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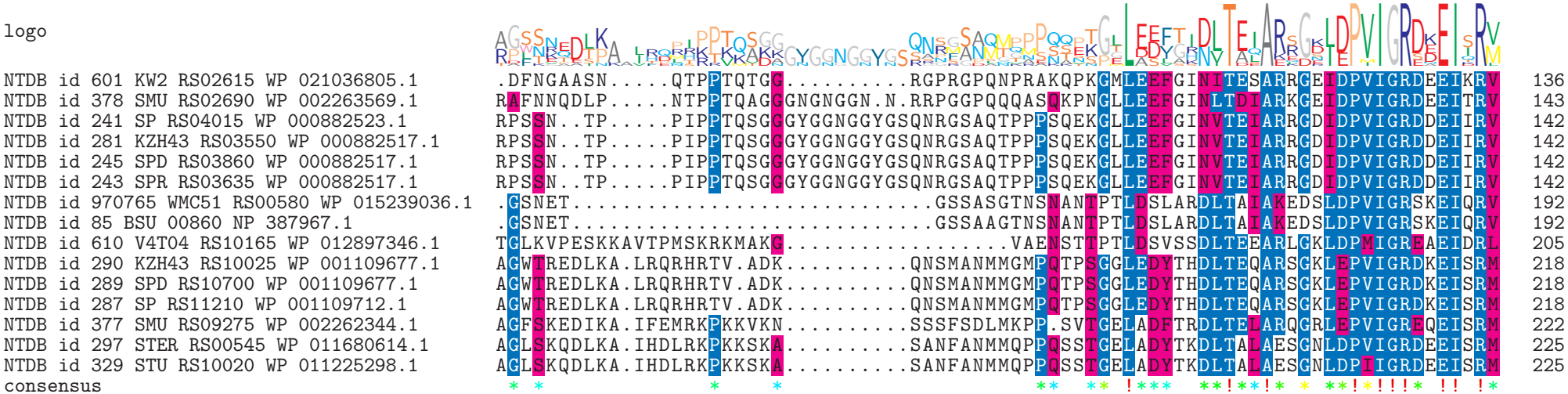
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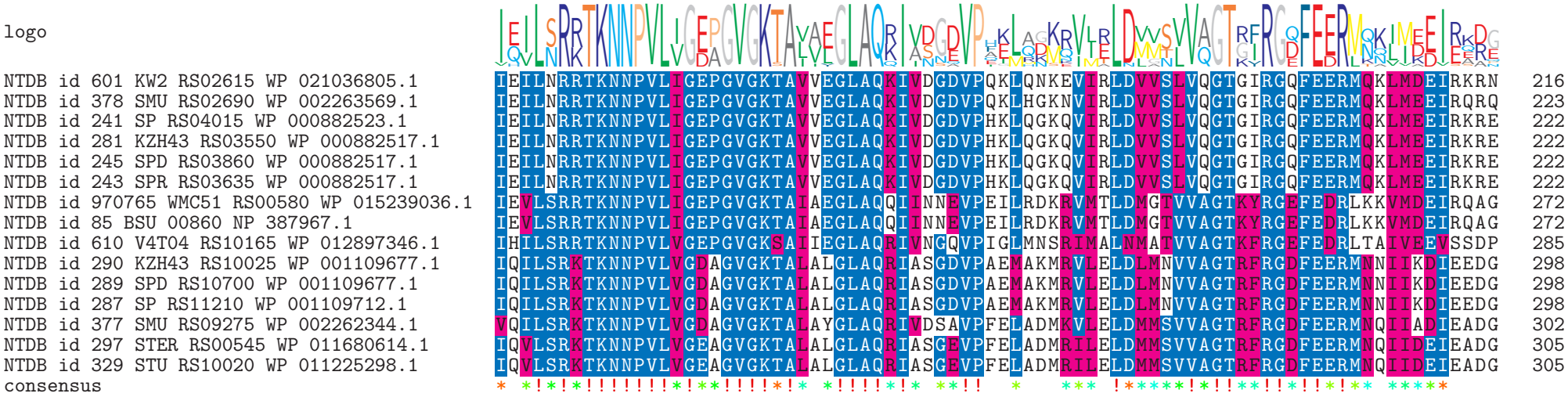
consensus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consensus

logo

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logo

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consensus                                     !** **!**! !  *! !***!* ** !!** * !!!*!!!!*!!** *** *  * ***  * *  !*
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consensus                                    *!!**!*  * *!*!!**  !* * *!!*!!!*!*!!!!!!!!!!*****!* *!* ** **!*!!!!*!!!*  **!*!*!
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consensus                                   !!!!!!*!*!*!!!*!! *!!!**!*!!*!!!!!!*****!!*!*!! *!!**!! !*!*****!!!!!*!** ** ***
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NTDB id 289 SPD RS10700 WP 001109677.1      VGFGAKDIRFDQENMEKRMFEELKKAYRPEFINRIDEKVVFHSLSSDHMQEVVKIMVKPLVASLAEKGIDLKLQASALKL  754
NTDB id 287 SP RS11210 WP 001109712.1       VGFGAKDIRFDQENMEKRMFEELKKAYRPEFINRIDEKVVFHSLSSDHMQEVVKIMVKPLVASLTEKGIDLKLQASALKL  754
NTDB id 377 SMU RS09275 WP 002262344.1      VGFGAKSFSQDYKAMESRILEELKKVYRPEFINRIDEKVVFHNLGQEDIRHIVKIMVAPLIAHLADQGITLKFQPSALKH  757
NTDB id 297 STER RS00545 WP 011680614.1     VGFGAQTISHNHQAMQARIMEELKKSYRPEFINRIDEKVVFHSLEEEQLHDIVKIMVKPLISALADKGISLKFQPAALKH  760
NTDB id 329 STU RS10020 WP 011225298.1      VGFGAQTISHNHQAMQARIMEELKKSYRPEFINRIDEKVVFHSLEEEQLHDIVKIMVKPLISALADKGISLKFQPAALKH  760
consensus                                   !!!**         *  *** *!** **!!!*!!*!* **!* !* *  *  *! *!*  *   * ** * * **  *
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NTDB id 601 KW2 RS02615 WP 021036805.1 LVDLGYNPAMGARPLRRTIQENIEDSIADFYIEHPEYKQ.LVADLIDDKIVISNQTQETAETTDEEVPAE 748
NTDB id 378 SMU RS02690 WP 002263569.1 LVDLGYDPKMGARPLRRTIQEHIEDAITDFYLEHPEQKE.LKATMTSNGKIVIKASNK.VETVTKATSD. 753
NTDB id 241 SP RS04015 WP 000882523.1 LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK. 752
NTDB id 281 KZH43 RS03550 WP 000882517.1 LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK. 752
NTDB id 245 SPD RS03860 WP 000882517.1 LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK. 752
NTDB id 243 SPR RS03635 WP 000882517.1 LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK. 752
NTDB id 970765 WMC51 RS00580 WP 015239036.1 VAEEGVDLEYGARPLRRAIQKHVEDRLSEELLRGNIDKGQHIVLDVEDGEFVVKTTAKTN..... 810
NTDB id 85 BSU 00860 NP 387967.1 VAEEGVDLEYGARPLRRAIQKHVEDRLSEELLRGNIHKGQHIVLDVEDGEFVVKTTAKTN..... 810
NTDB id 610 V4T04 RS10165 WP 012897346.1 IAEVGFDPEYGARPLRKALQKEVEDLLSEQLLSGEIKAGNHVSIGASNKKIKIAQIV..... 816
NTDB id 290 KZH43 RS10025 WP 001109677.1 LANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDLVAGSILKIGVKAGQLKFDIA..... 810
NTDB id 289 SPD RS10700 WP 001109677.1 LANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDLVAGSILKIGVKAGQLKFDIA..... 810
NTDB id 287 SP RS11210 WP 001109712.1 LANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDLVAGSTLKIGVKAGQLKFDIA..... 810
NTDB id 377 SMU RS09275 WP 002262344.1 LALAGYDAEMGARPLRRTLQTEVEDKLAELILSGKLASGQTLKIGIQKEKLKFDIV..... 813
NTDB id 297 STER RS00545 WP 011680614.1 LAKDGYDIEMGARPLRRTIQTQVEDKLSELLLGGQVVSGQTLKIGCSKDKLTFTVV..... 816
NTDB id 329 STU RS10020 WP 011225298.1 LAKDGYDIEMGARPLRRTIQTQVEDKLSELLLGGQVVSGQTLKIGCSKDKLTFTVV..... 816
consensus
non conserved
similar
≥ 50% conserved