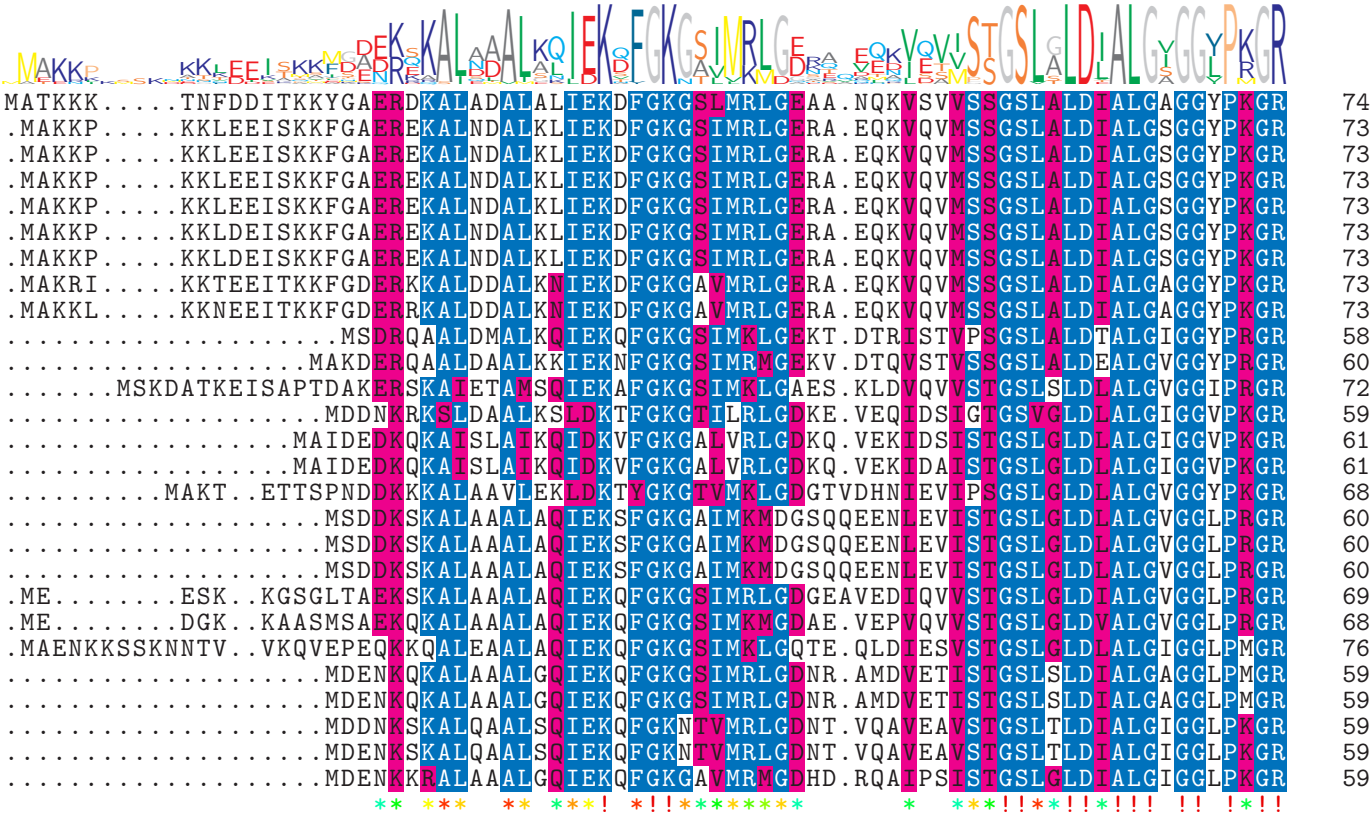


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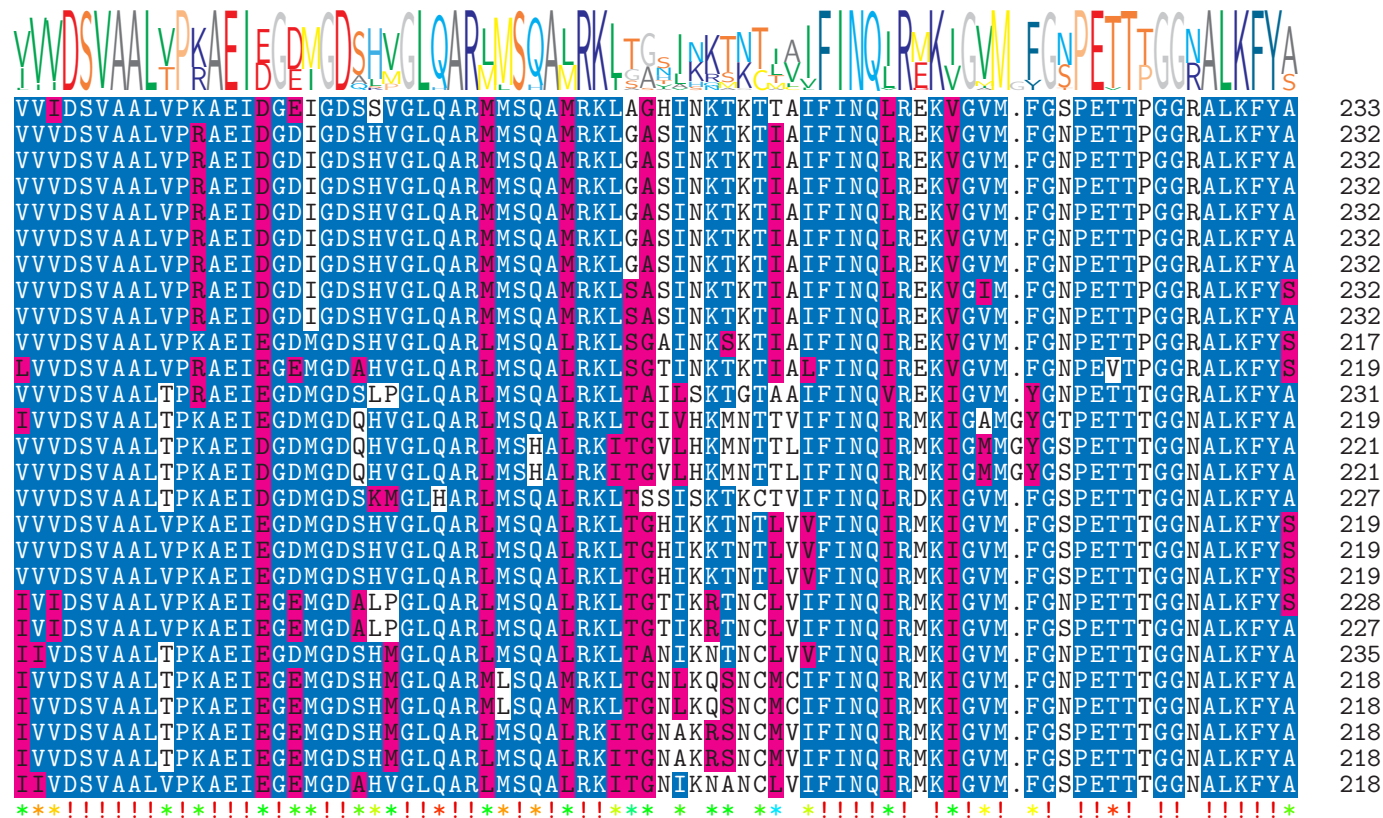
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NTDB id 453 Spy49 1753c ACI62003.1	SVRLDVRGTTQIKGTG.DQKDSS	IGKETKIKVVKNKVAPPFKVAEVEIMYEGEGISRTGELVKIASDLDI	ITKKAGAWFSYN	311
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	SVRLDIRRIGQ....V.KEG	DEIIGSETKVVVKNKMAPPFKEAIFQILYGKGTNQLGELVDLAVQQ	NIVQKAGAWYSYQ	293
NTDB id 1205 PSJM300 05545 AFN77183.1	SVRLDIRRTGA....V.KEG	DEVVGSETRVKVVKNKVAPPFRQAEFQILYGKGIYRNGEIIDLGVQQ	GLVEKSGAWAYK	293
consensus	*!!**!* *	* ** *****!*****!*****!*****	*****!!***** ***** *	*** ! !*****!

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NTDB id 600 KW2 RS01765 WP 011675413.1	DEKIGQGA	EKAKNYLKEHQDV	DEIDHKVRAAHG	LDDSEVAETEEETT	ASKTKAKAKKEEK	AVETEEIELELED		387		
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	DKKLGGG	RENAKALLKEDK	ALADEITLKI	KESIGSNEEIM	PLPD	..EPL	..EEME		347	
NTDB id 1334 RAOC RS04870 WP 004918236.1	DTKLGGG	RDITVKELIKDN	PELAEELEAKI	REKLQAK					343	
NTDB id 1130 NGFG RS03960 WP 003688695.1	GAKIGQG	KDNVRVWLKENPEIS	DEIDAKIRALNG	VEMHIT	EGTQD	ETD	..GER	PEE	348	
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	GTKIGQG	KNNAVKWLKENPEI	ADKLEQEI	RNTLASNPNSV	LKATAS	DSN	..DESD	VEEFSEEEEL	372	
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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved