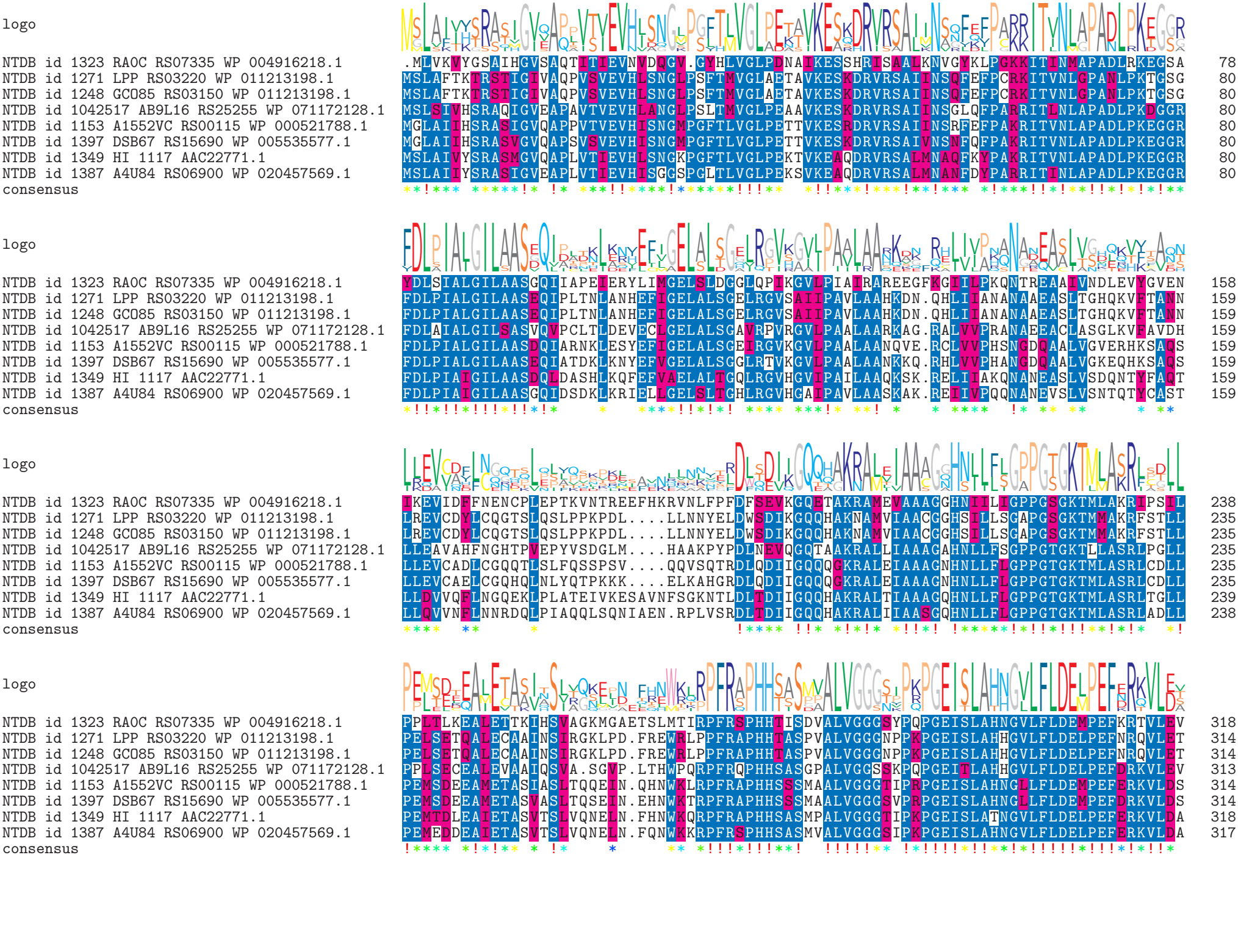
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NTDB id 1271 LPP RS03220 WP 011213198.1 MSLAFTKTRSTIGIVAQPVSVEVHLSNGLPSFTMVGLAETAVKESKDRVRSAIINSQFEFPCRKITVNLGPANLPKTGSG 80
NTDB id 1248 GCO85 RS03150 WP 011213198.1 MSLAFTKTRSTIGIVAQPVSVEVHLSNGLPSFTMVGLAETAVKESKDRVRSAIINSQFEFPCRKITVNLGPANLPKTGSG 80
NTDB id 1042517 AB9L16 RS25255 WP 071172128.1 MSLSIVHSRAQIGVEAPAVTVEVHLANGLPSLTMVGLPEAAVKESKDRVRSAIINSGLQFPARRITLNLAPADLPKDGGR 80
NTDB id 1153 A1552VC RS00115 WP 000521788.1 MGLAIIHSRASIGVQAPPVTVEVHISNGMPGFTLVGLPETTVKESRDRVRSAIINSRFEFPAKRITVNLAPADLPKEGGR 80
NTDB id 1397 DSB67 RS15690 WP 005535577.1 MGLAIIHSRASVGVQAPSVSVEVHISNGMPGFTLVGLPETTVKESKDRVRSAIVNSNFQFPAKRITVNLAPADLPKEGGR 80
NTDB id 1349 HI 1117 AAC22771.1 MSLAIVYSRASMGVQAPLVTIEVHLSNGKPGFTLVGLPEKTVKEAQDRVRSALMNAQFKYPAKRITVNLAPADLPKEGGR 80
NTDB id 1387 A4U84 RS06900 WP 020457569.1 MSLAIIYSRASIGVEAPLVTIEVHISGGSPGLTLVGLPEKSVKEAQDRVRSALMNANFDYPARRITINLAPADLPKEGGR 80
consensus
logo
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NTDB id 1248 GCO85 RS03150 WP 011213198.1 FDLPIALGILAASEQIPLTNLANHEFIGELALSGELRGVSAIIPAVLAAHKDN.QHLIIANANAAEASLTGHQKVFTANN 159
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consensus
logo
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consensus
logo
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consensus

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NTDB id 1042517 AB9L16 RS25255 WP 071172128.1   LREPLESGHIVIARAKDRVRFPARFQLVAAMNPCPCGYLGEPSGKCSCTPDMVQRYRNKLSGPLLDRIDLHLTVARE..A   391
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consensus                                       *!*!!!*****!*!!  ***  *!!*!*******!!*!*!  *  * *    **  *  !!* *!!*!*!!!  !* ***  *  *
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NTDB id 1248 GCO85 RS03150 WP 011213198.1       ELIKPNTHLEKQSLAIREKVTKMHEIQMARQD....SLNANLNSKTCEMVCELGSEEQLFLREVMSKLKLSARGYHRLLK   470
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NTDB id 1387 A4U84 RS06900 WP 020457569.1       ALQNNTDQRGETTEQARKRVLEARHLQFNRRG....KINAQLTTKEIEQDCKLTDRDALFLENALTKLGLSVRAYHRILK   471
consensus                                        *       *** *  **  *!  ****** ! *     *!! * *****  ! !   *  *** **  *! !!*!***!**!
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NTDB id 1248 GCO85 RS03150 WP 011213198.1       VSRTIADMNSSKKVLLNHLQQALSYKQNLHLPK.......   503
NTDB id 1042517 AB9L16 RS25255 WP 071172128.1   VARTLADLEQADAITREHLAEALQYRPATP..........   497
NTDB id 1153 A1552VC RS00115 WP 000521788.1     VARTIADLQGEAQIARPHLAEALGYRAMDRLLKQLSAQNV   507
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NTDB id 1349 HI 1117 AAC22771.1                 VSRTIADLQGEQQIFQPHLAEALGYRAMVRLLQKLSNM..   509
NTDB id 1387 A4U84 RS06900 WP 020457569.1       VSRTIADLAGEAQIQQPHLAEALGYRAMDRLLQRLQGES.   510
consensus                                       !*!!*!!*     **    !***!* !* * **    *
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X non conserved
X similar
X ≥ 50% conserved