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NTDB id 1122 OK783 RS09595 WP 003705341.1    ...MTEDERFAWLQLAFTPYIGAESFLLLMRSFGSAQNALSA.PAEQVAPAVRHKHALEAWRNAEKRALARQAAEAALEW   76
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NTDB id 1361 HI 0985 AAC22646.1              ....MNDITYTLLRLMQVPKLGGVGIDKILSNITLNEL.LNY.DDVAFRQMGWGAIQIRRWFKPEAKF.....IEPALVW   69
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NTDB id 1152 A1552VC RS00200 WP 000654772.1  ...MKDQDLAAWLALCFTPKLGSKTISHLLATRLPAQL.QSF.TPKQWLACGLKPEQLVFLTTQAAKQ.....AEQCLQW   70
NTDB id 203976 K10K4 RS16395 WP 054579448.1  MNRRNNDDLAAWLKLSCLPGIGGVKMNKLLARDTPCNI.VNF.SPEQLQQLGLTTKQIQAW.ASTDKE.....VDACLSW   72
NTDB id 1399 DSB67 RS15600 WP 010645755.1    MTPQNDIDLAAWLKLSCLPGIGGVKMNKLLSKDTPSNI.VQY.STEQLQLLGLTAKQLQAW.SQVDKE.....VDACLTW   72
consensus                                       * *** **** * *** *** ** *** * ** * * *** ** * * ** * * ***! *
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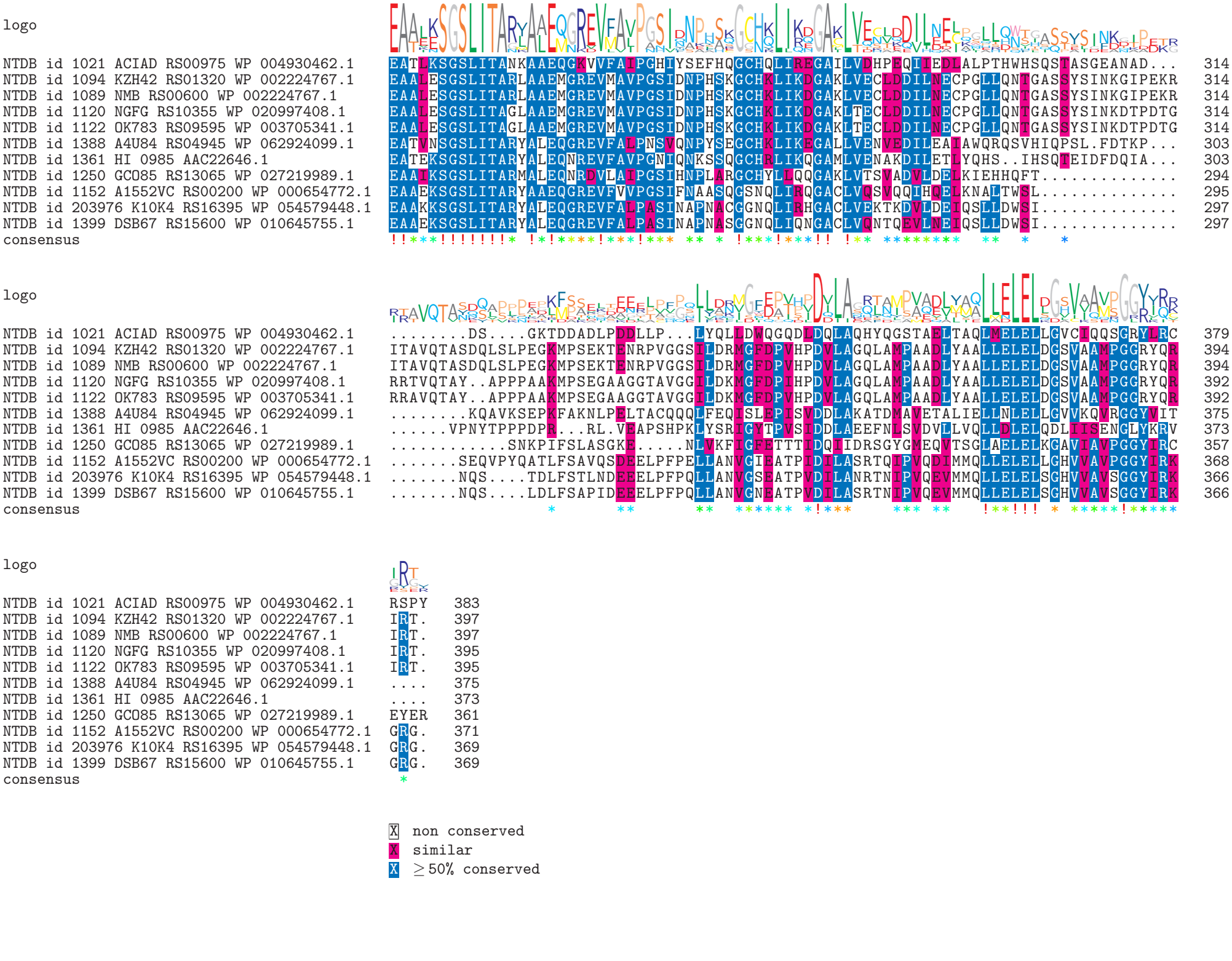


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NTDB id 1089 NMB RS00600 WP 002224767.1      EMRD.GCRLMLLQDEDFPEMLTQGLTAPPVLFLRGNVQLLHKPSAAIVGSRHATPQAMRIAKDFGKSLGGKGIPVVSGMA   155
NTDB id 1120 NGFG RS10355 WP 020997408.1     EMRD.GCRLMLLQDEDFPEMLTQGLTAPPVLFLRGNVRLLHKPSAAIVGSRHATPQAMRIAKDFGRALGGKGIPTVSGMA   155
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NTDB id 1250 GCO85 RS13065 WP 027219989.1    LSSSENHHILTWDSPYYPALLKEIADPPFILYAKGELSALTQPSLAIVGSRNASVTGNENARAFSREISRHGVSIVSGLA   149
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consensus                                       *    **** *  *! *!** ** !**!***!**  *  * *!*!!!! **  ** *** !* **   ** **!!*!
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NTDB id 1089 NMB RS00600 WP 002224767.1      SGIDTAAHQGALQAEGGTIAVWGTGIDRIYPPVNKNLAYEIAE.KGLIVSEFPIGTRPYAGNFPRRNRLIAALSQVTLVV   234
NTDB id 1120 NGFG RS10355 WP 020997408.1     SGIDTAAHQGALEAEGGTIAVWGTGIDRIYPPANKNLAYEIAE.KGLIVSEFPIGTRPYAGNFPRRNRLIAALSQVTLVV   234
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NTDB id 1388 A4U84 RS04945 WP 062924099.1    IGIDGFCHQRAANENGITIAVLGSGLDQVYPARHKKLAEQIVETSGALVSEFFPSQPPIAENFPRRNRIISGLSLGTLVI   227
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NTDB id 1152 A1552VC RS00200 WP 000654772.1  LGIDGYAHDGALQAQGQTVAVLGSGLAQVYPKQHQGLAERIIA.QGALVSEFAPHTPPKADHFPRRNRIISGLSLGVVVV   229
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consensus                                    *!!!* *!******* * **!!*!*!** *!!  ** !*  * *  !***!!! * **! ***!!*!!!****!!****!*
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NTDB id 1120 NGFG RS10355 WP 020997408.1 EAALESGSLITAGLAAEMGREVMAVPGSIDNPHSKGCHKLIKDGAKLTECLDDILNECPGLLQNTGASSYSINKDTPDTG 314
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consensus
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consensus
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X non conserved
X similar
X ≥50% conserved