

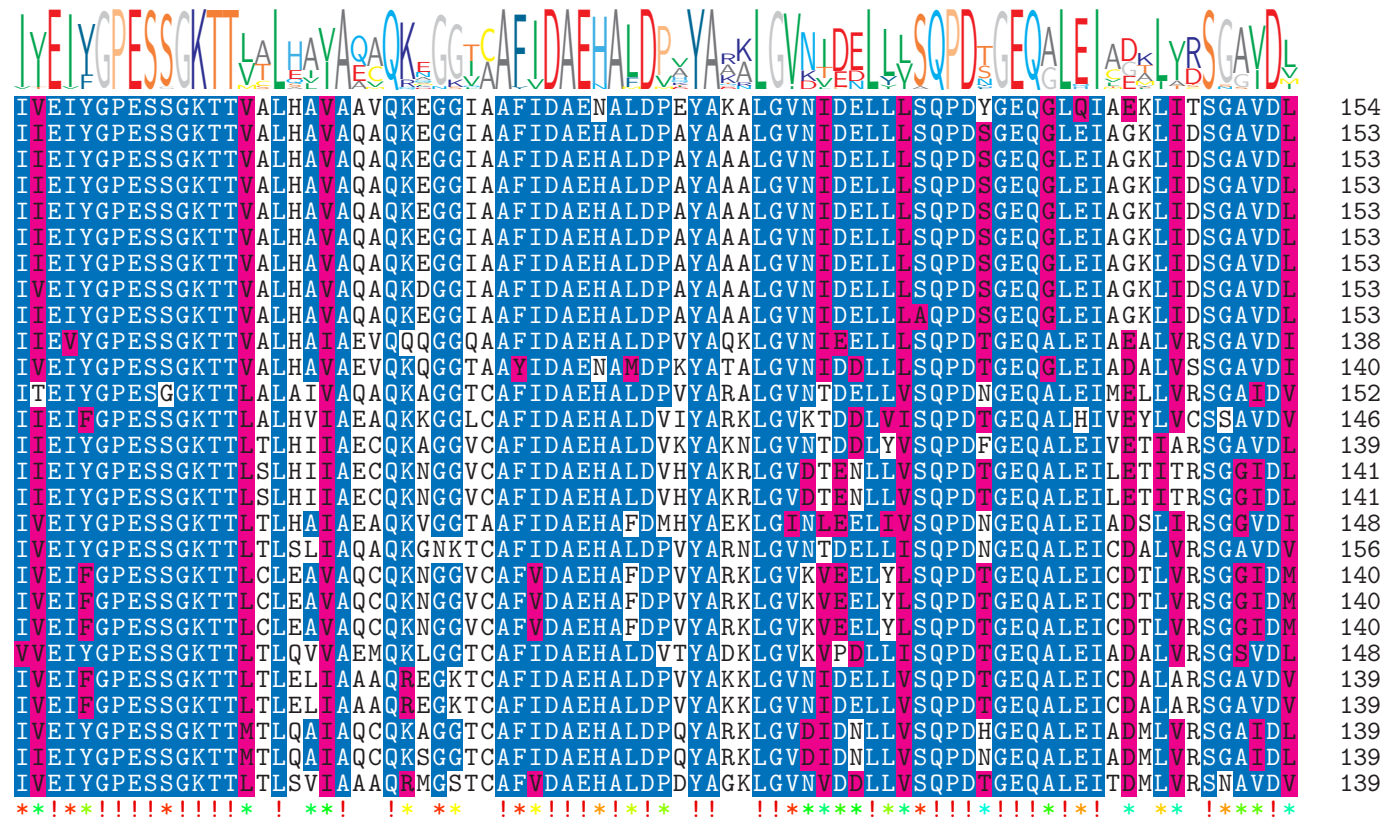
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NTDB id 93382 wHmt RS03190 WP 007302323.1	MA..HNPEERN	NDKQKALDNAISQIEK	AFGKGAIMKLKQN	.PVEKIDT	ISTGSTALD	S	ALGVGGL	PKGR	66
NTDB id 1245 Cj1673c YP 002345041.1	MDDN	KRKSLSLDAALKSLDK	TFGKGTILRLGDK	.EVEQIDS	IGTGSVGLD	L	ALGIGGV	PKGR	59
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NTDB id 1211 C694 RS00795 WP 000963128.1	MAIDE	DKQKATISLAIKQID	KVFGKGALVRLGDK	.QVEKIDS	ISTGSLGLD	L	ALGIGGV	PKGR	61
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NTDB id 1130 NGFG RS03960 WP 003688695.1	MSD	DKSKALAAALAQIEKS	FGKGAIMKMDGSQ	QEEENLEV	ISTGSLGLD	L	ALGVGGL	PRGR	60
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NTDB id 1123 OK783 RS03845 WP 003688695.1	MSD	DKSKALAAALAQIEKS	FGKGAIMKMDGSQ	QEEENLEV	ISTGSLGLD	L	ALGVGGL	PRGR	60
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....KKAASMSA	EKQKALAAALAQIEKQ	FGKGSIMKMGDA	.EVEPVQV	VSTGSLGLD	V	ALGVGGL	PRGR	68
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NTDB id 1205 PSJM300 05545 AFN77183.1	MDEN	KKRALAAALGQIEKQ	FGKGAVMRMGDH	.DRQAIPS	ISTGSLGLD	I	ALGIGGL	PKGR	59
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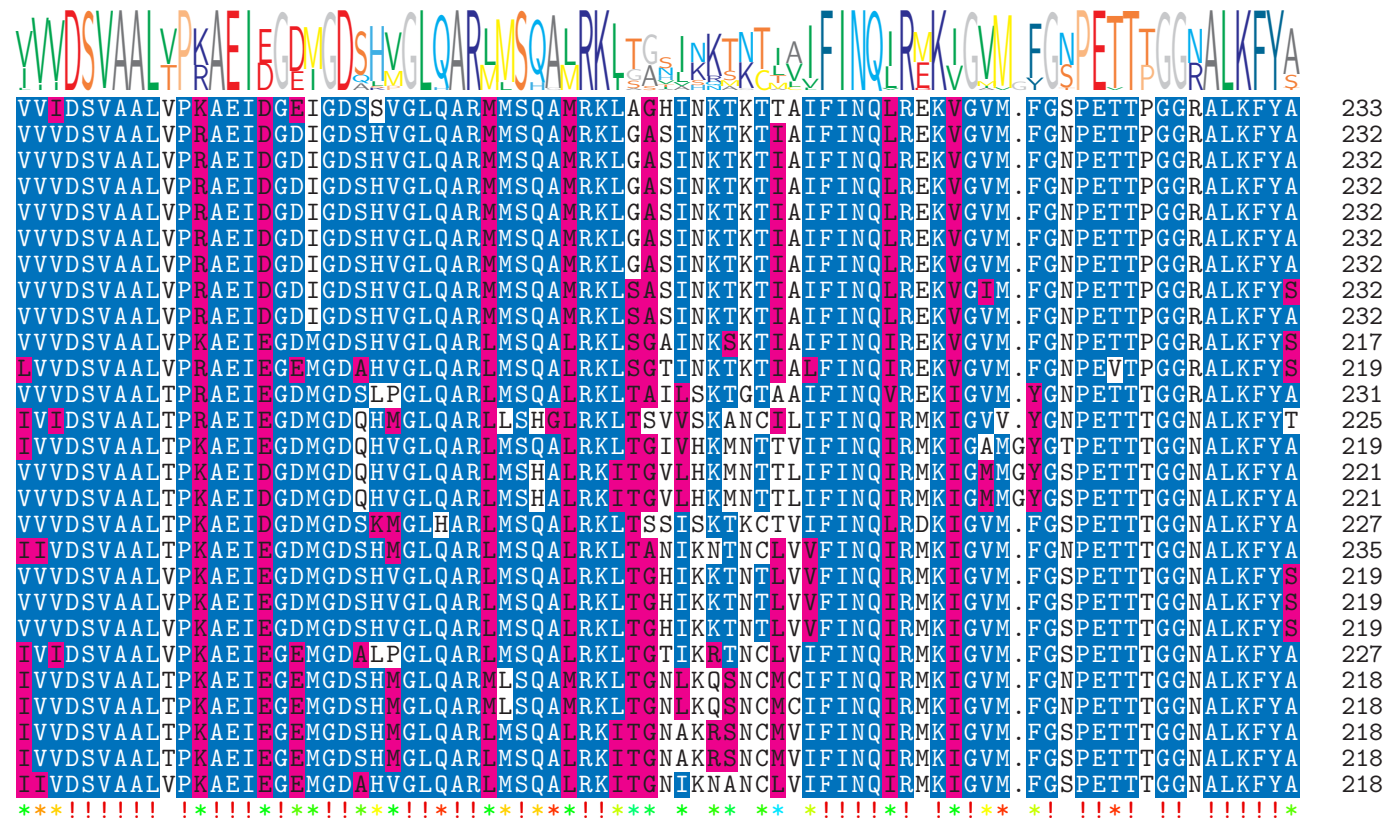
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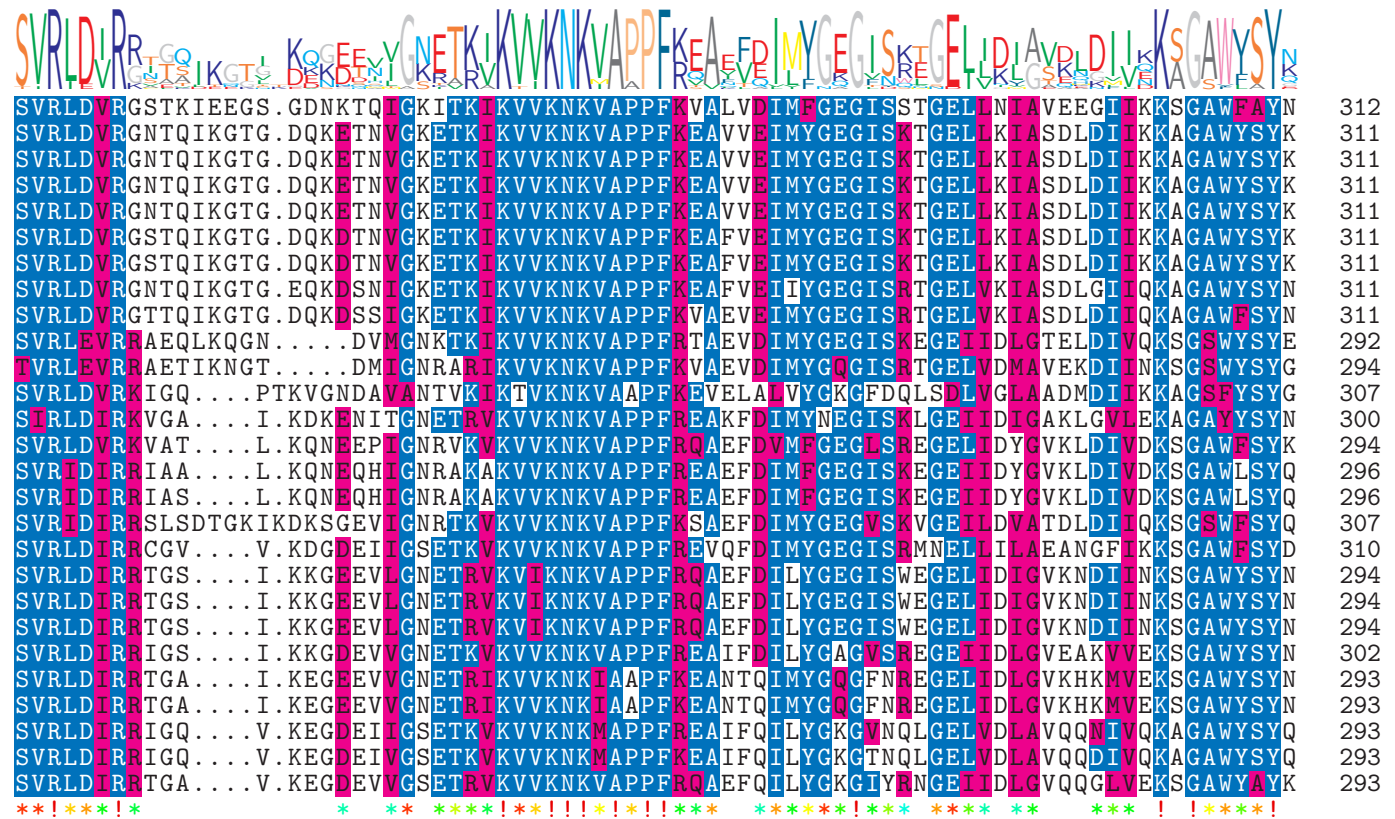
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consensus

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NTDB id 194 SPR RS08825 WP 001085462.1	DEKIGQGS	ENAKKYLA	EHPEIF	DEIDKQVRSKFGL	LIDGEEVSE		QDTENKK	DE....	PKKEEA..VNEEVPLDLGD	380
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consensus	***!!!	*****	**	*****	*****	**	**	*	*	

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consensus			

- X non conserved
- X similar
- X ≥ 50% conserved