

NTDB id 587 KW2 RS10850 WP 042211510.1	MQTDLRLLRPKGKKLALIKQAKLIQLMGNLLNNGFNLGEVINFLALSKEVEKVFVKM	59
NTDB id 411385 GQY29 RS04095 WP 159340986.1	MPEKISKTIIRPTGISSWKVWLNKDVSRLRGISKGKKLKISQQVKVTQLFKQLLKAGFTLTETIIVAFLEERSHLLKETSLSLM	80
NTDB id 395 FSA28 RS09400 WP 257638827.1	MTFLLQDISLKEKHKKQKKLKIKKQRKVVIQLFNLFSGFNLTEIIDFLERSHLLAEIYTCKM	62
NTDB id 386 SMU RS09030 WP 255262721.1	MTFLLQDISLKEKHKKQKKLKIKKQRKVVIQLFNLFSGFNLTEIIDFLERSHLLAEIYTCKM	62
NTDB id 430 SGO RS09415 WP 012130941.1	MISFLQDISILSKQKKKLGTSKQKQVIELFNLFSSGFHLAETVDFLGRSALLKKNYKQVM	63
NTDB id 271 KZH43 RS09260 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 230 SPD RS09890 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 196 SPR RS09425 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 161 SP RS10375 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 529 SMSK321 RS09235 WP 080550783.1	MDISQVFRLRRKKLATAKQKNIITLFNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
NTDB id 501 SM12261 RS00950 WP 004238896.1	MDISQVFRLRRKKLATAKQKNIITLFNLFSSGFHLVETISFLDRSALLDKQCVTQM	57
consensus	*!*****!!*****!*!!!!*!!!*!***!****!	

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consensus



☒ non conserved
☒ similar
☒ ≥ 50% conserved