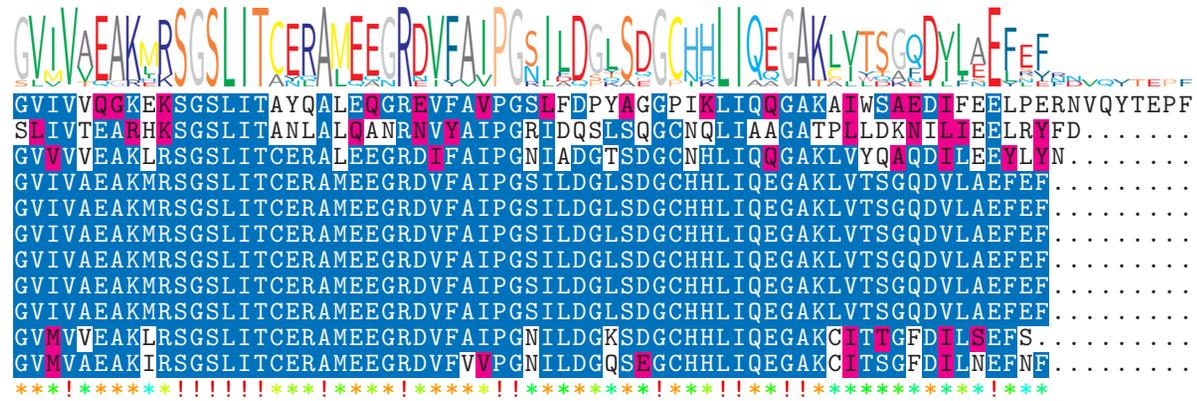


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NTDB id 114 BSU 16110 NP 389493.1	GVIVVQCKEKSGSLITAYQALEQGREVFAVPGSLFD ^X DPYAGGPIKLIQQGAKAIWSAEDIFEELPERNVQYTEPF	297
NTDB id 626 LCA RS04960 WP 011374694.1	SLIVTEARHKSGSLITANLALQANRNVYAIPGRIDQSLSQGCNQLIAAGATPLLDKNILTEELRYFD.....	288
NTDB id 599 KW2 RS05940 WP 021037268.1	GVVVVEAKLRSGSLITCERALEEGRDIFAIPGNIADGTS ^X DCNHLIQGAKLVYQAQDILEEYLYN.....	282
NTDB id 525 SMSK321 RS06295 WP 000705298.1	GVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLAEF ^X EF.....	282
NTDB id 267 KZH43 RS05590 WP 000705306.1	GVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLAEF ^X EF.....	282
NTDB id 226 SPD RS05990 WP 000705306.1	GVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLAEF ^X EF.....	282
NTDB id 192 SPR RS05715 WP 000705306.1	GVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLAEF ^X EF.....	282
NTDB id 157 SP RS06205 WP 000705306.1	GVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLAEF ^X EF.....	282
NTDB id 497 SM12261 RS05565 WP 000705318.1	GVIVAEAKMRSGSLITCERAMEEGRDVFAIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLAEF ^X EF.....	282
NTDB id 223282 A6J87 RS09950 WP 080611997.1	GVVVVEAKLRSGSLITCERAMEEGRDVFAIPGNILDGKSDGCHHLIQEGAKCIT ^X TGFDILSEFS.....	279
NTDB id 407 SMU RS04605 WP 002262865.1	GVVVVEAKLRSGSLITCERAMEEGRDVFAIPGNILDGKSDGCHHLIQEGAKCIT ^X TGFDILNEFNF.....	280
consensus	***!*****!!!!!!***!*****!*****!*****!*****!*****!*****!*****!*****!*****	

- X non conserved
- X similar
- X ≥ 50% conserved