

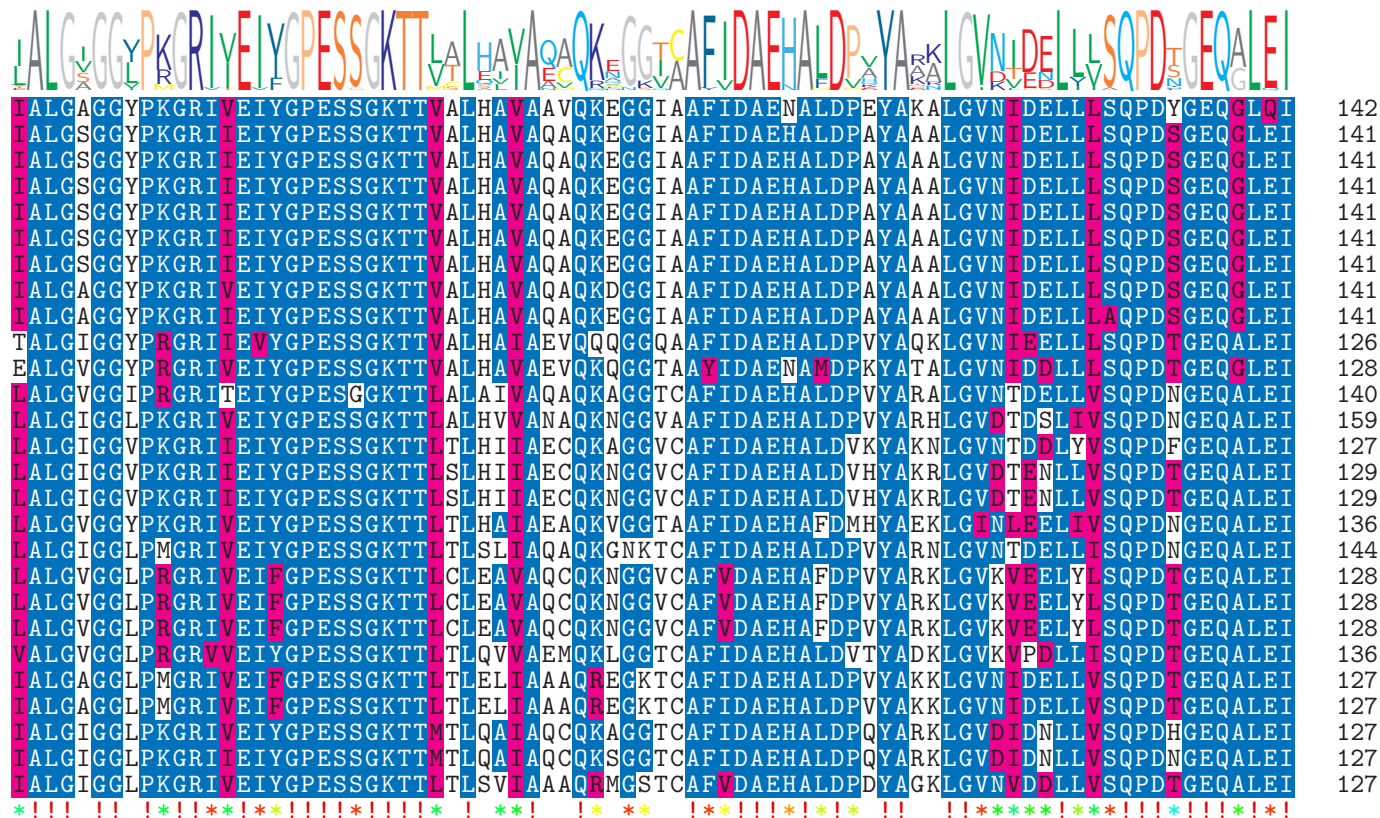
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consensus		***	***	*	***!	***!	*****	*	***!	***!							

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NTDB id 600 KW2 RS01765 WP 011675413.1
NTDB id 269 KZH43 RS08665 WP 001085462.1
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NTDB id 194 SPR RS08825 WP 001085462.1
NTDB id 159 SP RS09750 WP 001085462.1
NTDB id 527 SMSK321 RS11065 WP 001085454.1
NTDB id 499 SM12261 RS08315 WP 078228268.1
NTDB id 408 SMU RS09530 WP 002262392.1
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consensus

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NTDB id 1389 A4U84 RS07560 WP 021112712.1	F I K K S G A W F S Y D G T K I G Q G K N N A V K W L K E N P E I A D K L E Q E I R N T L A S N P N S V L K A T A S D S N D E S . D V E E F S E E L	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	I I N K S G A W S Y N G A K I G Q G K D N V R V W L K E N P E I S D E I D A K I R A L N G V E M H I T E G T Q D E T D G E R P E E	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	I I N K S G A W S Y N G A K I G Q G K D N V R V W L K E N P E I S D E I D A K I R A L N G V E M H I T E G T Q D E T D G E R P E E	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	I I N K S G A W S Y N G A K I G Q G K D N V R V W L K E N P E I S D E I D A K I R A L N G V E M H I T E G T Q D E T D G E R P E E	348
NTDB id 1427 RS RS02760 WP 011000510.1	V V E K S G A W S Y N G E R I G Q G R D N C R E F L R E N A E L A R E I E N K V R E H L G V T P M G A V T L A . E E V E E D	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	M V E K S G A W S Y N G D K I G Q G K A N A C K Y L K E N P E I A K T L D K K L R E M L L N P E N M Q L I A E T S S A A D D V . E F G A V P E E F	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	M V E K S G A W S Y N G D K I G Q G K A N A C K Y L K E N P E I A K T L D K K L R E M L L N P E N M Q L I A E T S S A A D D V . E F G A V P E E F	354
NTDB id 1022 ACIAD RS06390 WP 004925639.1	I V Q K A G A W S Y Q G N K I G Q G K N N V I R Y F E E N N Q I A E E I E R N I R E Q L L T T G T N G A V Q . I . E D E E E P . D L L . . L E S	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	I V Q K A G A W S Y Q G N K I G Q G K N N V I R Y F E E N N Q I A E E I E R N I R E Q L L T T G T N G A V Q . I . E D E E E P . D L L . . L E S	349
NTDB id 1205 PSJM300 05545 AFN77183.1	L V E K S G A W Y A Y K G S K I G Q G K A N A A K Y L E E N P E I G Q E I E Q Q I R A K L L V T S A N T K A T P V A E D L A D A . D I	347
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NTDB	id	1205	PSJM300	05545	AFN	77183.1		347

consensus

-  non conserved
-  similar
-  >50% conserved