

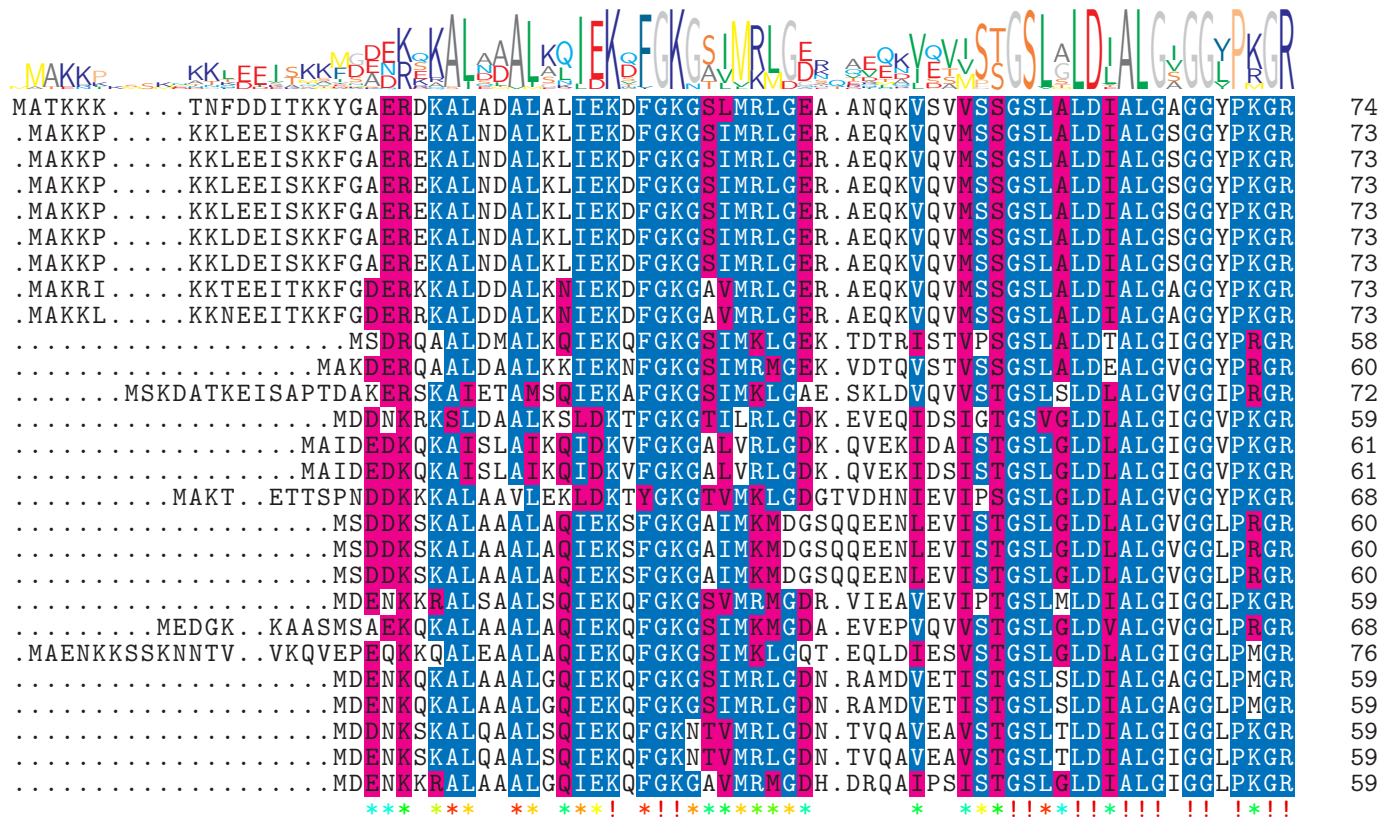
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NTDB id 1165 A1552VC RS01565 WP 000344154.1
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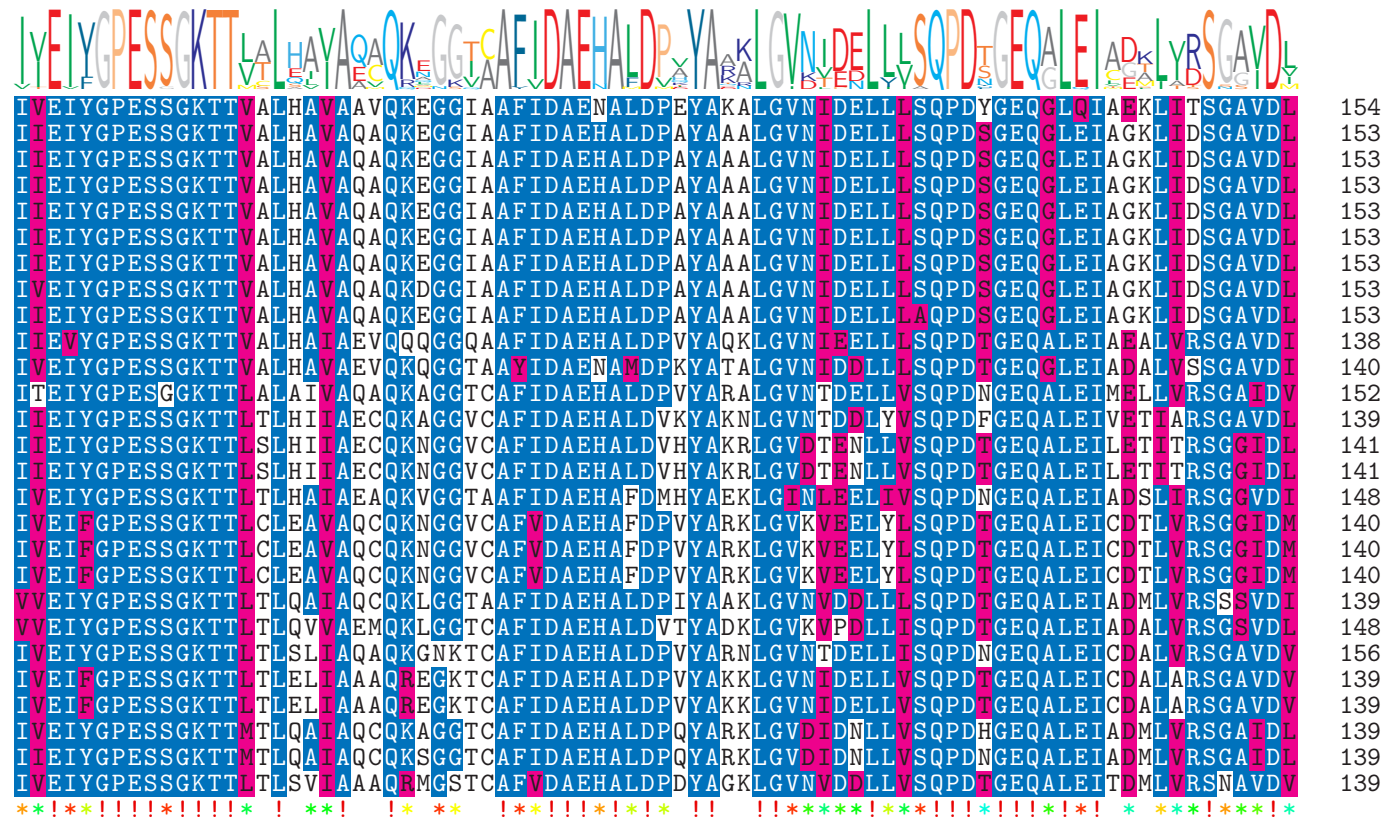
consensus



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consensus

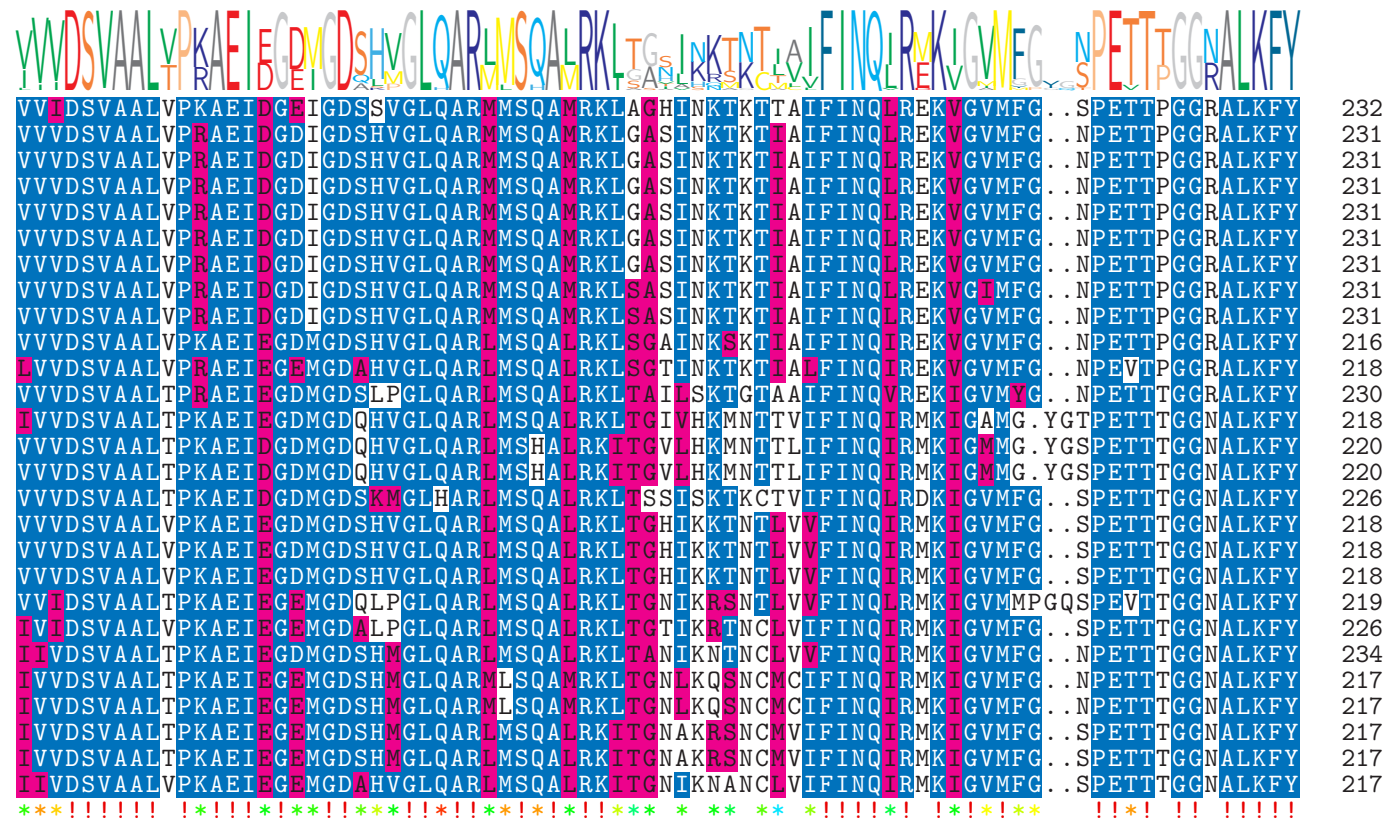
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consensus

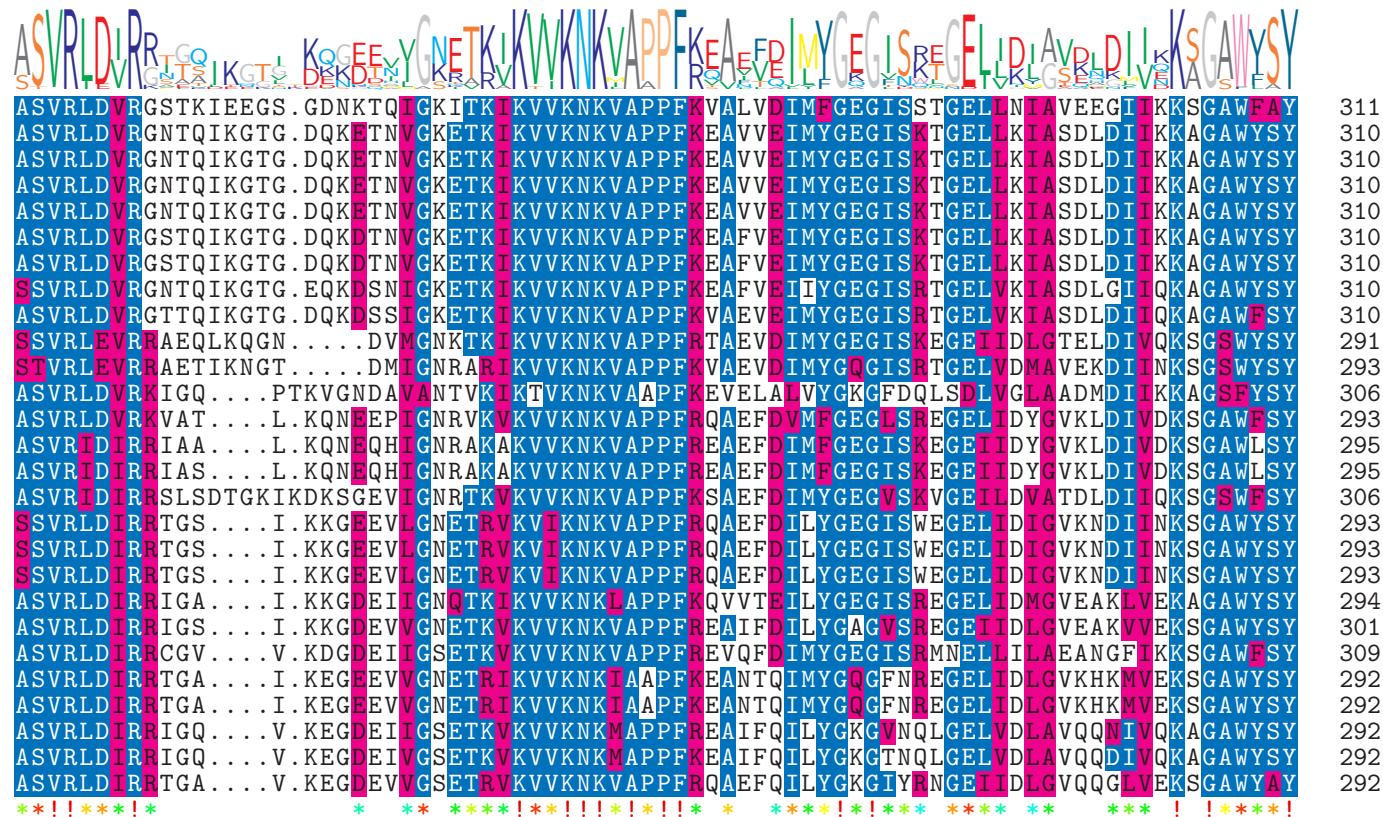
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consensus

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logo

NTDB id 600 KW2 RS01765 WP 011675413.1	NDEKIGQGA	EKAKNYLKEHQDV	FD	DEIDHKVRAAHG	LDDSEVAETE	EEETTAS	TKAKAKKEEKAVETEE	IELELED...	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	KDEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKKDE...	PKKEEA..VNEEVPLDLGDELEI				384
NTDB id 228 SPD RS09265 WP 001085462.1	KDEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKKDE...	PKKEEA..VNEEVPLDLGDELEI				384
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NTDB id 159 SP RS09750 WP 001085462.1	KDEKIGQGS	ENAKKYLAEHPEIF	DEIDKQVRSKFGL	IDGEEVSEQDTENKKDE...	PKKEEA..VNEEVPLDLGDELEI				384
NTDB id 527 SMSK321 RS11065 WP 001085454.1	KDEKIGQGS	ENAKKYLA	DNPEVFAEIDHQVRVQFGL	IDGEEAAE...	SKKDE...VAQVDS..VNEEVTLDLGDELEI				380
NTDB id 499 SM12261 RS08315 WP 078228268.1	KDEKIGQGS	ENAKKYLA	DNPEIFDEIDHQVRVQFGL	IDGEEIAE...	SKKDE...AAQVDS..VNEEVTLDLGDELEI				380
NTDB id 408 SMU RS09530 WP 002262392.1	NGEKIGQGS	ENAKKFLA	DNPEIFDDIDHKVVRVQYGL	IEEDNTEEEKQSSKEK.....	ETDEK..ADKNLVLELDDTIEL				381
NTDB id 453 Spy49 1753c ACI62003.1	NGEKIGQGS	ENAKRYLA	DHPQLFDEIDRKVVRVKFGL	LEESEESAMAVASE.....	E.....TDDLALDL	DNNGIEI			376
NTDB id 115 BSU 16940 NP 389576.2	EEERLGGG	REN	AKQFLKENKDIMLM	IQEQIREHYGL	DNNGVVQQAEETQEEL.EFE.E.....				348
NTDB id 625 LCA RS02525 WP 011374196.1	GSEIRIGGG	REN	AKNYLA	DHEDVEDEVRLKVRAAYG	ISDVPEEDLPTTEDE.QI.NILPDDST.....	EE.....			355
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NTDB id 1211 C694 RS00795 WP 000963128.1	QDKKLGGG	REN	AKALLKEDKALA	DEITLKI	KESIGSNEEI.....MPLPDEPLEEME.....				347
NTDB id 1334 RAOC RS04870 WP 004918236.1	QDTKLGGGR	DTVKELIK	DNPELAEELEAKIREKLQAK.....						343
NTDB id 1130 NGFG RS03960 WP 003688695.1	NGAKIGQGGK	DNVRVWLKENPEIS	DEIDAKIRALNGVEMHI.....	TEGTQDETGERP.....EE.....					348
NTDB id 1124 NGFG RS03960 WP 003688695.1	NGAKIGQGGK	DNVRVWLKENPEIS	DEIDAKIRALNGVEMHI.....	TEGTQDETGERP.....EE.....					348
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	QGNKIGQGGK	NNVIRY	LEENPQISTEIEAVIREQLLT	TKASD.....QTAA.HDETEEEP	DLL...ES.....				349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	QGNKIGQGGK	NNVIRY	FEENNQIAEIEIERNIREQLLT	TTGTN.....GAVQ.I.EDEE	EPDLL...LES.....				349
NTDB id 1205 PSJM300 05545 AFN77183.1	KGSKIGQGGK	ANA	KYLEENPEIGQIEIEQQIRAKLLV	TSAN.....TKATPVAEDLADADI.....					347
consensus	**	!!!	*****	**	*****	*****	**	*	

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consensus

 non conserved
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 ≥ 50% conserved