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NTDB id 1361 HI 0985 AAC22646.1              QKE.GNHLVNYFSPFYPFLLKQTASFPPLLFVKGNLTALSQRQMAMVGSRYCTTYGEYWAKHFATELSLAGFTITSGLAL   149
consensus                                     ** *  **** **** *!** ***!**!***!**  ** ***!*!!!! **  ** *** !***** ** * !!*!*
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NTDB id 1120 NGFG RS10355 WP 020997408.1     GIDTAAHQGALEAEGGTIAVWGTGIDRIYPPANKNLAYEIAE.KGLIVSEFPIGTRPYAGNFPRRNRLIAALSQVTLVVE   235
NTDB id 1122 OK783 RS09595 WP 003705341.1    GIDTAAHQGALEAEGGTVAVWGTGIDRIYPPANKNLAYEIAE.KGLIVSEFPIGTRPYAGNFPRRNRLIAALSQVTLVVE   235
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NTDB id 1361 HI 0985 AAC22646.1              GIDGHCHQAVVNIQGQTIAVLGSGLEQIYPSKHQRLSAQIIENNGALVSEFLPNQAPIAANFPRRNRIISGLSVGTLVVE   229
consensus                                    !!!* *!***** ! !*!!*!*!****!! *** !*  *** *! **!!!** **! !**!!*!!!*!**!!****!*!
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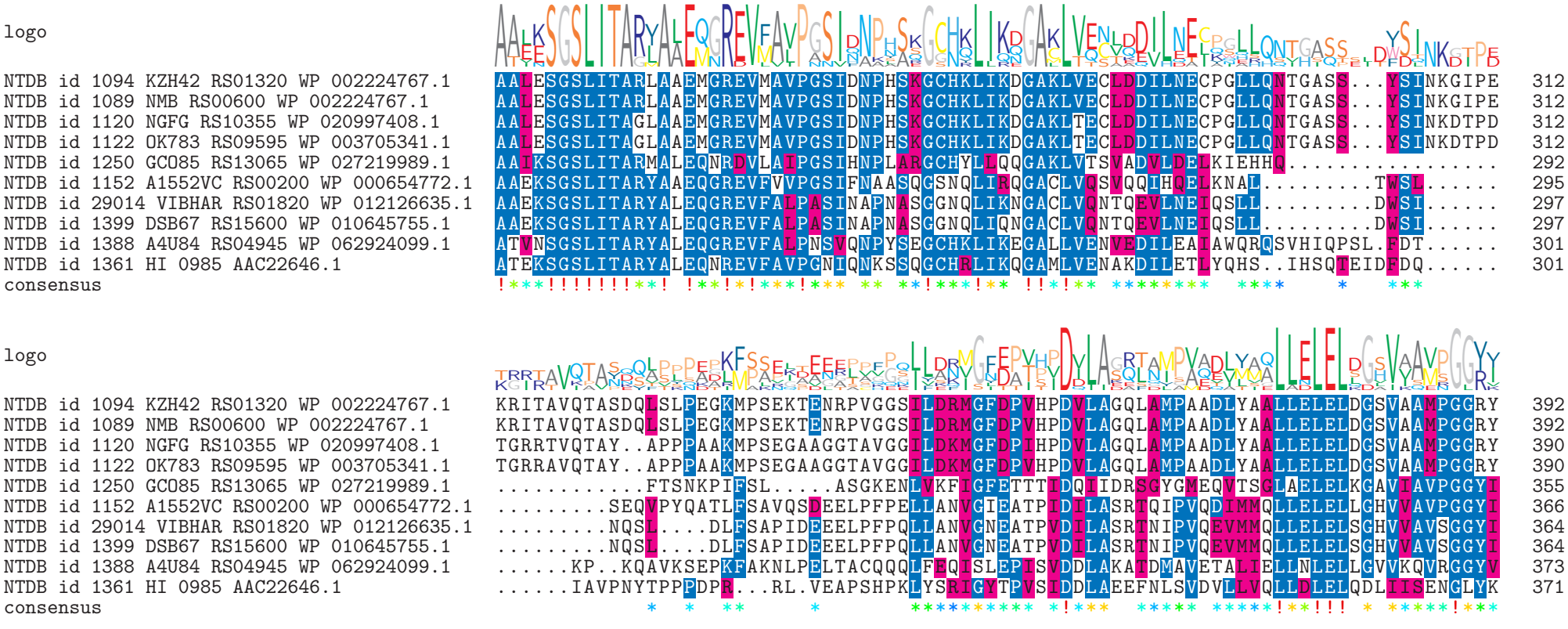
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consensus
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consensus
X non conserved
X similar
X ≥50% conserved