

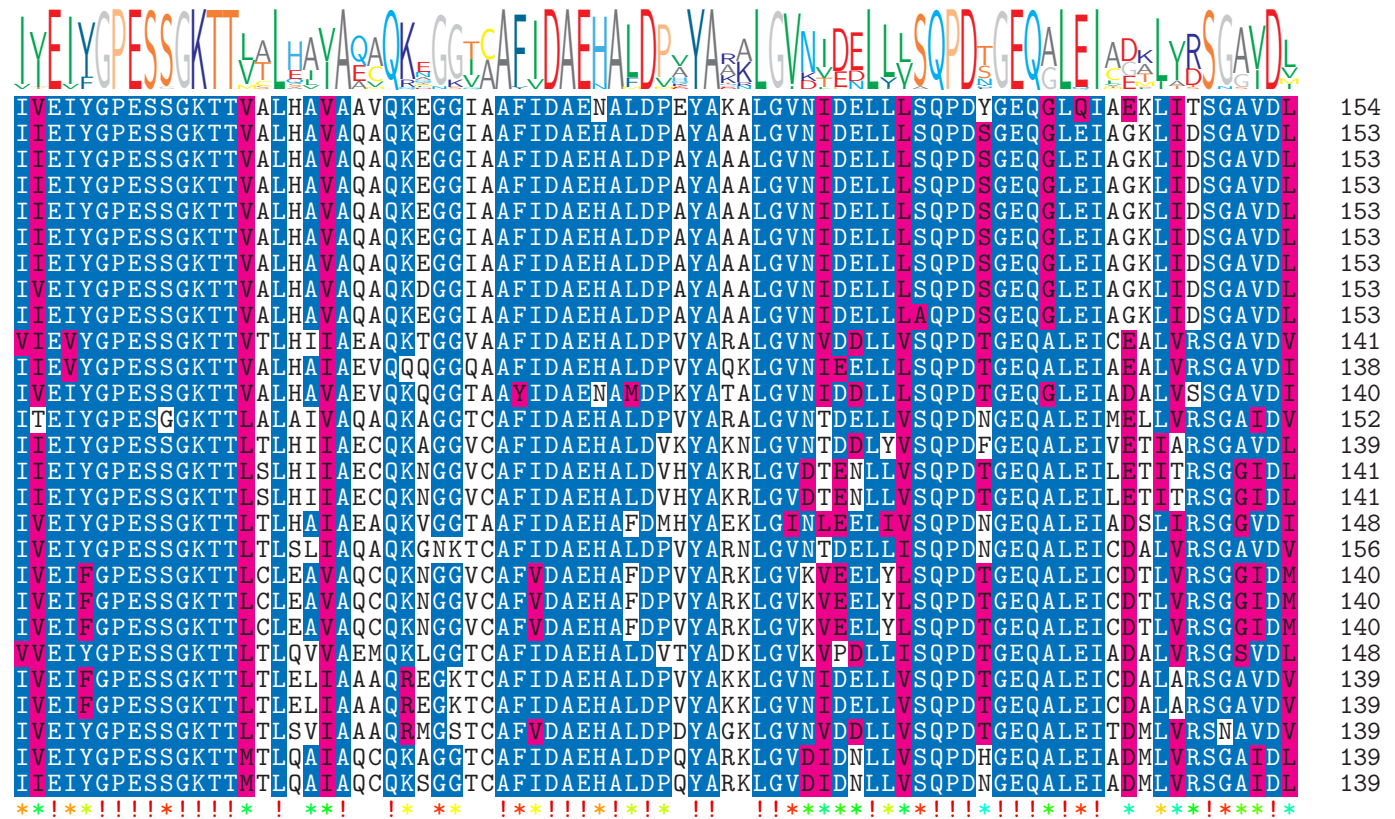
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NTDB id 35603 DHBDCA RS01595 WP 015042386.1	MATPDKLKALDIALS	QIEKQFGKGA	AIMKLG	EASDRMAVET	ISTGSLALD	L	ALGVGGVPR	GR	61
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NTDB id 625 LCA RS02525 WP 011374196.1	MAKDERQAALDAALKK	IEKNFGKGSIMRMGE	KVDTQVSTV	SSGSLALD	E	ALGVGGYPR	GR	60	
NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTD	AKERSKATETAMSQIEKA	FGKGSIMKLGAE	.SKLDVQVV	STGSLSLD	L	ALGVGGIPR	GR	72
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NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....K..KAASMSA	EKQKALAAALAQIEKQ	FGKGSIMKMGDA	.EVEPVQVV	STGSLGLD	V	ALGVGGLPR	GR	68
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	MDENKQKALAAALGQIEKQ	FGKGSIMRLGDN	.RAMDVETISTGSLSLD	I	ALGAGGLPM	GR	59		
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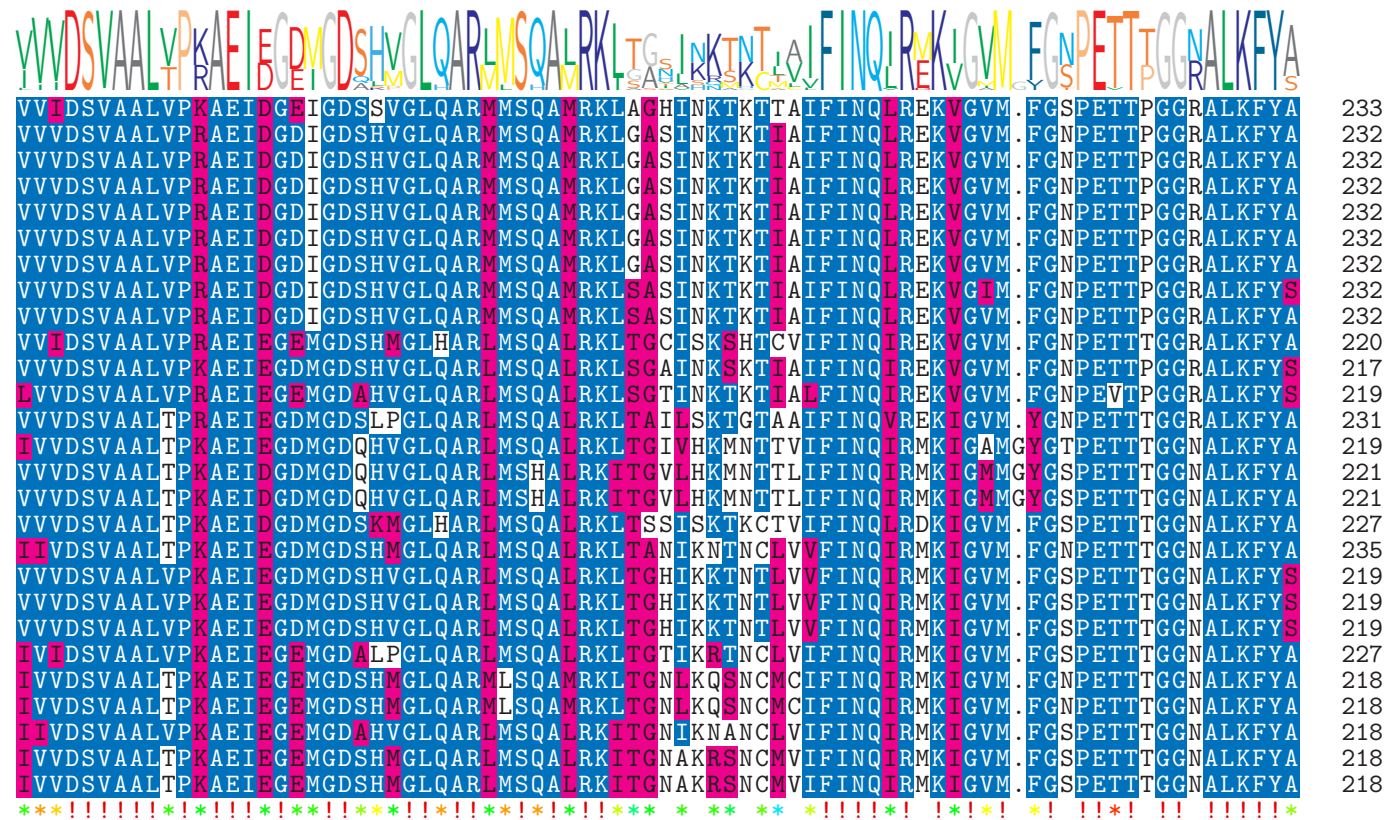
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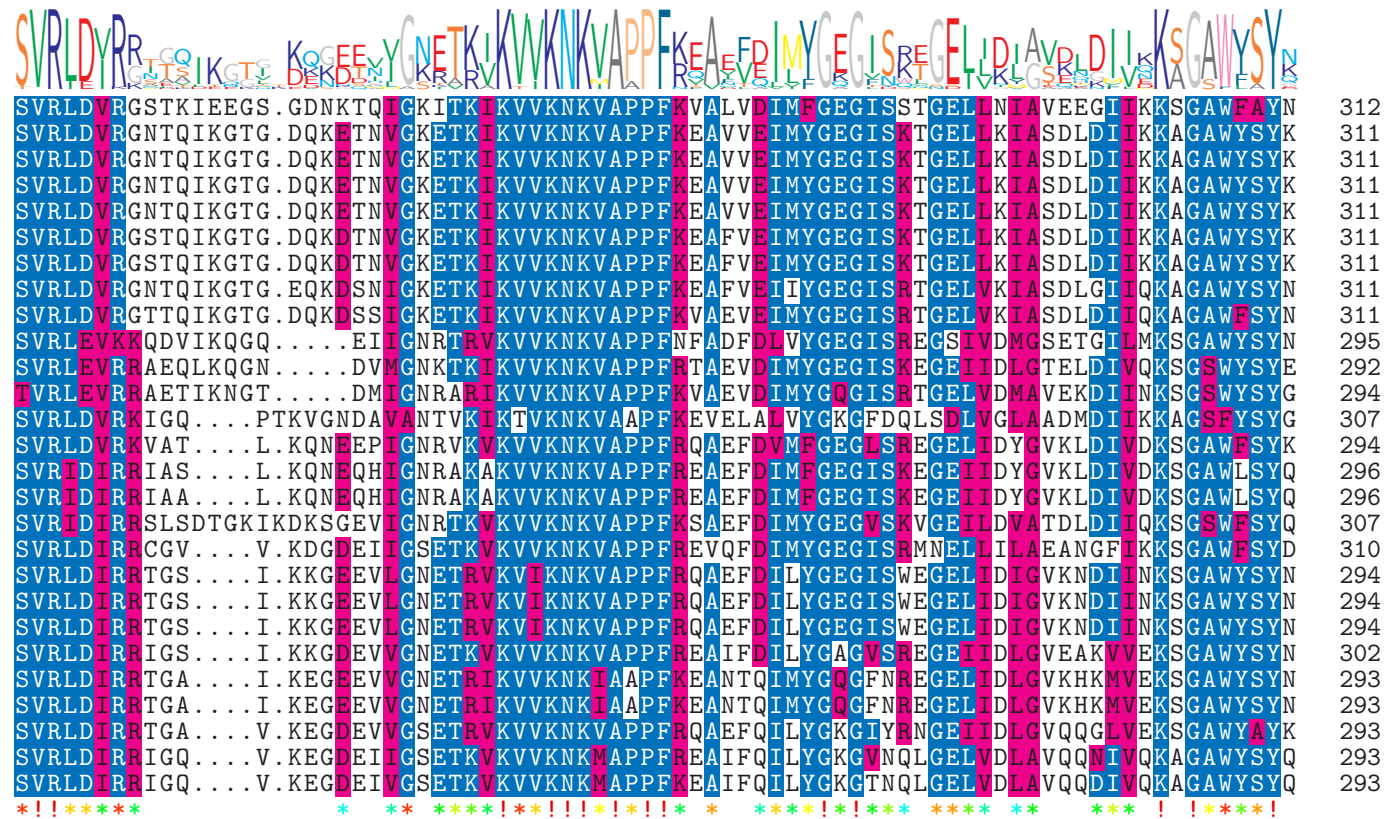
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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NTDB	id	194	SPR	RS08825	WP	001085462.1	DEKIGQGS	ENAKKYLAEHPEI	FDKQVRSKFGL	LIDGEEVSEQDTENKK...	DE....PKKEEA..VNEEVPLDLGDE	381
NTDB	id	159	SP	RS09750	WP	001085462.1	DEKIGQGS	ENAKKYLAEHPEI	FDKQVRSKFGL	LIDGEEVSEQDTENKK...	DE....PKKEEA..VNEEVPLDLGDE	381
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NTDB	id	1074	ABD1	RS10195	WP	000344167.1	GNKIGQCKN	NVIRYLEENPQI	STEIEAVIREQLLT	GTGN....GAVQ.I.EDE	EEPDLL..LE.....S.....	349
consensus							*** I *****	***	*****	*****	*****	

logo

NTDB id 600 KW2 RS01765 WP 011675413.1		387
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved