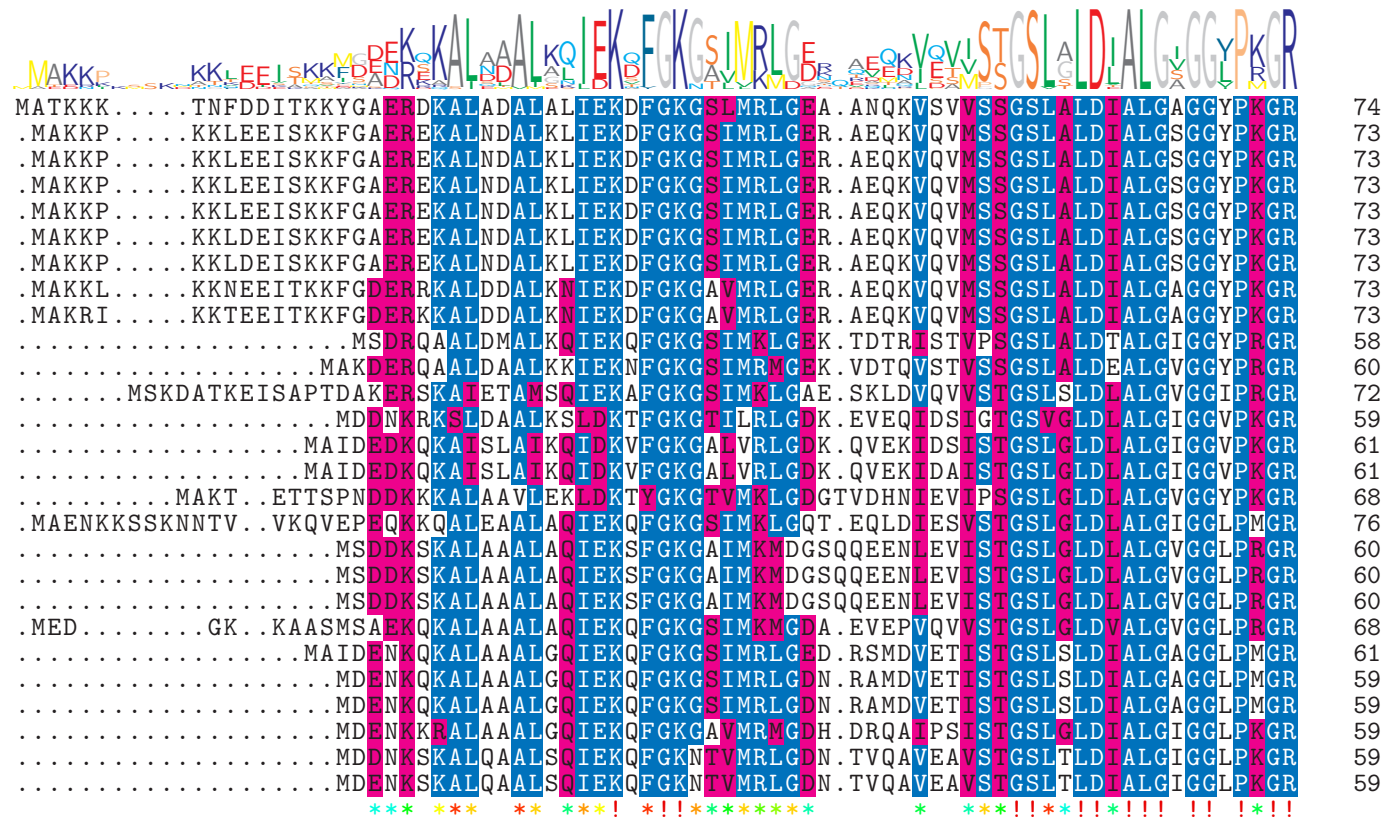


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consensus

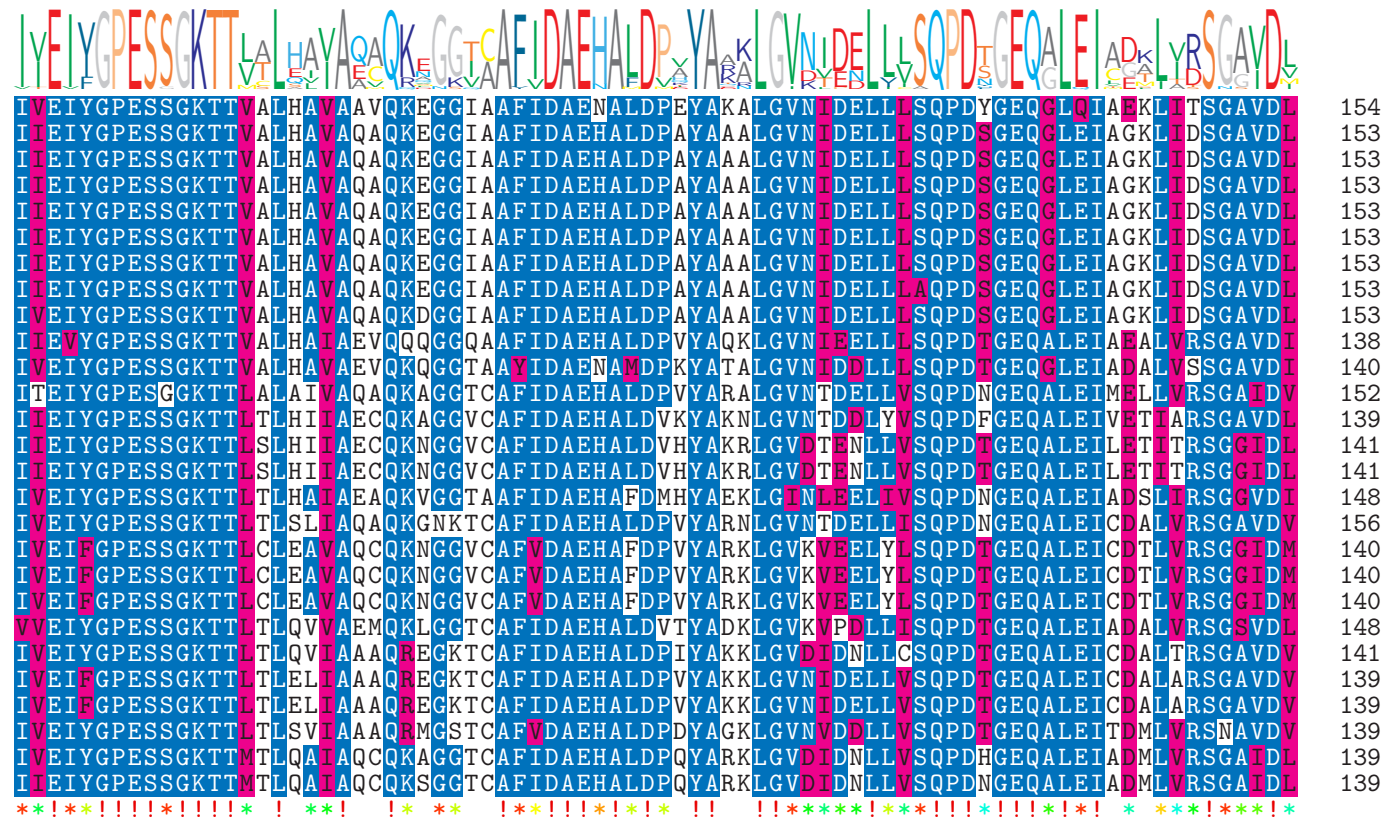
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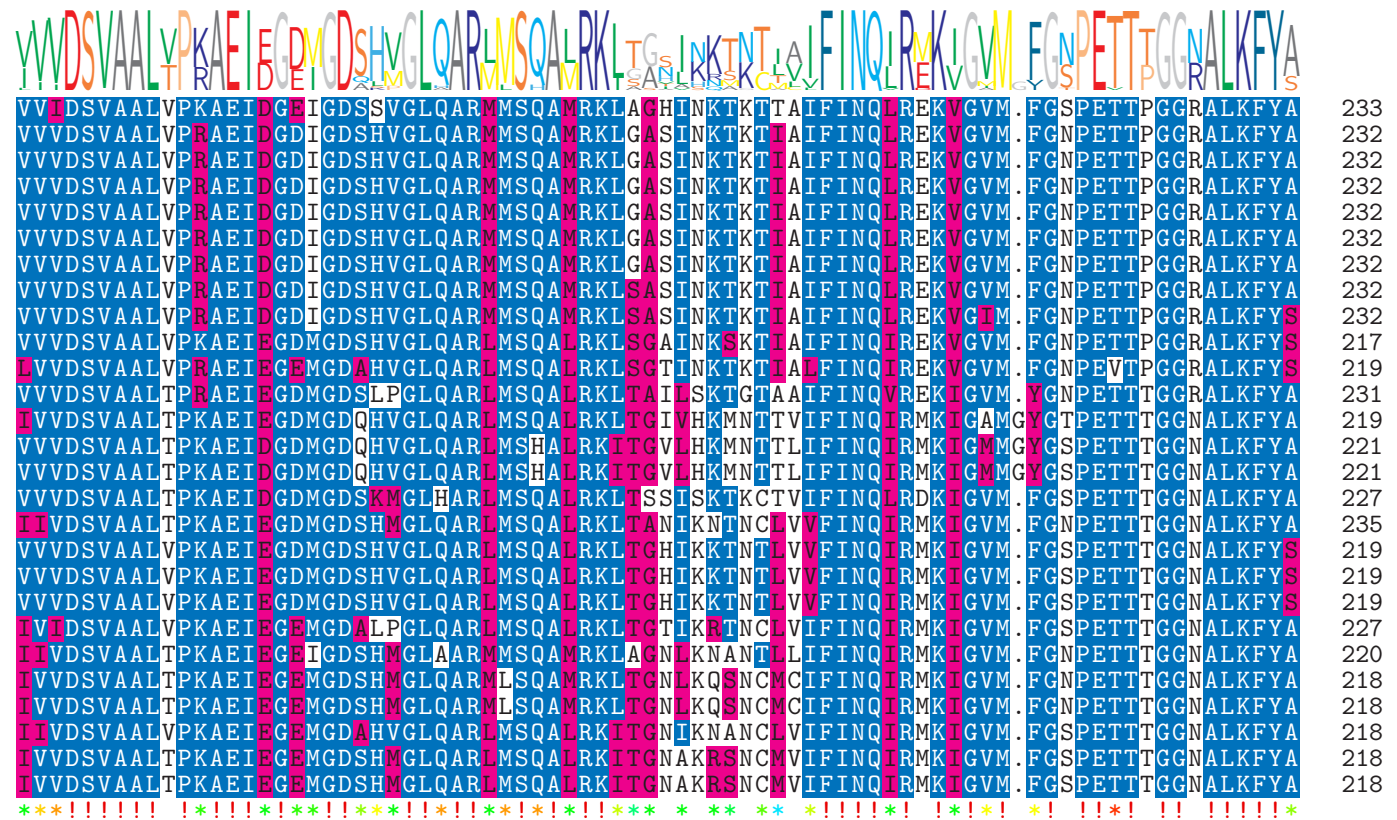
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NTDB id 1022 ACIAD RS06390 WP 0004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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logo

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consensus

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Sequence	Score
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DEKIGQGS ENAKKYLA DNPEIFDEIDHQVRVQFGLIDGEEAAE SKK . DE AAQVDS . . VNEEVTLDLGDELEI	380
GEKIGQGS ENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE E TDDLALDLNGIEI	376
GEKIGQGS ENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK ETDEK . . ADKNLVLELDDTIEL	381
EERLGGG RENAKQFLKENKDIMLMIQE QIREHYGLDNGGVVQQAEETQE . EL . EFE . E	348
SERIGGG RENAKNYLADHEDVEDEVRLKVRAAAYGISDVPEEDLPTTEDE . . QI . NILPDDST EE .	355
DERIGGG KEKTIAYIA ERPMEQEITDRVMAAIRAGNAGEAPALAPAPAA PEEA . E A .	363
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DKKLGQGR ENAKALLKEDKALADEITLKIKESIGSNEEIMP . LPDE PLEEME	347
DKKLGQGR ENAKALLKEDKALADEITLKIKESIGSNEEIMP . LPDE PLEEME	347
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GAKIGQK KDNVRVWLKENPEISDEIDAKIRALNGVEMHITE GTQDET DG . ER PEE .	348
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GERIGQGRD NCREFLRENAELAREIENKVR EHLGVTPMGAV TL . AEEVEED	352
GEKIGQK KANACNFLKENPATIAAELDKKLRDLLLHSGGELV AASGDDFDE A . ET SEQF .	354
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GDKIGQK KANACKYLKENPEIAKTLDKKLRMLLNPNENMQL IAETSSAADVFEFG . AV PEEF .	354
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logo

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NTDB id 1074 ABD1 RS10195 WP 000344167.1		349
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved