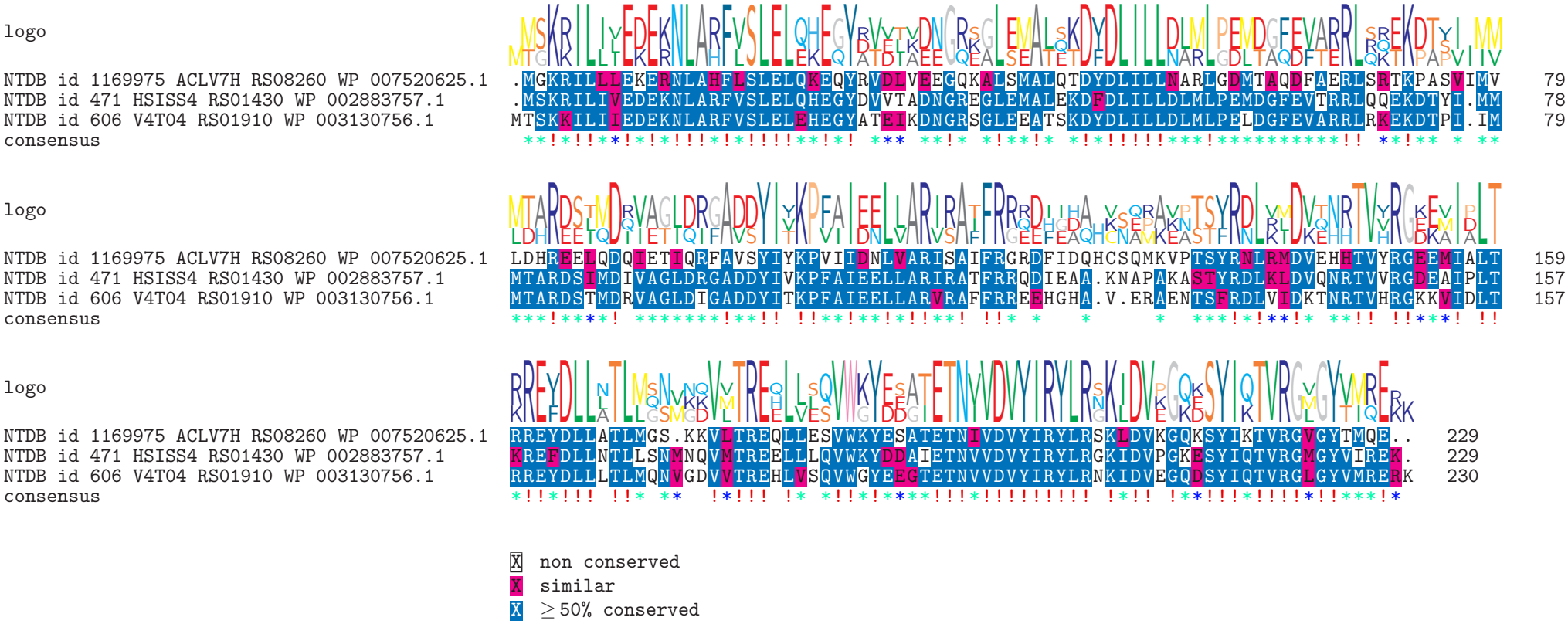
logo
NTDB id 1169975 ACLV7H RS08260 WP 007520625.1 .MGKRILLLEKERNLAHFLSLELQKEQYRVDLVEEGQKALSMALQTDYDLILLNARLGDMTAQDFAERLSRTKPASVIMV 79
NTDB id 471 HSISS4 RS01430 WP 002883757.1 .MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNGREGLEMALEKDFDLILLDLMLPEMDGFEVTRRLQQEKDTYI.MM 78
NTDB id 606 V4T04 RS01910 WP 003130756.1 MTSKKILIIEDEKNLARFVSLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKEKDTPI.IM 79
consensus **!*!!**!*!*!!!*!*!!!!**!*! *** **!* *!**!* *!*!!!!!***!********** **!** * **
logo
NTDB id 1169975 ACLV7H RS08260 WP 007520625.1 LDHREELQDQIETIQRFAVSYIYKPVIIDNLVARISAIFRGRDFIDQHCSQMKVPTSYRNLRMDVEHHTVYRGEEMIALT 159
NTDB id 471 HSISS4 RS01430 WP 002883757.1 MTARDSIMDIVAGLDRGADDYIVKPFAIEELLARIRATFRRQDIEAA.KNAPAKASTYRDLKLDVQNRTVVRGDEAIPLT 157
NTDB id 606 V4T04 RS01910 WP 003130756.1 MTARDSTMDRVAGLDIGADDYITKPFAIEELLARVRAFFRREEHGHA.V.ERAENTSFRDLVIDKTNRTVHRGKKVIDLT 157
consensus ***!*****! *******!**!! !!**!**!*!!**! !!* * * * *** !*!**!* **!! !!***! !!
logo
NTDB id 1169975 ACLV7H RS08260 WP 007520625.1 RREYDLLATLMGS.KKVLTREQLLESVWKYESATETNIVDVYIRYLRSKLDVKGQKSYIKTVRGVGYTMQE.. 229
NTDB id 471 HSISS4 RS01430 WP 002883757.1 KREFDLLNTLLSNMNQVMTREELLLQVWKYDDAIETNVVDVYIRYLRGKIDVPGKESYIQTVRGMGYVIREK. 229
NTDB id 606 V4T04 RS01910 WP 003130756.1 RREYDLLLTLMQNVGDVVTREHLVSQVWGYEEGTETNVVDVYIRYLRNKIDVEGQDSYIQTVRGLGYVMRERK 230
consensus *!!*!!! !!* ** !*!!! !* *!!*!*****!!!*!!!!!!!!! !*!! !**!!!*!!!!*!!***!*
X non conserved
X similar
X ≥ 50% conserved