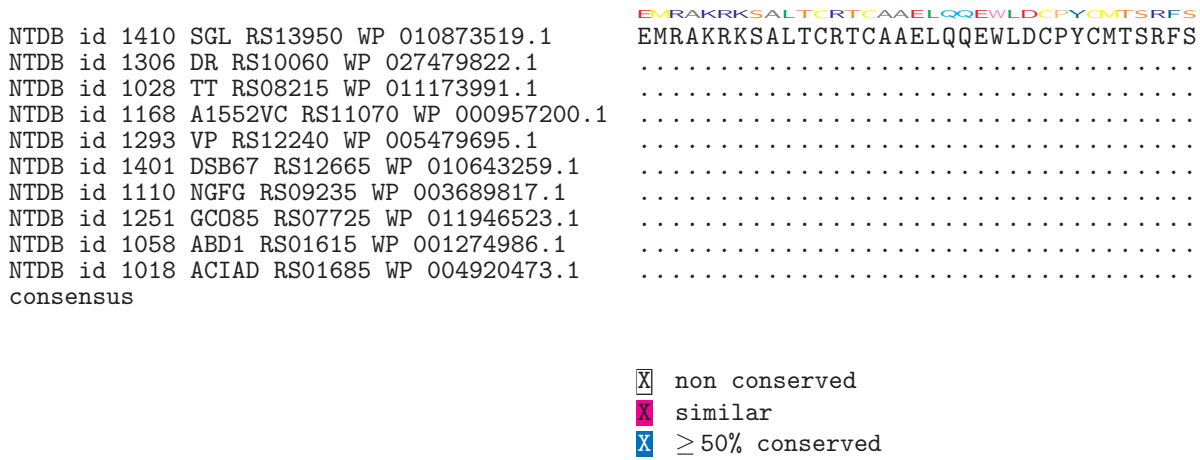
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consensus
X non conserved
X similar
X ≥ 50% conserved