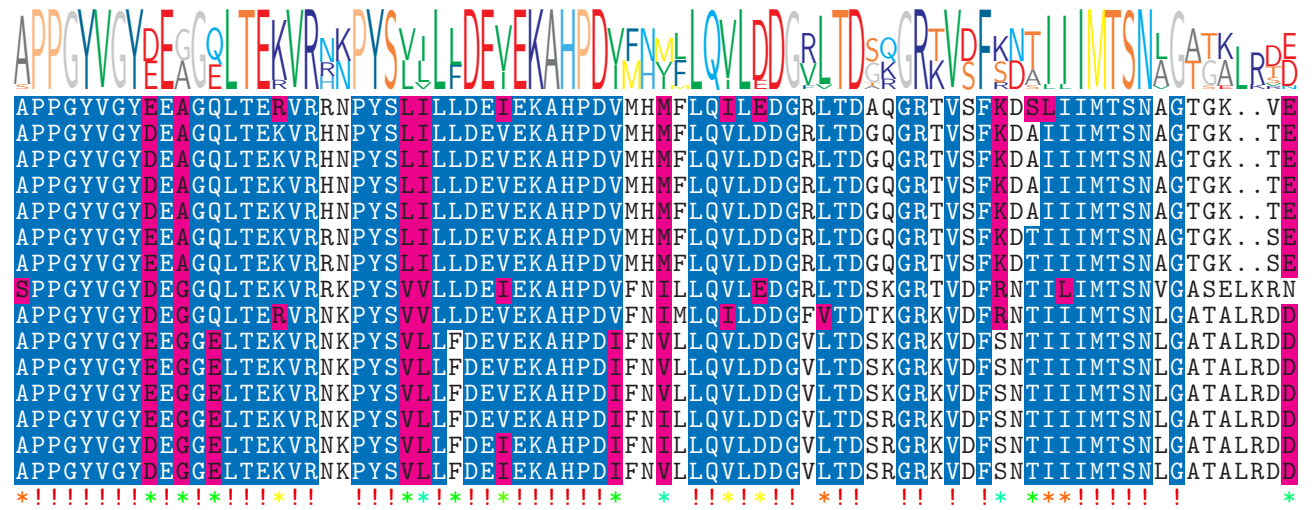


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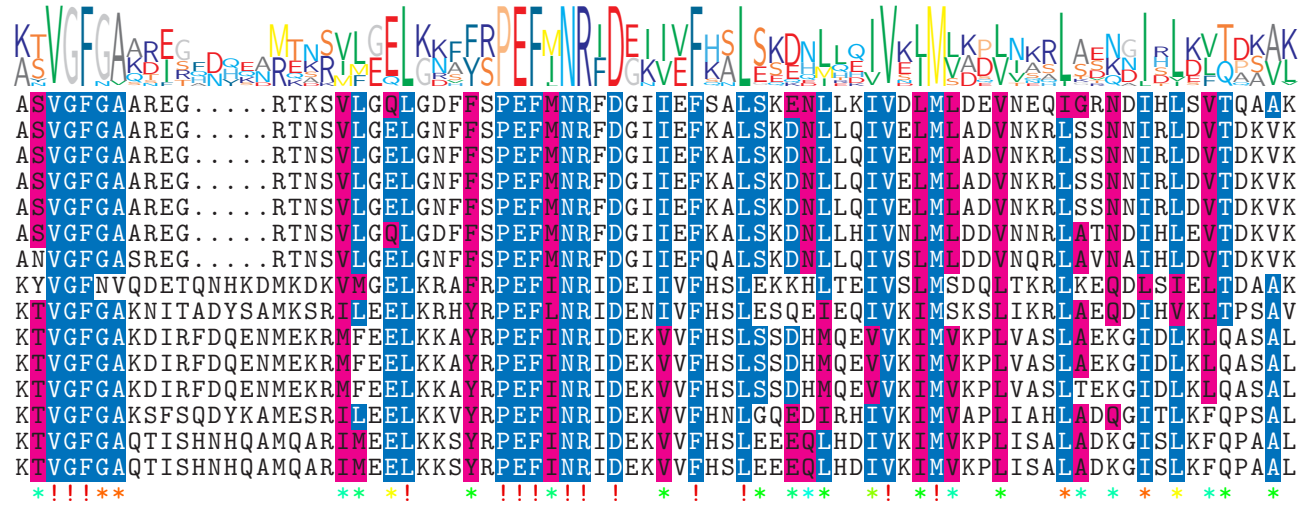
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 NTDB id 281 KZH43 RS03550 WP 000882517.1
 NTDB id 245 SPD RS03860 WP 000882517.1
 NTDB id 243 SPR RS03635 WP 000882517.1
 NTDB id 241 SP RS04015 WP 000882523.1
 NTDB id 409447 GPZ88 RS07130 WP 074563274.1
 NTDB id 378 SMU RS02690 WP 002263569.1
 NTDB id 85 BSU 00860 NP 387967.1
 NTDB id 610 V4T04 RS10165 WP 012897346.1
 NTDB id 290 KZH43 RS10025 WP 001109677.1
 NTDB id 289 SPD RS10700 WP 001109677.1
 NTDB id 287 SP RS11210 WP 001109712.1
 NTDB id 377 SMU RS09275 WP 002262344.1
 NTDB id 297 STER RS00545 WP 011680614.1
 NTDB id 329 STU RS10020 WP 011225298.1
 consensus



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NTDB id 601 KW2 RS02615 WP 021036805.1
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 NTDB id 245 SPD RS03860 WP 000882517.1
 NTDB id 243 SPR RS03635 WP 000882517.1
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 NTDB id 287 SP RS11210 WP 001109712.1
 NTDB id 377 SMU RS09275 WP 002262344.1
 NTDB id 297 STER RS00545 WP 011680614.1
 NTDB id 329 STU RS10020 WP 011225298.1
 consensus



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NTDB id 601 KW2 RS02615 WP 021036805.1	EKLV	DLGY	NPAMGARPLRRTIQ	ENTIEDS	IADFY	TEHPEYKQL	.	VADLIDDKIV	ISNQTQ	..ETAETTTDEEVPAE	748		
NTDB id 281 KZH43 RS03550 WP 000882517.1	EKLV	DLGYDPKMGARPLRRTIQDY	IEDTITDYYLENPSEKDL	.	KAVMTSKGNI	QIKSAKKA	AEVKSSEKEK			752		
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NTDB id 243 SPR RS03635 WP 000882517.1	EKLV	DLGYDPKMGARPLRRTIQDY	IEDTITDYYLENPSEKDL	.	KAVMTSKGNI	QIKSAKKA	AEVKSSEKEK			752		
NTDB id 241 SP RS04015 WP 000882523.1	EKLV	DLGYDPKMGARPLRRTIQDY	IEDTITDYYLENPSEKDL	.	KAVMTSKGNI	QIKSAKKA	AEVKSSEKEK			752		
NTDB id 409447 GPZ88 RS07130 WP 074563274.1	EKLV	DLGYDPKMGARPLRRTIQDHI	IEDAITDFYLEHPTEKDL	.	KAVMTSNGKI	IIKSAKK	VEKEETVSAE			761		
NTDB id 378 SMU RS02690 WP 002263569.1	EKLV	DLGYDPKMGARPLRRTIQEHI	IEDAITDFYLEHPEQKEL	.	KAIMTSNGKI	VIKASNK	VETVTKATSD			753		
NTDB id 85 BSU 00860 NP 387967.1	AKV	AEEGVDLEYGARPLRRAIQKHV	EDRLSEELLRGNIHKGQHTVLD	V	EDG	GEFVVKT	TAKTN			810		
NTDB id 610 V4T04 RS10165 WP 012897346.1	KL	IAEVGF	DPEYGARPLR	KALQKEVEDL	LSEQLLS	SGEIKAGNHV	SIGAS	NKKTKIAQIV		816		
NTDB id 290 KZH43 RS10025 WP 001109677.1	KLL	ANQGYDPEM	GARPLRRTLQTEVEDK	LAELL	LKGD	LVAGSILKIGVK	AGQLKFDIA			810		
NTDB id 289 SPD RS10700 WP 001109677.1	KLL	ANQGYDPEM	GARPLRRTLQTEVEDK	LAELL	LKGD	LVAGSILKIGVK	AGQLKFDIA			810		
NTDB id 287 SP RS11210 WP 001109712.1	KLL	ANQGYDPEM	GARPLRRTLQTEVEDK	LAELL	LKGD	LVAGSTL	KIGVKAGQLKFDIA			810		
NTDB id 377 SMU RS09275 WP 002262344.1	KHL	ALAGYDAEM	GARPLRRTLQTEVEDK	LAEL	LSGKL	ASGQTLKIGIQKEK	LKFDIV			813		
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NTDB id 329 STU RS10020 WP 011225298.1	KHL	LAKDGYDIE	MGARPLRRTIQTQVEDK	LSELL	GGQV	VSGQTLKIGCSKDKL	TFTVV			816		
consensus	*	*	!***	*!!!!!!***!	*!!	***	*		*	*	*	*	*

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