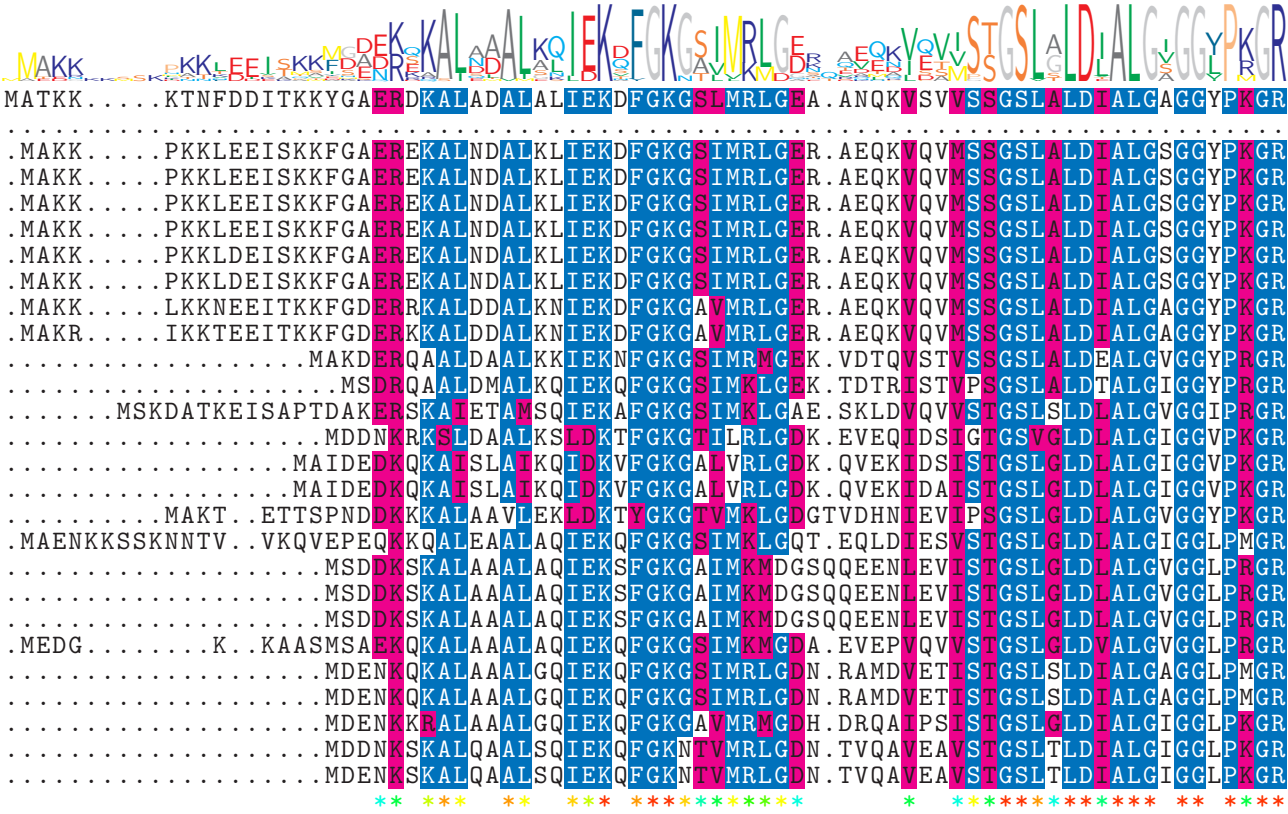


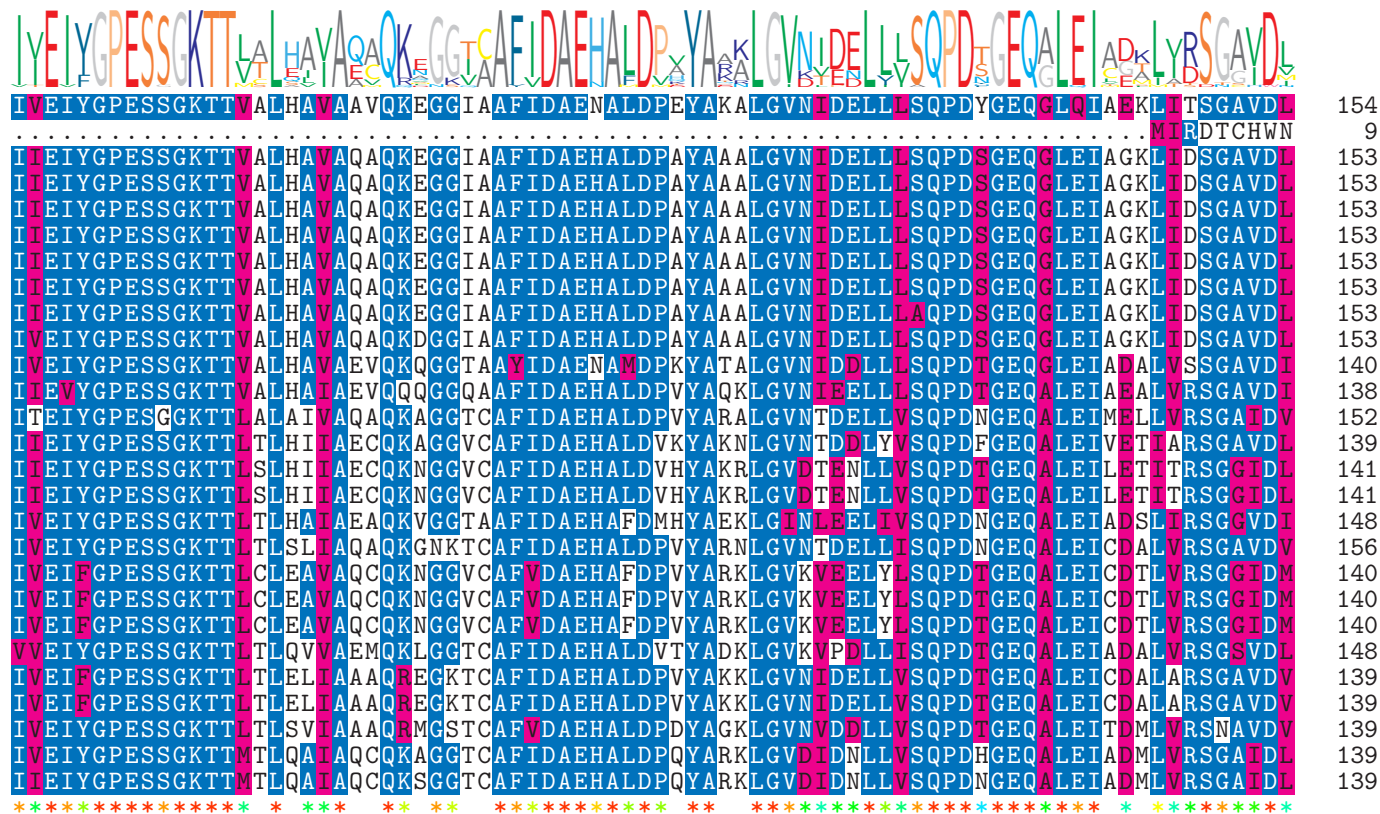
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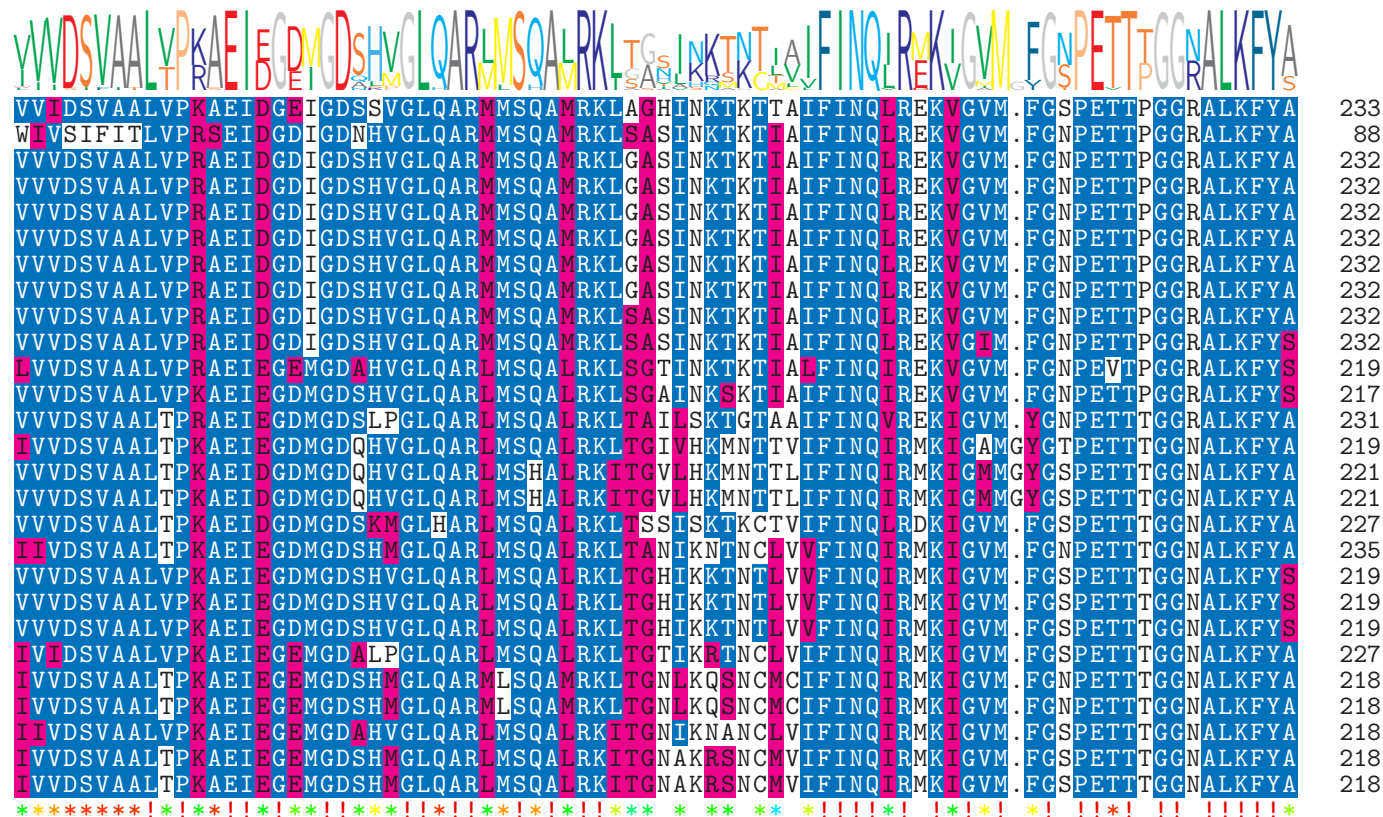


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consensus



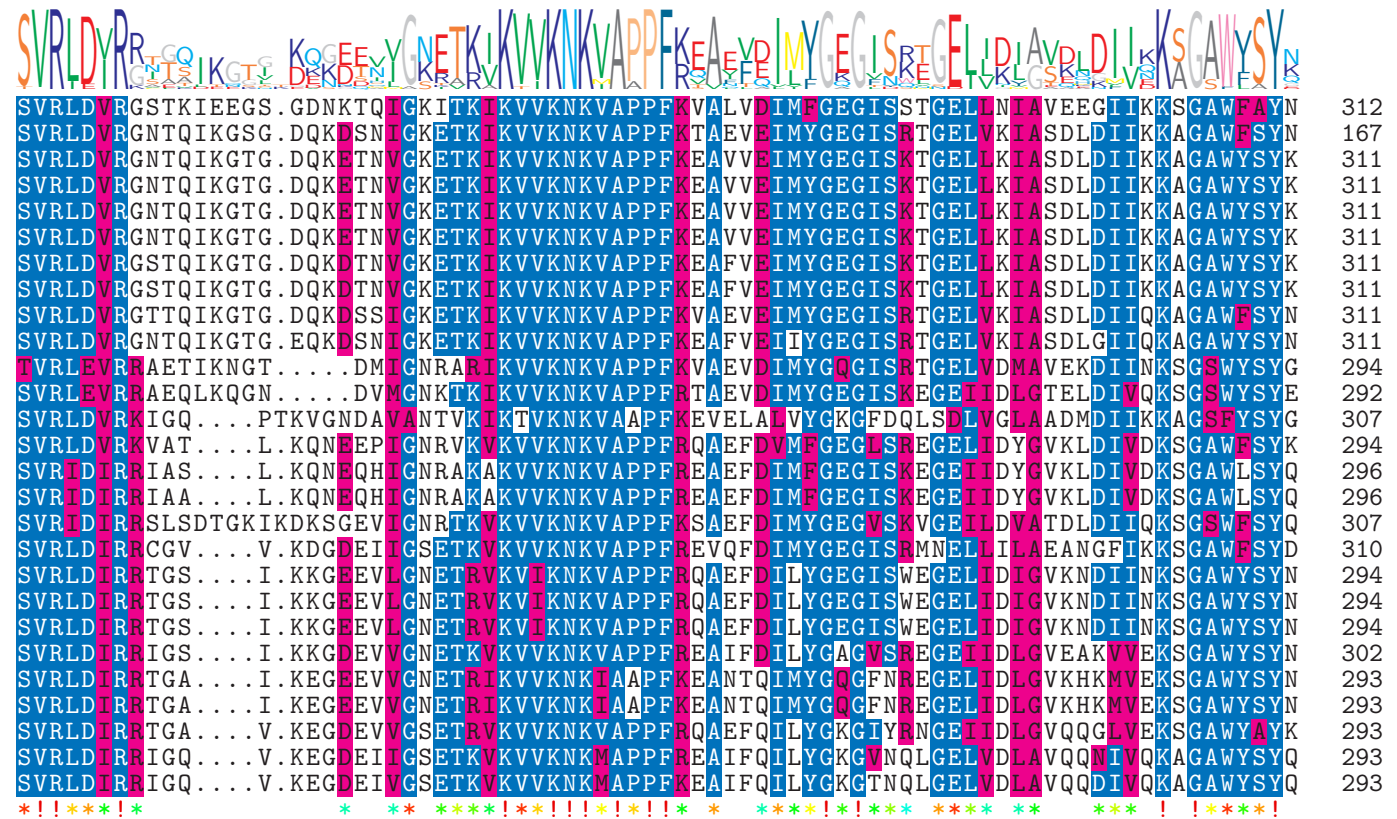
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consensus

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DEKIGQGSSENAKKYLAADNPEVFAEIDHQRVVFGLIDGEEAAES...KKDEV...AQVDS...VNEEVTLDLGDLEIE	381
GEKIGQGSSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSSENAKKFLADNPEIFDDIDHKVVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
SERIGQGRENAKNYLADHEDVEDEVRLKVRRAAYGISDVPPEEDLPTTEDE.QIN.....ILPDDSTEE.	355
EERLGGQRENAKQFLKENKDMLMTEQEIREHYGLDNNGVVQQQAETQEELE.....FE.E.....	348
DERIGQGEKKTIAIYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEA..AEA.....	363
DKKLGGQRENSKAFLENPEIADAITKAIQNSMGTIEGMI.....SGSEDDE...GEE.....	343
DKKLGGQRENAKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP...LEEME.....	347
DKKLGGQRENAKALLKEDKALADEITLKIKESIGSNEEI...MPLPDEP...LEEME.....	347
DTKLGGQGRDTPVKELIKDNPELAELEAKIREKLQAK.....	343
GTKIGQGGKNNAVKWLKENPEIADKLEQEIRNTLASNPNS...VLKATASDS..NDES DVEEF...SEEEL...	372
GAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGT.....QDET DGER...PEE.....	348
GAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGT.....QDET DGER...PEE.....	348
GAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGT.....QDET DGER...PEE.....	348
GERIGQGRDNCREFLENARLEIENKVRHGLVTPMG...AV.TLAEV..EED.....	352
GDKIGQGGKANACKYLKENPEIAKTLDDKLREMLLNPNEM...QLIAETSSA..ADDVEFGAV...PEEF....	354
GDKIGQGGKANACKYLKENPEIAKTLDDKLREMLLNPNEM...QLIAETSSA..ADDVEFGAV...PEEF....	354
GSKIGQGGKANAAKYLEENPEIGQEIEQQIRAKLLVTSAN...TKATPVAED..LADADI.....	347
GNKIGQGGKNNVIRYLEENPEIISTEIEAVIREQLLTAKSD...QTAA.HDET..EEEPDLL...ES.....	349
GNKIGQGGKNNVIRYLEENPEIISTEIEAVIREQLLTGTN...GAVQ.I.ED..EEEPDLL...LES.....	349

logo

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NTDB id 1074 ABD1 RS10195 WP 000344167.1	...		349
consensus			

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