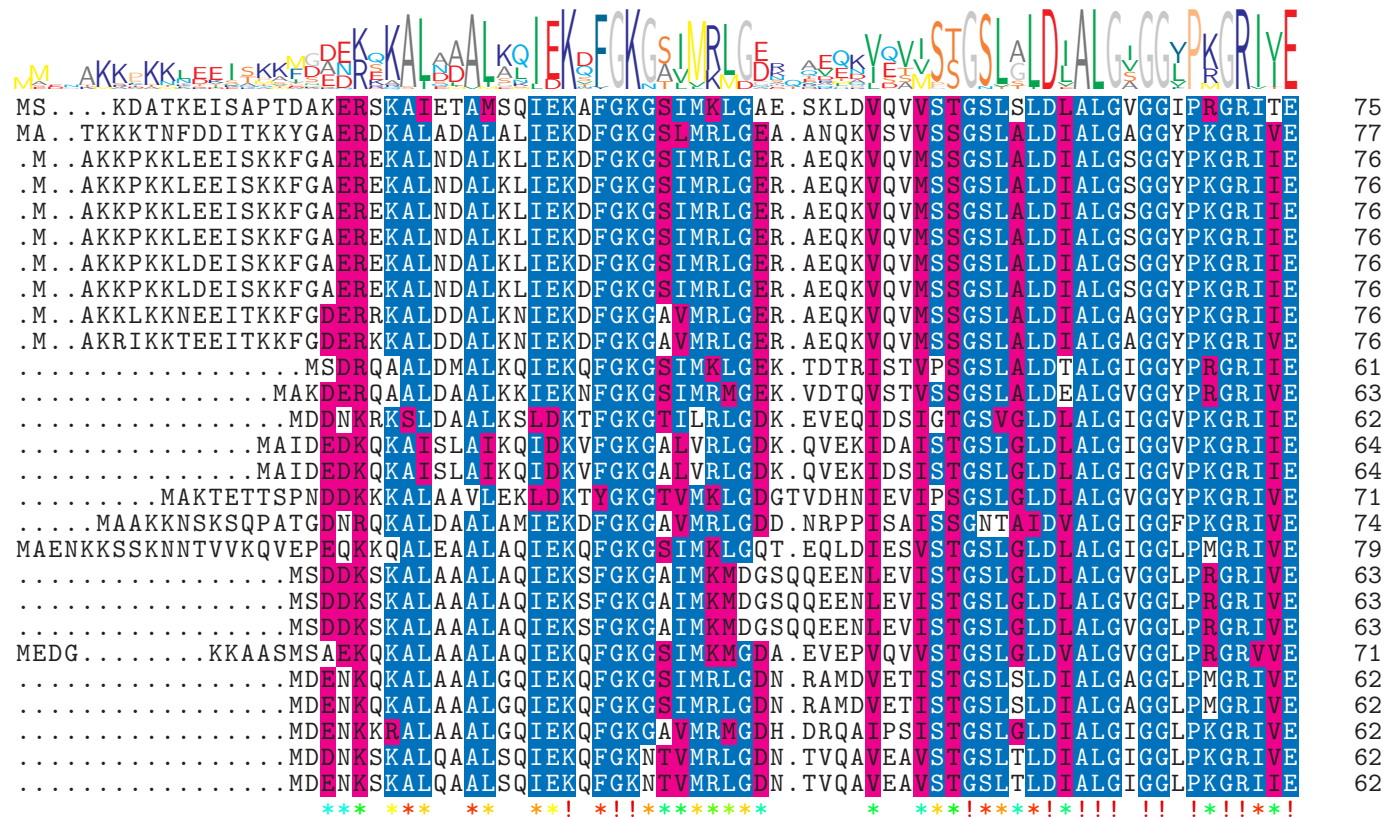


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consensus

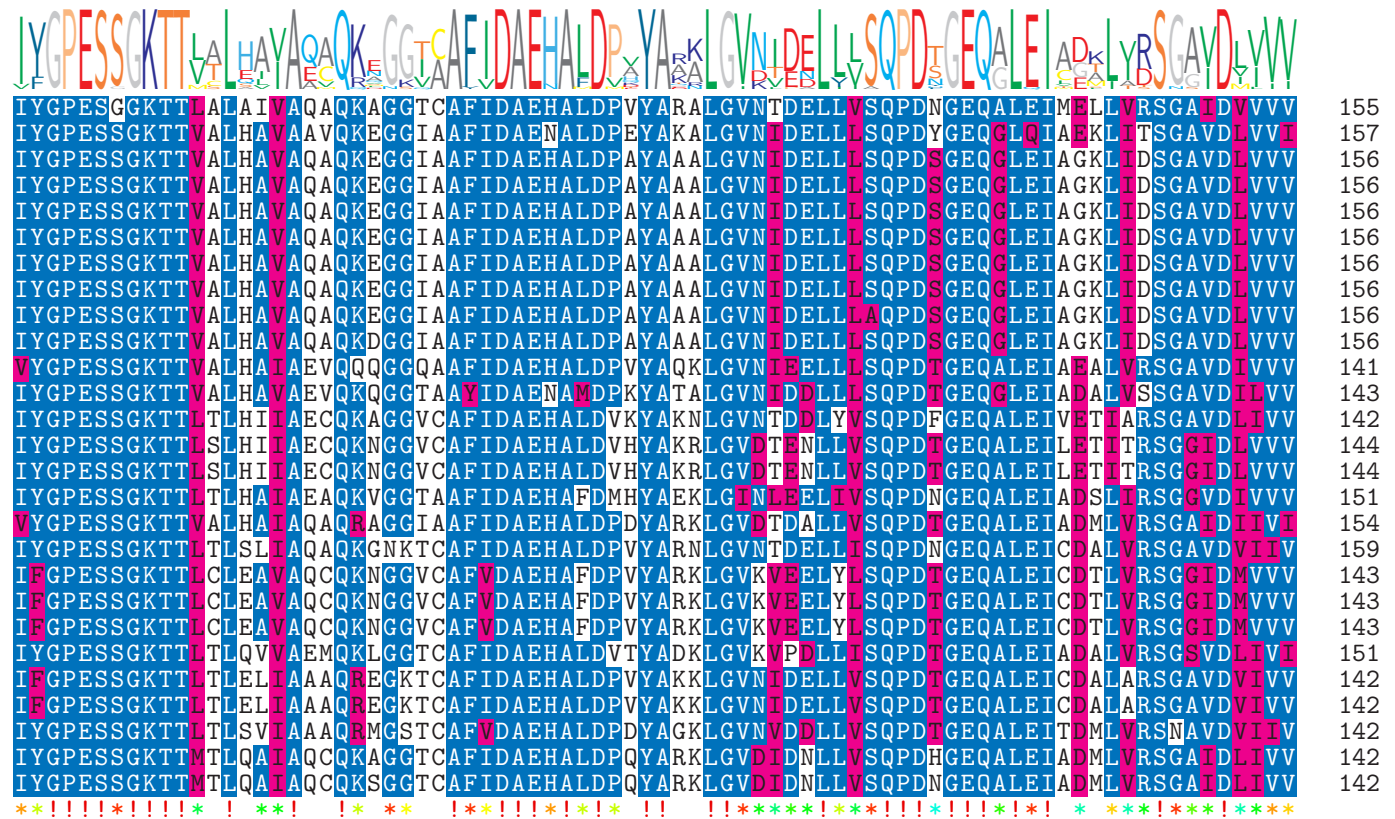
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consensus

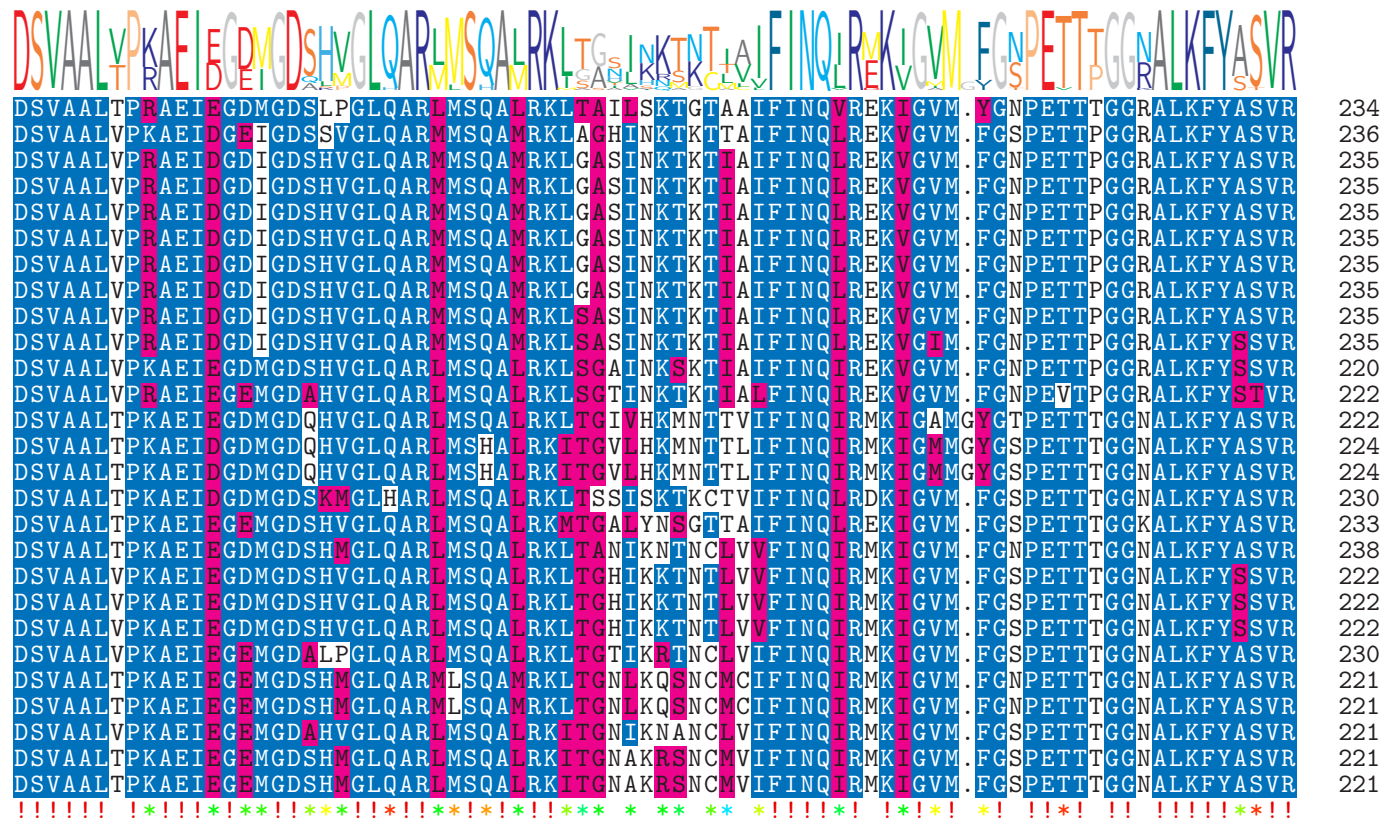
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consensus

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NTDB id 269 KZH43 RS08665 WP 001085462.1	LDVRGNTQIKGTG.DQKETN	VGKETKIKVVKNKVAPPFKEAVVE	IMY	GEGISKT	GELLK	IASDL	DIKK	AGAWYSYKDEK	314
NTDB id 228 SPD RS09265 WP 001085462.1	LDVRGNTQIKGTG.DQKETN	VGKETKIKVVKNKVAPPFKEAVVE	IMY	GEGISKT	GELLK	IASDL	DIKK	AGAWYSYKDEK	314
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NTDB id 159 SP RS09750 WP 001085462.1	LDVRGNTQIKGTG.DQKETN	VGKETKIKVVKNKVAPPFKEAVVE	IMY	GEGISKT	GELLK	IASDL	DIKK	AGAWYSYKDEK	314
NTDB id 499 SM12261 RS08315 WP 078228268.1	LDVRGSTQIKGTG.DQKDTN	VGKETKIKVVKNKVAPPFKEAFVE	IMY	GEGISKT	GELLK	IASDL	DIKK	AGAWYSYKDEK	314
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NTDB id 453 Spy49 1753c ACI62003.1	LDVRGTTQIKGTG.DQKDSS	IGKETKIKVVKNKVAPPFKVAEVE	IMY	GEGISRT	GELVK	IASDL	DIQK	AGAWFSYNGEK	314
NTDB id 408 SMU RS09530 WP 002262392.1	LDVRGNTQIKGTG.EQKDSN	IGKETKIKVVKNKVAPPFKEAFVE	IIY	GEGISRT	GELVK	IASDL	GIIQK	AGAWYSYNGEK	314
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	IDIRRIAA....L.KQNEQH	IGNRAKAKVVKNKVAPPFREA	AEFD	IMF	GEGISKEGE	IIDYGVKL	DIVDK	SGAWLSYQDKK	299
NTDB id 1211 C694 RS00795 WP 000963128.1	IDIRRIAS....L.KQNEQH	IGNRAKAKVVKNKVAPPFREA	AEFD	IMF	GEGISKEGE	IIDYGVKL	DIVDK	SGAWLSYQDKK	299
NTDB id 1334 RAOC RS04870 WP 004918236.1	IDIRRLSDTGKIKDKSGEV	IGNRTKVVKNKVAPPFKSA	AEFD	IMY	GEGVSKVGE	ILDVATDL	DIIQK	SGSWFSYQDTK	310
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NTDB id 1427 RS RS02760 WP 011000510.1	LDIRRIGS....I.KKGDEV	VGNETKVVKNKVAPPFREA	IFD	DILY	GAGVSREGE	IIDLGV	EAKVVEK	SGAWYSYNGER	305
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NTDB id 1313 DR RS12030 WP 010888966.1	IGQGKEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPA.....APEAAEA.....	363
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NTDB id 625 LCA RS02525 WP 011374196.1	IGQGRENAKKNYLADHEDVEDEVRLKVRAAYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	LGQGRENAKALLKEDKALADEITLKIKEISIGSNEEIMPLP.....DEPLEE.....ME.....	347
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NTDB id 1205 PSJM300 05545 AFN77183.1	IGQGKANAAKYLEENPEIGQEIEQQIRAKLLVTSANTKATPVAEDLADADI.....	347
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	IGQGKNNVIRYFEENNQIAEEIERNIREQLLTGTNGAVQ.I.EDEEPPDLL.....LES.....	349
consensus	*!!!!***** ** * *** ***** ** ** *	

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