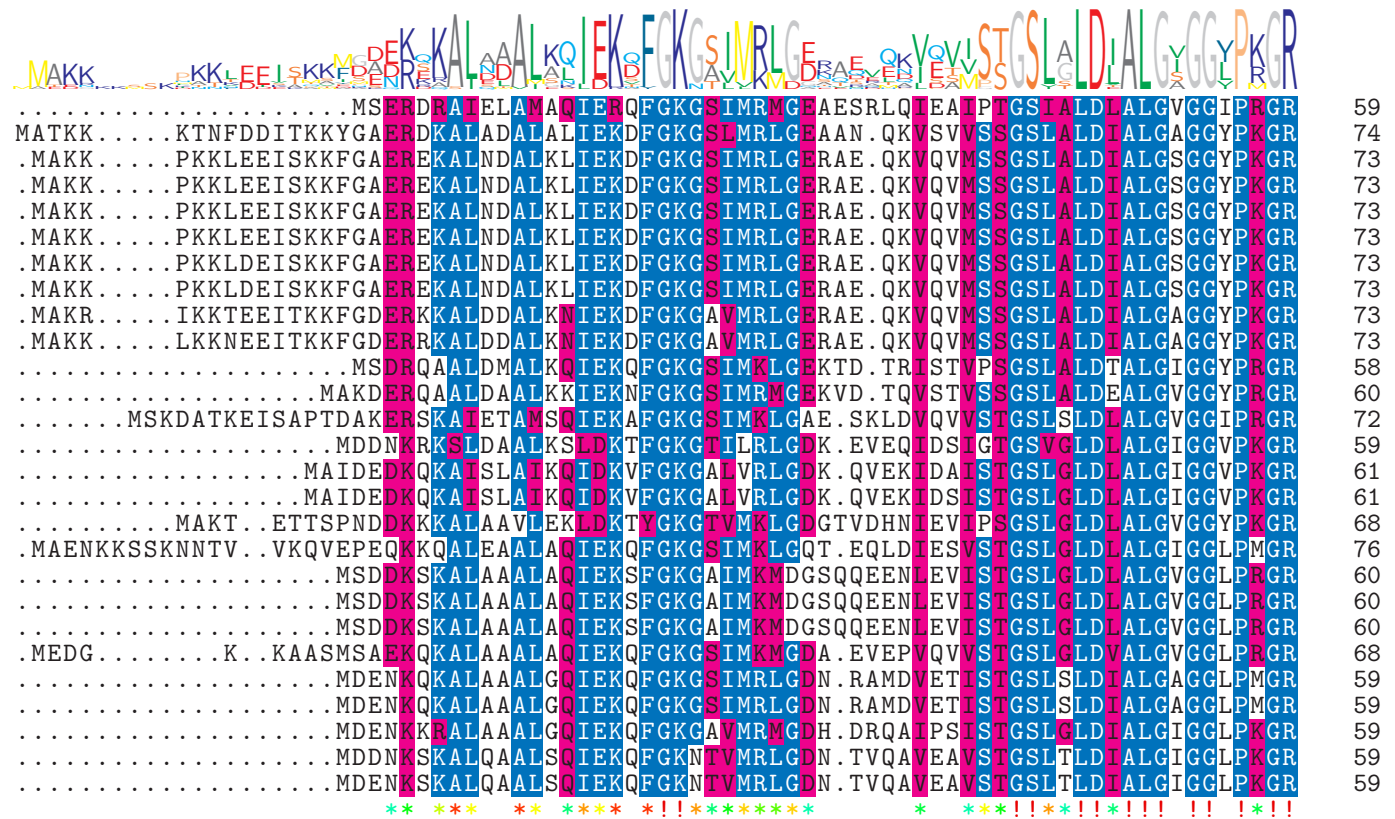


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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

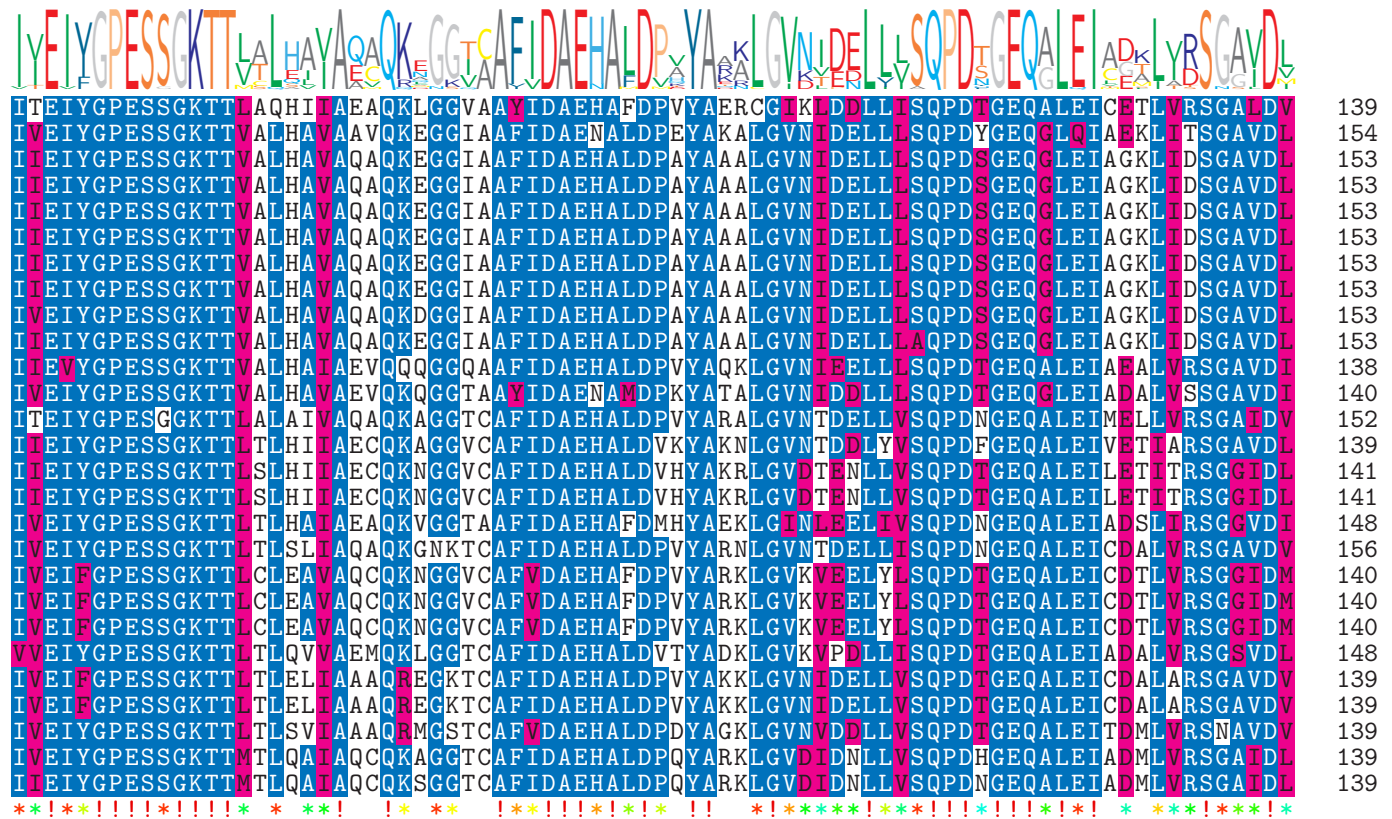
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consensus

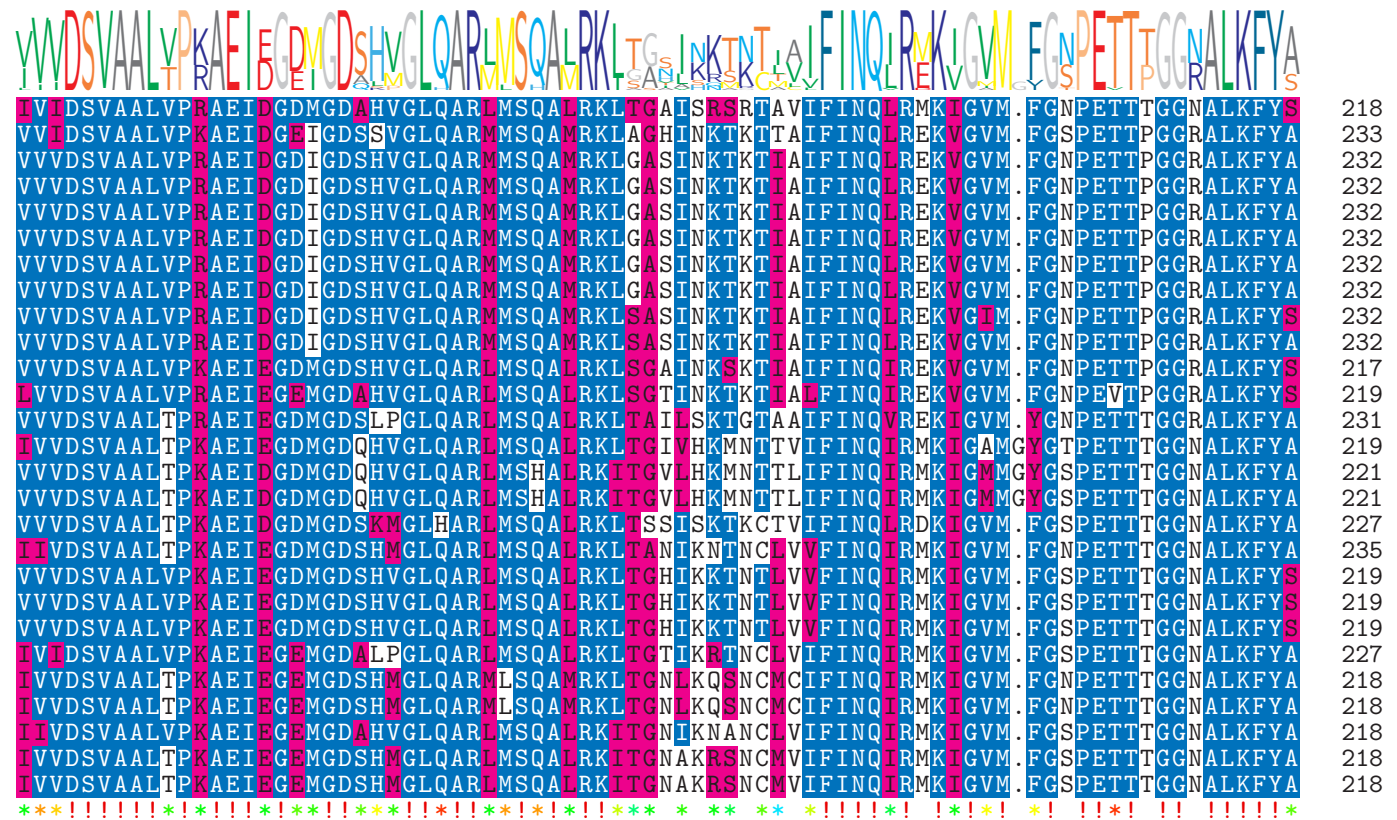
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

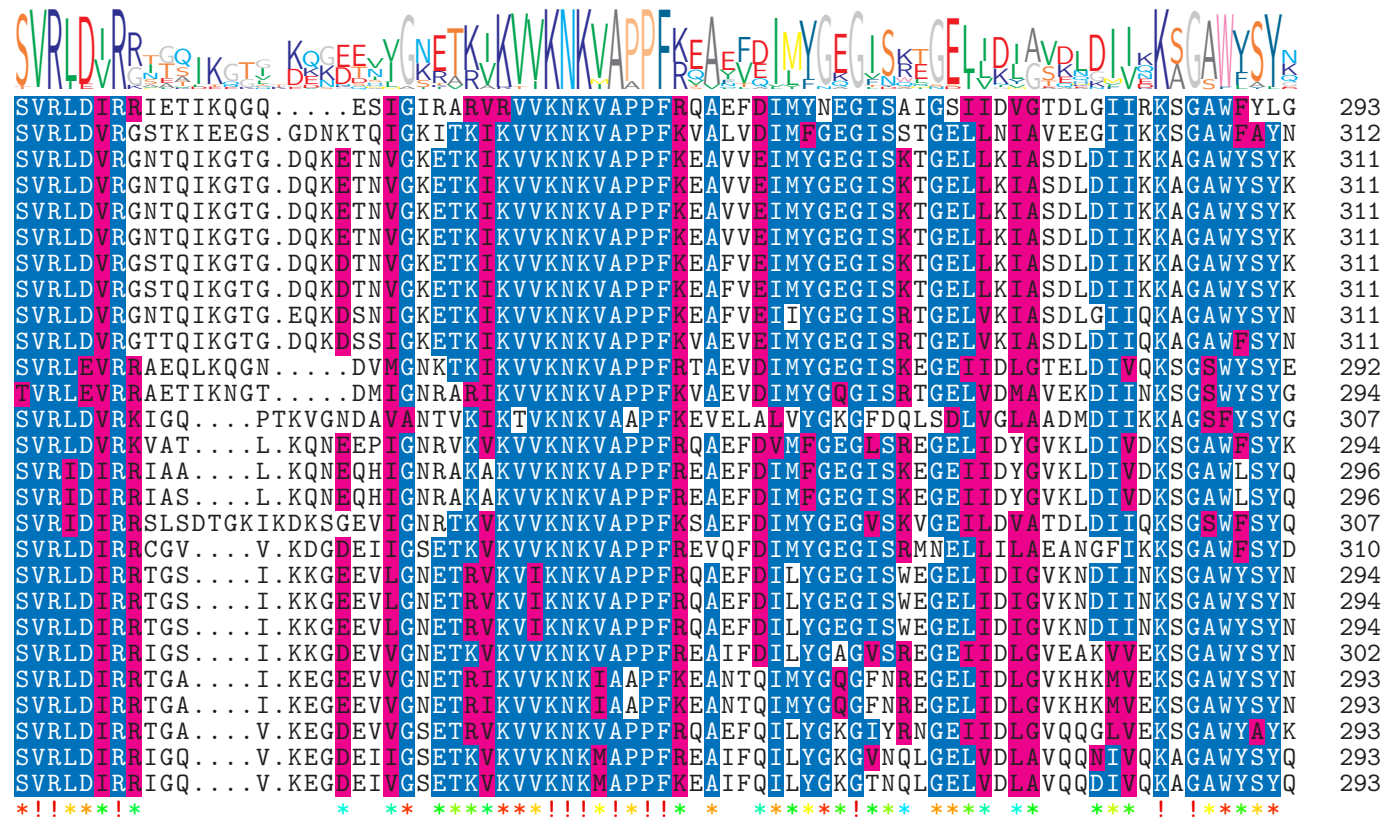
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consensus

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NTDB id 1205 PSJM300 05545 AFN77183.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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Sequence	Score
DEERLGGGRENKFLKANPDI LRDI EQQIRAA SPELRSRINYDGTSAEAEEL... A... DE.VLS.....	352
DEKIGQGAEKAKNYLKHQDVDFEIDHKVRAAHGLDDSEVAETEETASTKTAKAKKEEKAVETEEIELEED....	387
DEKIGGQSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE... PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGQSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE... PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGQSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE... PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGGQSENAKKYLAHNPEIFDEIDHQRVVRVQGLIDGEEIAES... KKDE...AAQVDS..VNEEVTLDLGDLEIE	381
DEKIGGQSENAKKYLAHNPEVFAEIDHQRVVRVQGLIDGEEAAES... KKDE...AAQVDS..VNEEVTLDLGDLEIE	381
GEKIGGQSENAKKFLADNPEIFDDIDHKVVRVQGLIEEDNTEEKQSSKEK... ETDEK..ADKNLVLELDDTIELE	382
GEKIGGQSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE... E...TDDLALDLNGIEIE	377
EERLGGGRENAKQFLKENKDIIMLMIQEQIREHYGLDNNGVVQQAAETQEEL.EFE.E...	348
SERIGGQRENAKNYLADHEDVEURLKVRAAYGISDVPEEDLPTTEDE.QI.NILPDST...EE...	355
DERIGGQKEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAAPEAA...EA...	363
DKKLGQGRENSKAFLENPEIAD EITKAIQNSMGI EGMI...SGSEDDE..G..EE...	343
DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP..L..EEME...	347
DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP..L..EEME...	347
DTKLGGGRDITVKELIKDNPELAEELEAKIREKLQAK.....	343
GKIGGQKNNAVKWLKENPEIADKLEQEI RNTLASNPNS...VLKATASDSN..DESDVEEFSEEEEL...	372
GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGT.....QDET DGERPEE...	348
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GERIGGGRDNCREFLRENAELAR E IENKVREHLGVTPMG...AV.TLAEVE..ED...	352
GDKIGGQKANACKYLKENPEIAKTLDKKLREMLLNPNEM...QLIAETSSAA..DDVEFGAVPEEF...	354
GDKIGGQKANACKYLKENPEIAKTLDKKLREMLLNPNEM...QLIAETSSAA..DDVEFGAVPEEF...	354
GSKIGGQKANAAKYLEENPEIGQIEEQIRAKLLVTSAN...TKATPVAEDL..ADADI...	347
GNKIGGQKNNVIRYLEENPQISTEIEAVIREQLLTKASD...QTAA.HDETE..EEPDLL..ES...	349
GNKIGGQKNNVIRIFYENNQIAEIERNIREQLLTGTN...GAVQ.I.EDE..EEPDLL..LES...	349

logo

		IEE	
NTDB id 35775	STHE RS04640 WP 012871408.1	...	352
NTDB id 600	KW2 RS01765 WP 011675413.1	...	387
NTDB id 269	KZH43 RS08665 WP 001085462.1	IEE	388
NTDB id 228	SPD RS09265 WP 001085462.1	IEE	388
NTDB id 194	SPR RS08825 WP 001085462.1	IEE	388
NTDB id 159	SP RS09750 WP 001085462.1	IEE	388
NTDB id 499	SM12261 RS08315 WP 078228268.1	IEE	384
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NTDB id 1205	PSJM300 05545 AFN77183.1	...	347
NTDB id 1022	ACIAD RS06390 WP 004925639.1	...	349
NTDB id 1074	ABD1 RS10195 WP 000344167.1	...	349
consensus			

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