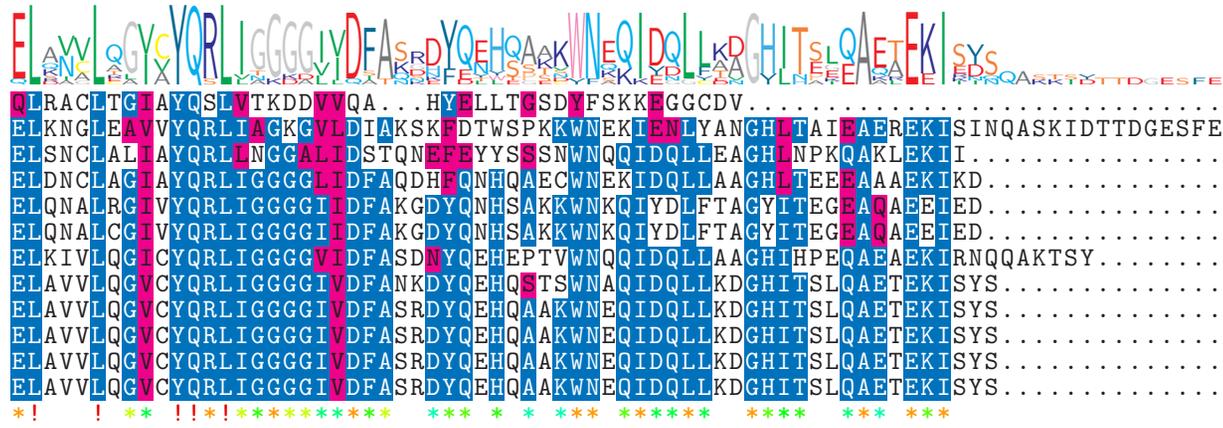


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NTDB id 617 LCA RS06480 WP 011375001.1	QLRACLTGTAYQSLVTKDDVVQA...HYELLTGSDFYFSKKEGGCDV.....	296
NTDB id 586 KW2 RS10855 WP 011836048.1	ELKNGLEAVVYQRLIAGKGVLDIAKSKFDTWSPKKWNEKIENTLYANGHLTAIEEAEREKISINQASKIDTTDGESFE	326
NTDB id 137310 SD89 RS00650 WP 002993471.1	ELSNCLALIAVYQRLINGGALIDSTQNEFEYYSSTSNWNQQIDQLLEAGHLNPKQAKLEKII.....	312
NTDB id 584 DLJ51 RS00890 WP 019777327.1	ELDNCLAGTAYQRLIGGGGLIDFAQDHFQNHQAECWNEKIDQLLAAGHLTEEEA AAAEKIKD.....	315
NTDB id 385 SMU RS09035 WP 002263435.1	ELQNALRGIVYQRLIGGGGIDFAKGDYQNHS AKKWNKQIYDLFTA GYITEGEAQAE EIED.....	313
NTDB id 394 FSA28 RS09405 WP 002291652.1	ELQNALCGIVYQRLIGGGGIDFAKGDYQNHS AKKWNKQIYDLFTA GYITEGEAQAE EIED.....	313
NTDB id 429 SGO RS09420 WP 012130942.1	ELKIVLQGITCYQRLIGGGGVIDFASDNYQEHEPTVWNQQIDQLLAAGHIHPEQAEAEKIRNQQAKTSY.....	319
NTDB id 500 SM12261 RS00945 WP 000249545.1	ELAVVLQGVICYQRLIGGGGIVDFANKDYQEHQSTSWNAQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 270 KZH43 RS09265 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQA AKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 229 SPD RS09895 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQA AKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 195 SPR RS09430 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQA AKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
NTDB id 160 SP RS10380 WP 000249564.1	ELAVVLQGVICYQRLIGGGGIVDFASRDYQEHQA AKWNEQIDQLLKDGHITSLQAETEKISYS.....	313
consensus	*! ! ** !*!*!***** ** * * ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** **	

non conserved
 similar
 >= 50% conserved