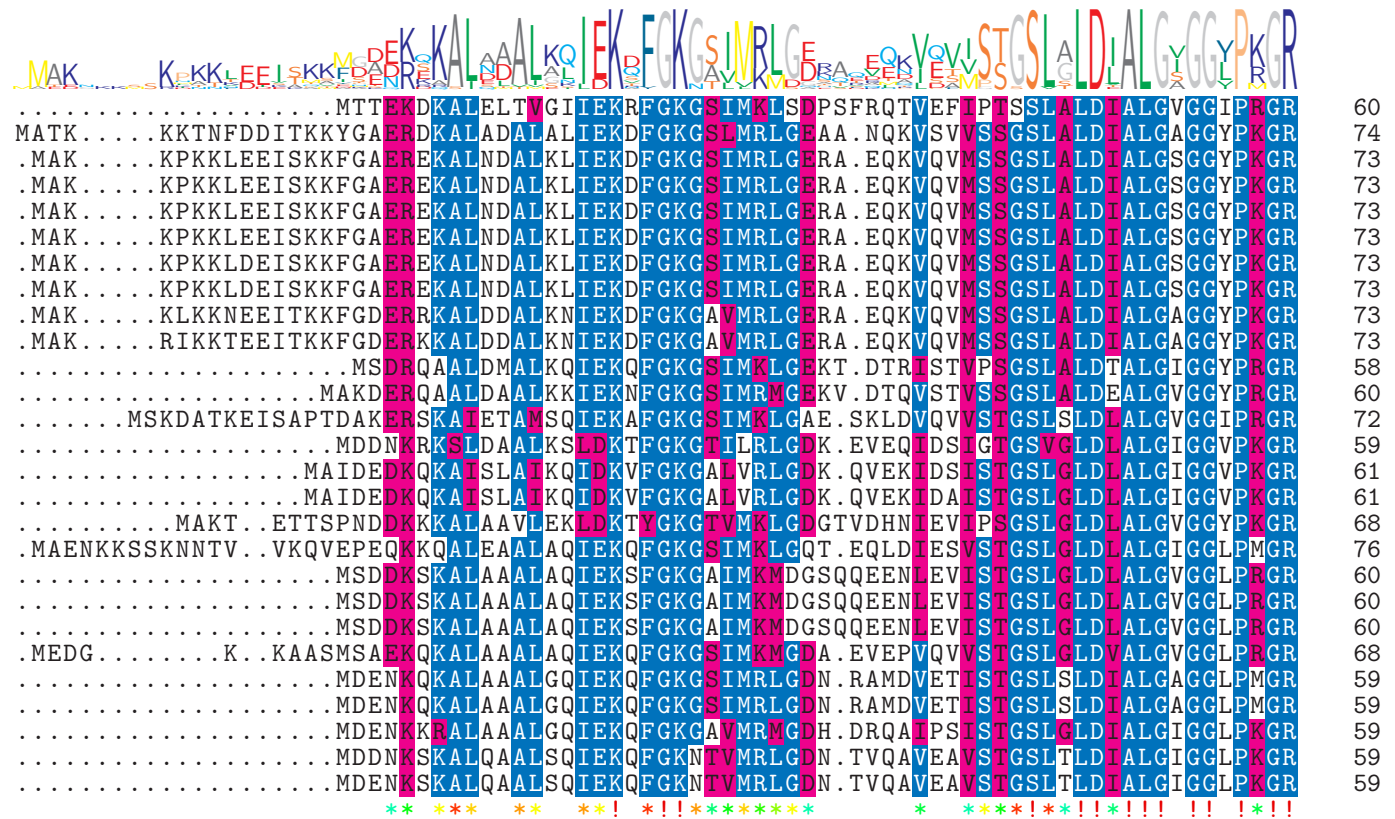


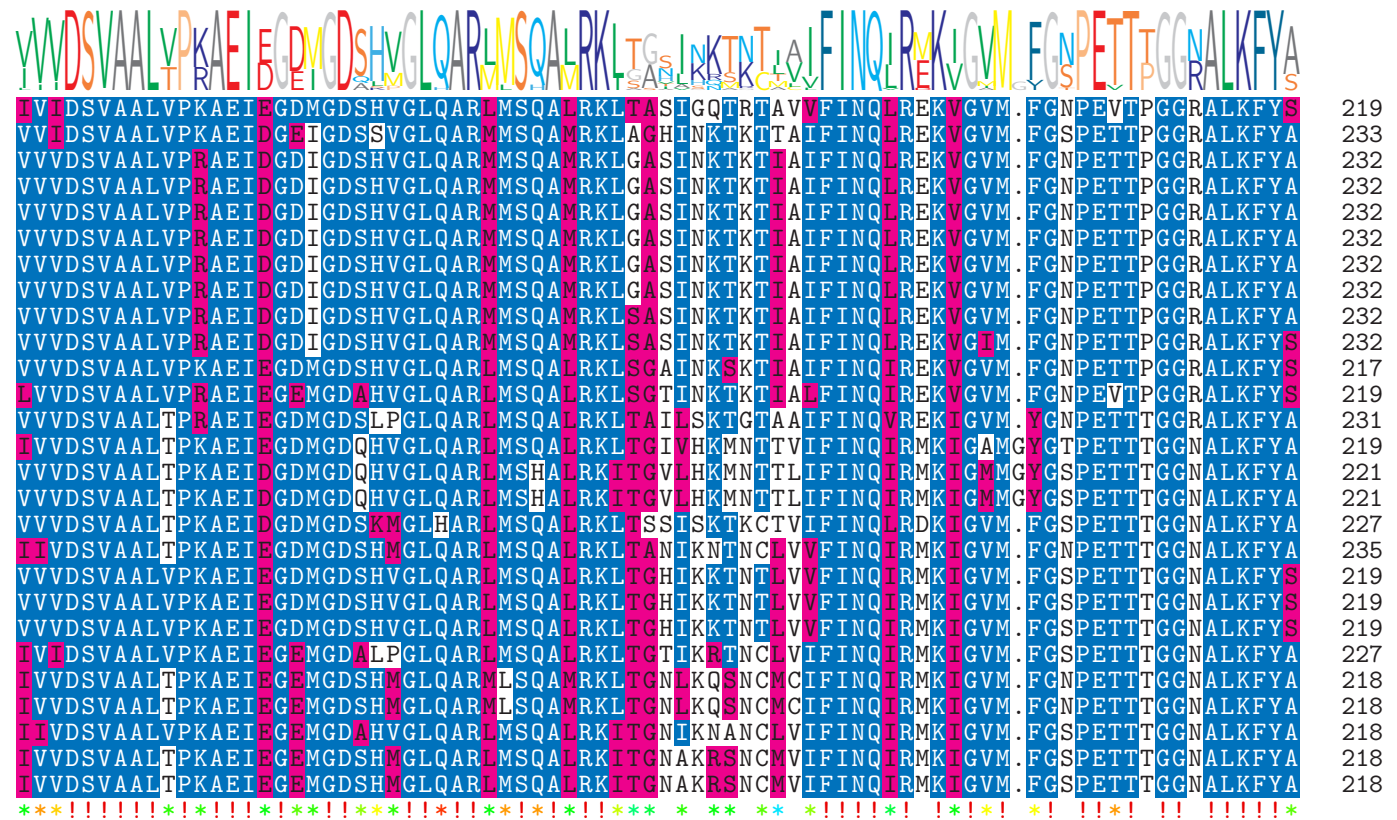
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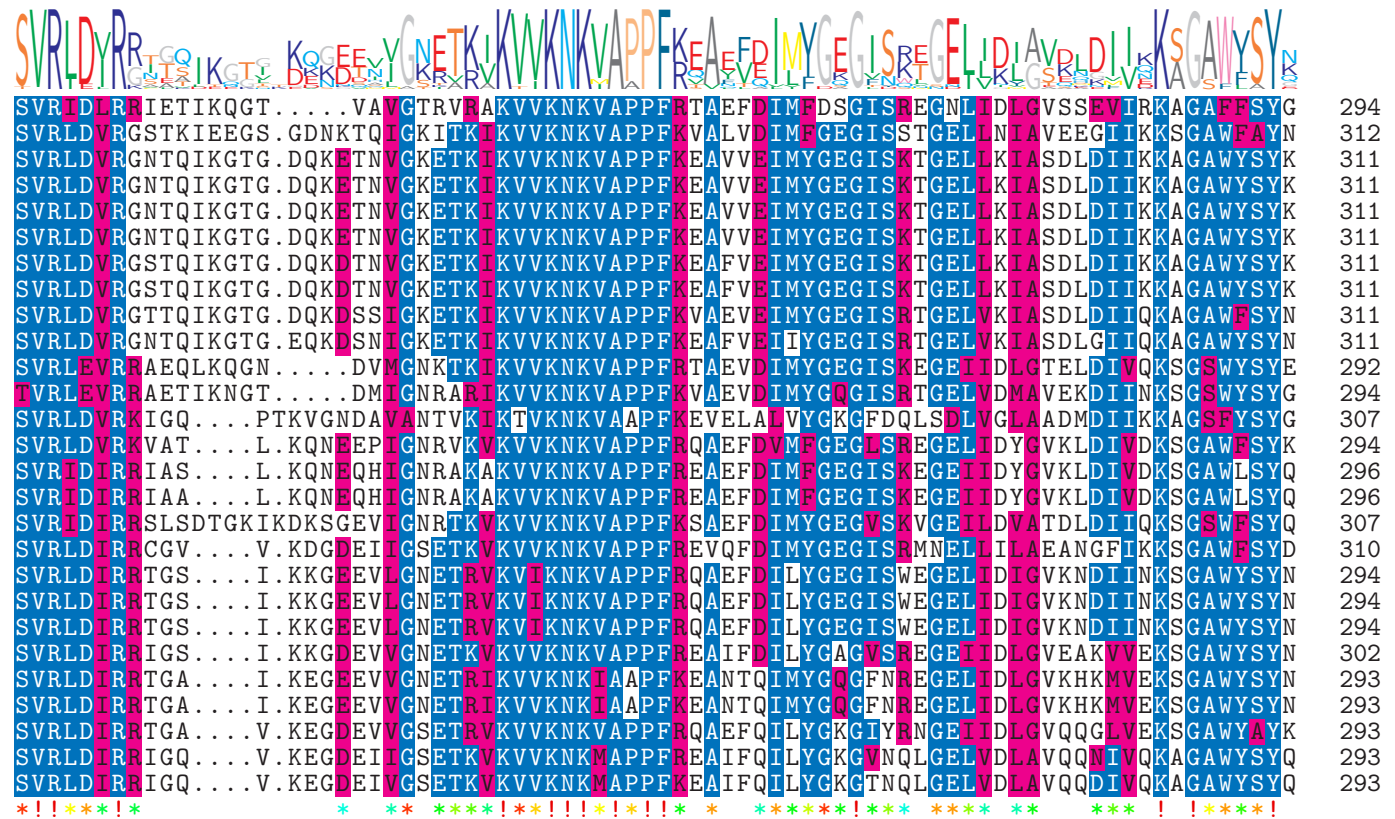
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consensus	**	!	*	*	*	!	!	*	*	!	!	!	!	!	*	*	*	*	!	!	*	*	*	!	!	*	*	!	!	*	*	*	!	!	*	*	*	!	!	*	*	*																																								

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consensus

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Sequence	Score
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DE K I G Q G S E N A K K Y L A E H P E I F D E I D K Q V R S K F G L I D G E E V S E Q D T E N K K D E . . . P K K E E A . V N E E V P L D L G D E L E I E	385
DE K I G Q G S E N A K K Y L A E H P E I F D E I D K Q V R S K F G L I D G E E V S E Q D T E N K K D E . . . P K K E E A . V N E E V P L D L G D E L E I E	385
DE K I G Q G S E N A K K Y L A E H P E I F D E I D K Q V R S K F G L I D G E E V S E Q D T E N K K D E . . . P K K E E A . V N E E V P L D L G D E L E I E	385
DE K I G Q G S E N A K K Y L A E H P E I F D E I D K Q V R S K F G L I D G E E V S E Q D T E N K K D E . . . P K K E E A . V N E E V P L D L G D E L E I E	385
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GE K I G Q G S E N A K R Y L A D H P Q L F D E I D R K V R V K F G L L E E S E E S A M A V A S E E T D D L A L D L D N G I E I E	377
GE K I G Q G S E N A K K F L A D N P E I F D D I D H K V R V Q Y G L I E E D N T E E K Q S S K E K E T D E K . . A D K N L V L E L D D T I E L E	382
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-  non conserved
-  similar
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