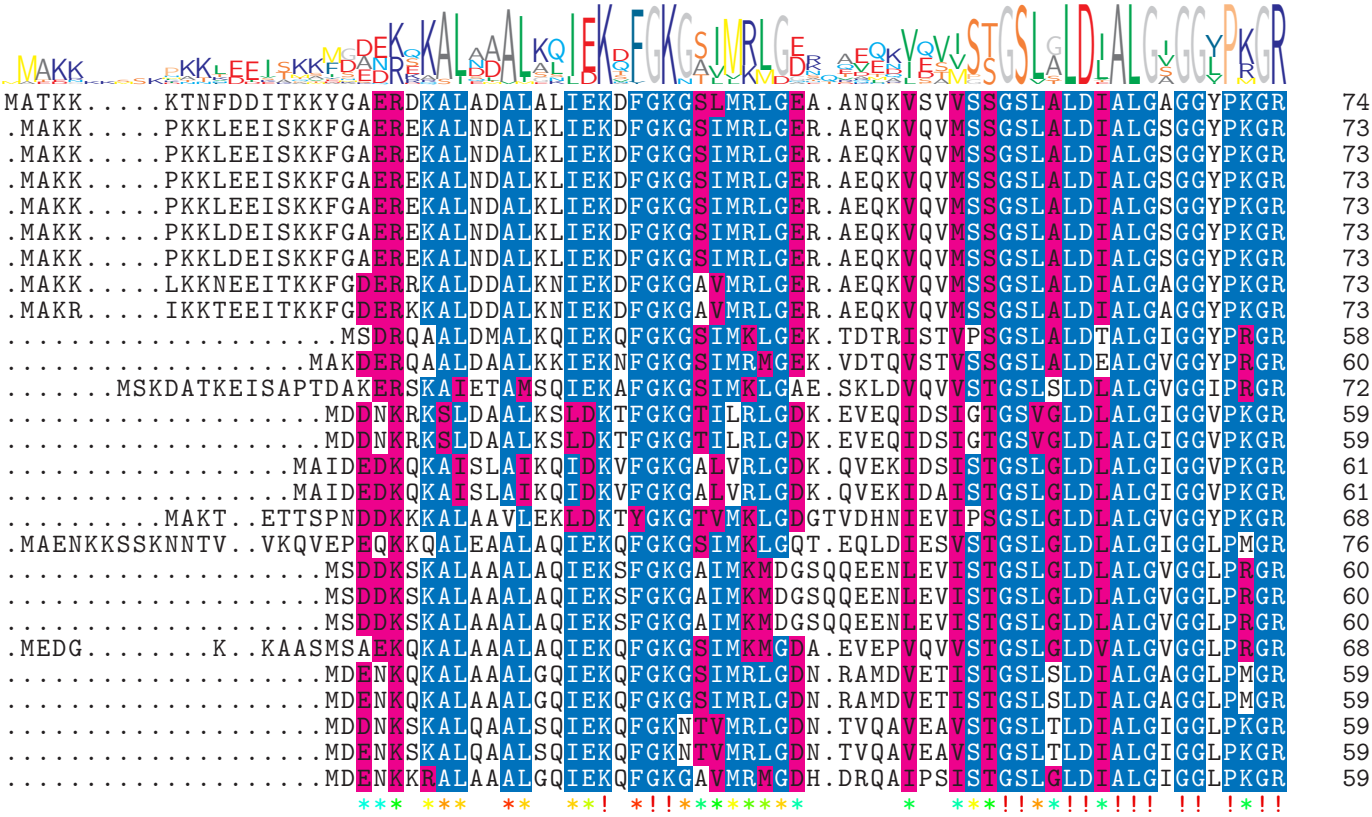


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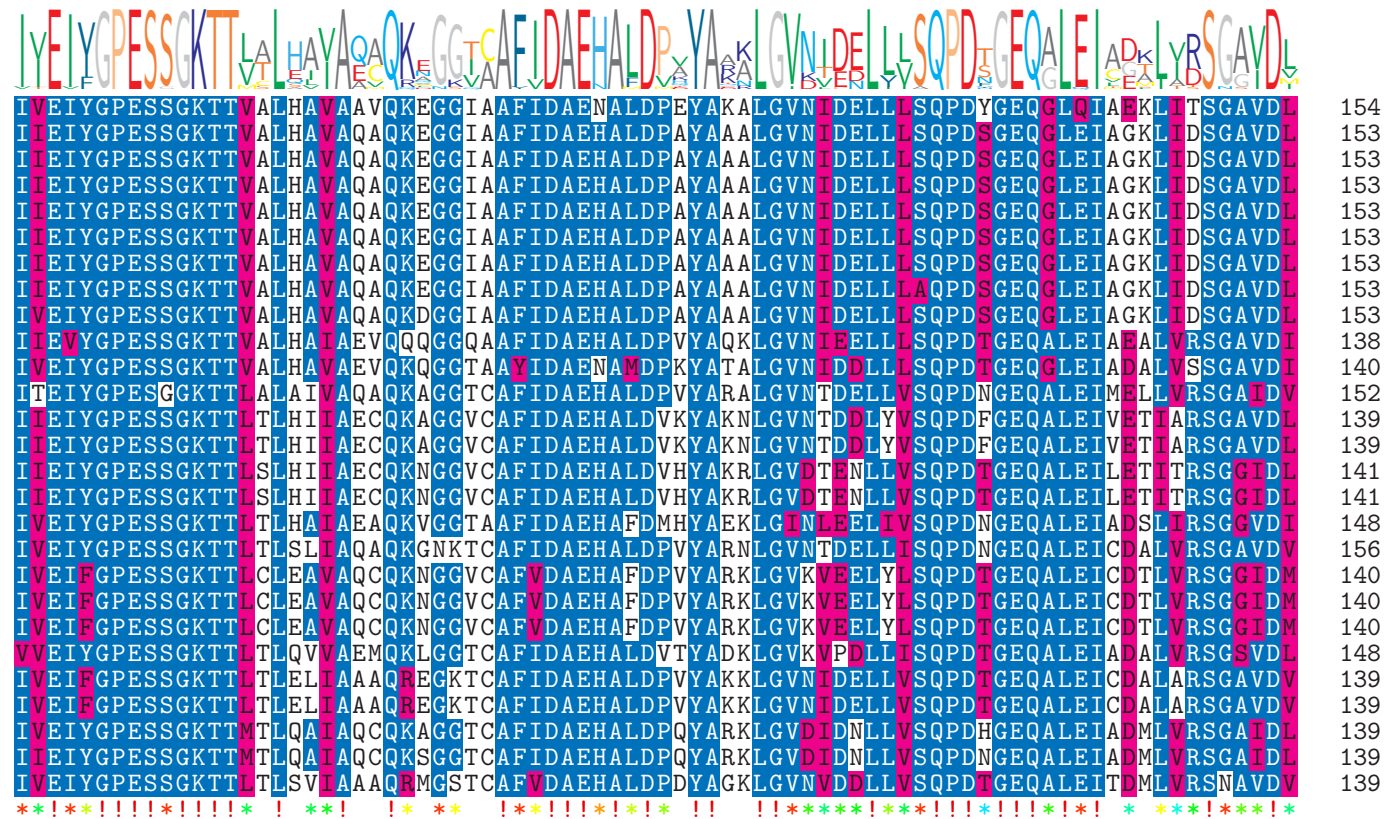
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consensus



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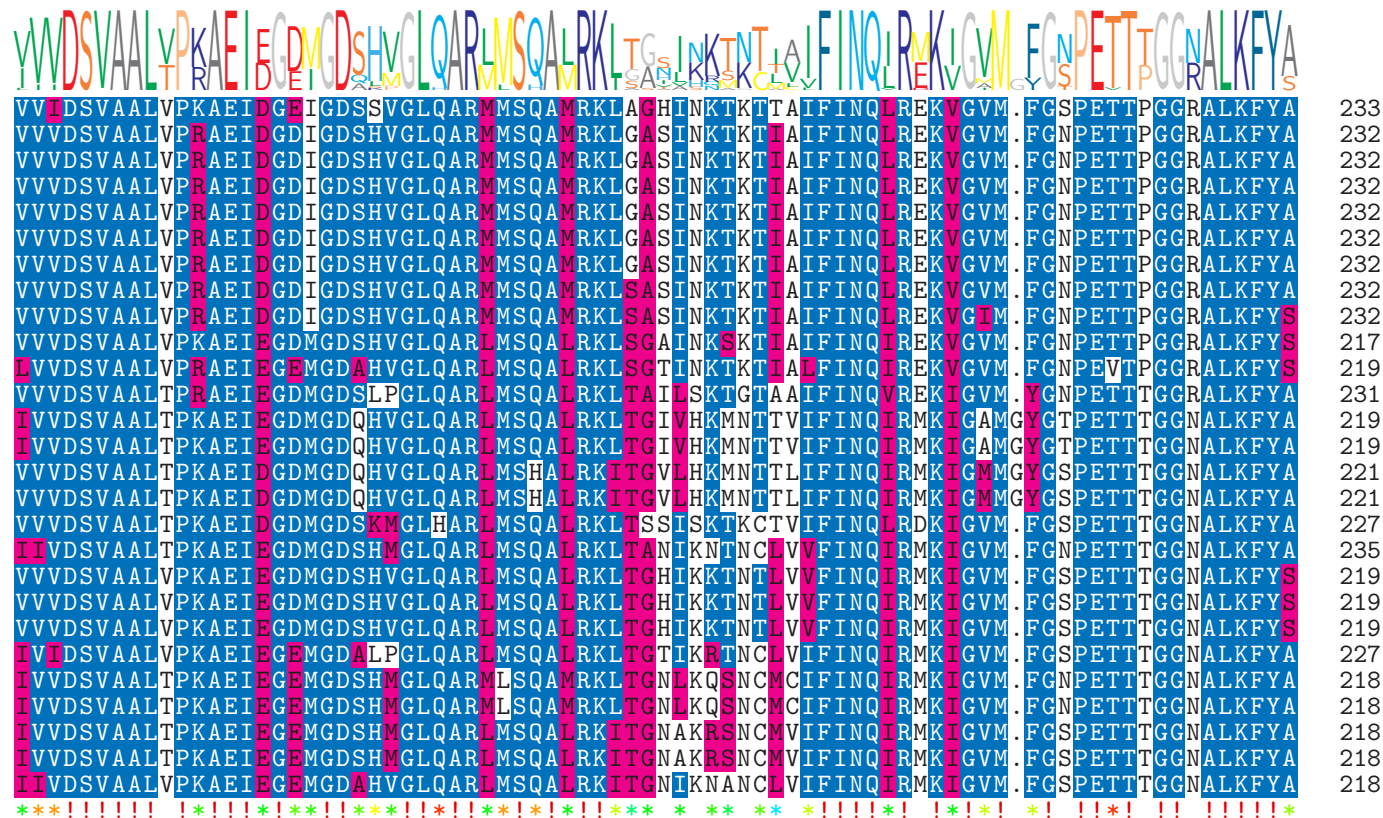
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consensus

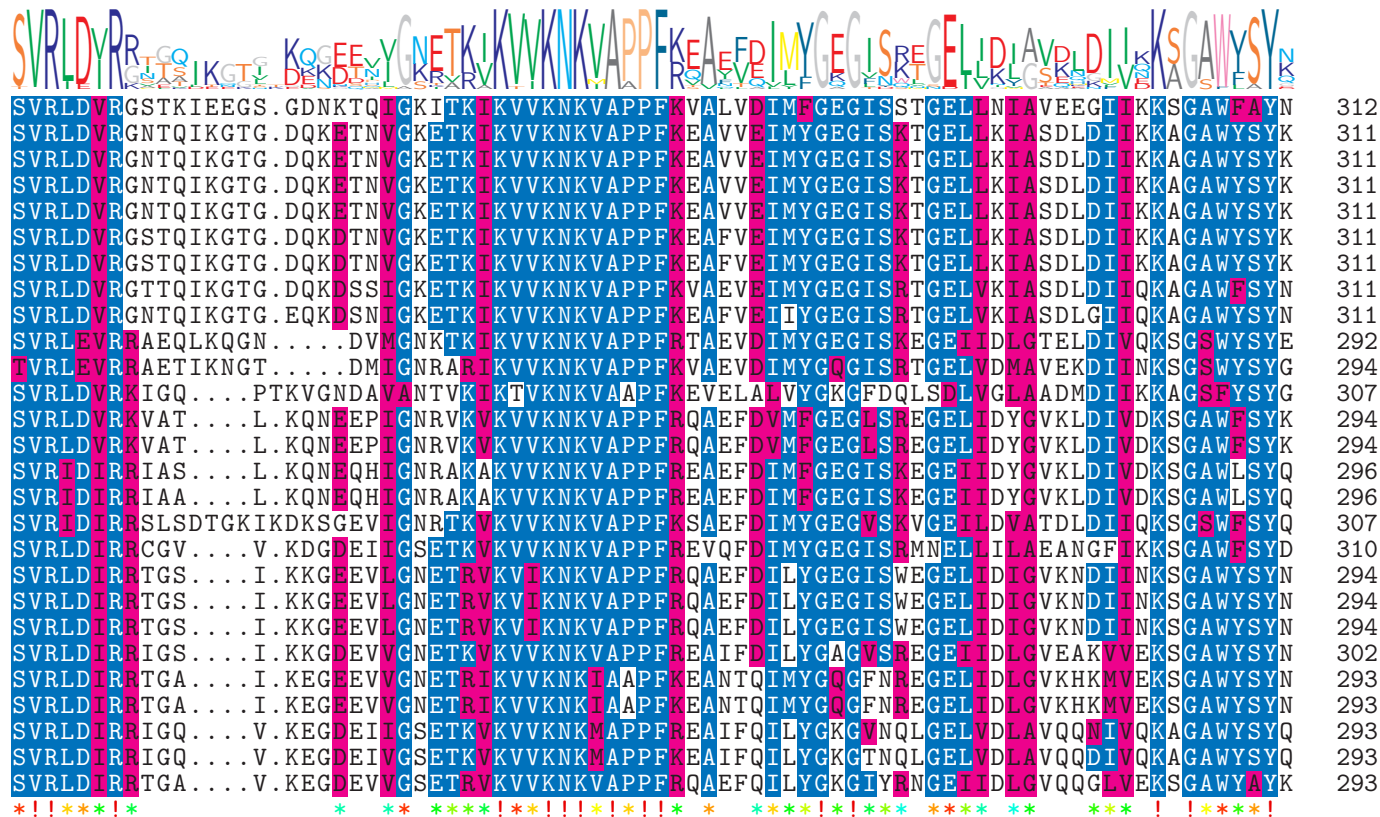
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consensus

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consensus

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DEKIGQGSENAKKYLAADNPFIIDHQRVVFGLIDGEEIAES...KKDE...AAQVDS..VNEEVTLDLGDLEIE	381
GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGSENAKKFLADNPFIIDHQRVVFGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
EERLGQGRENAKQFLKENKDIMLMIQEQUIREHYGLDNGGVVQQAEETQEEL.EFE.E.....	348
SERIGQGRENAKNYLADHEDVEDEVRLKVRAAAYGISDVPPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DERIGGQKEKTIATYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAE..A.....	363
DKKLGQGRENSKAFLKENPEIADITKAIQNSMGIEGMI....SGSEDD..EGE..E.....	343
DKKLGQGRENSKAFLKENPEIADITKAIQNSMGIEGMI....SGSEDD..EGE..E.....	343
DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDE..PLE..EME.....	347
DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDE..PLE..EME.....	347
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GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVEMHI....TEGTQDETGE..R.....PEE.....	348
GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNGVEMHI....TEGTQDETGE..R.....PEE.....	348
GERIGGQGRDNCREFLRNARLARIENKVRREHLGVTPMG....AVTLA.EEEVE..D.....	352
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GDKIGGQKANACKYLKENPEIAKTLDDKLREMLLNPNEM....QLIAETSSAAD..DVEFGAVPEEF..	354
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GNKIGGQKNNVIRYFEENNQIAEIEIERNIREQLLTTGTN....GAVQ..I..EDEE..EPDLL..LES.....	349
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logo

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NTDB id 1205 PSJM300 05545 AFN77183.1	...		347
consensus			

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