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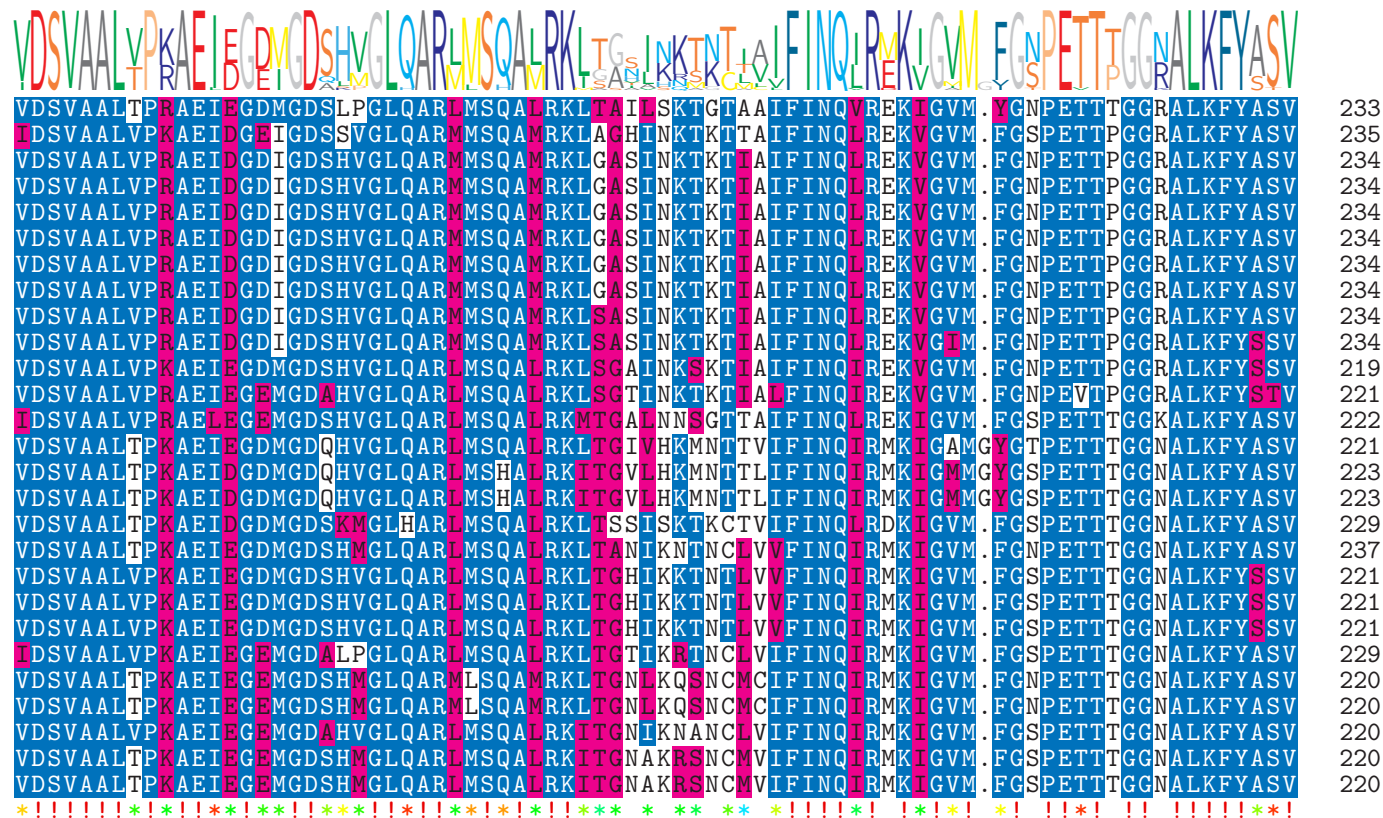
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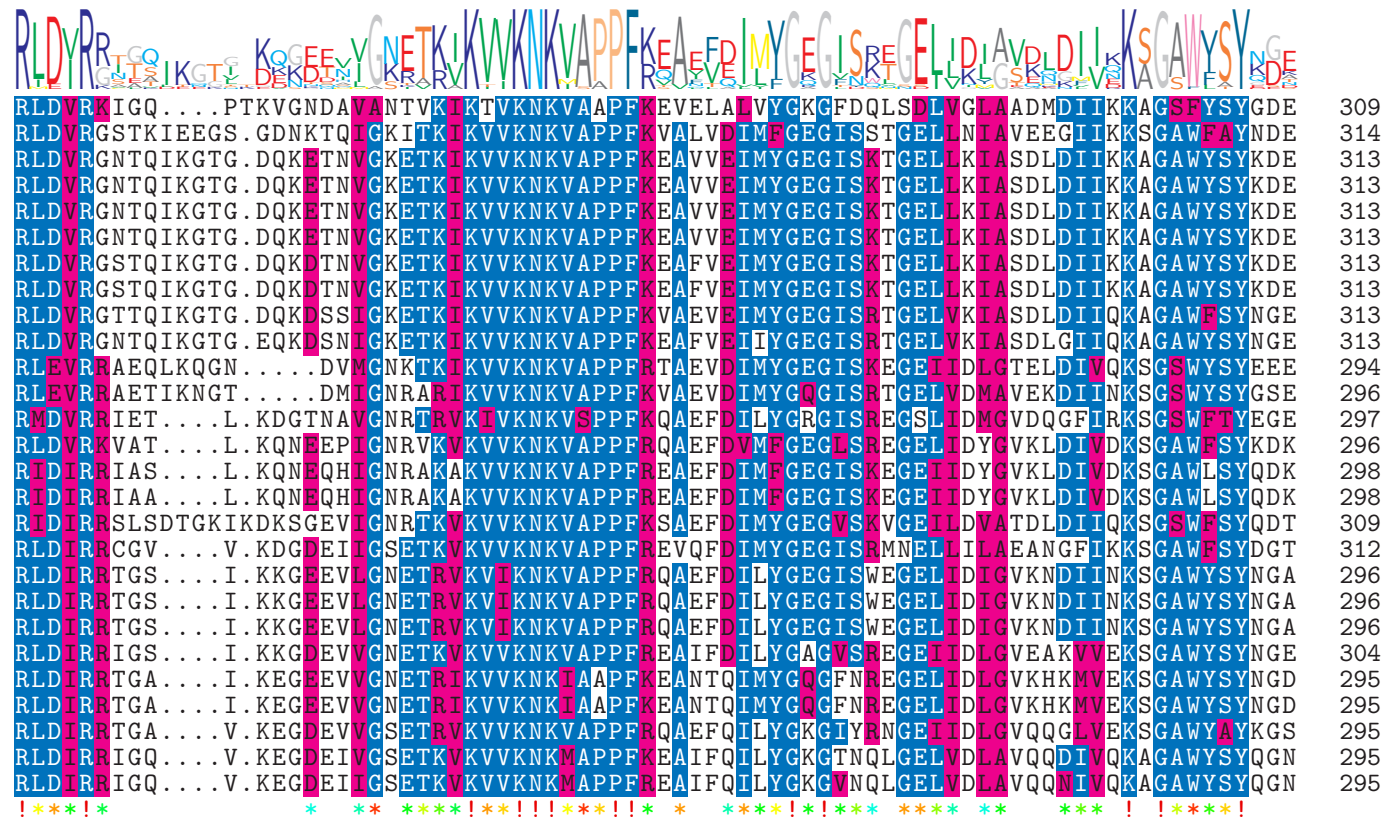
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NTDB id 1205 PSJM300 05545 AFN77183.1
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NTDB id 1022 ACIAD RS06390 WP 004925639.1
consensus

Sequence	Score
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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 I	386
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R L G Q G G R E N A K Q F L K E N K D I M L M I Q E Q I R E H Y G L D N N G V V Q Q A E E T Q E E L E F E . E	348
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K L G Q G G R E N S K A F L K E N P E I A D E I T K A I Q N S M G I E G M I S G S E D D E G E E	343
K L G Q G G R E N A K A L L K E D K A L A D E I T L K I K E S I G S N E E I M P L P D E P L E E M E	347
K L G Q G G R E N A K A L L K E D K A L A D E I T L K I K E S I G S N E E I M P L P D E P L E E M E	347
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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 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 E L	372
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consensus

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