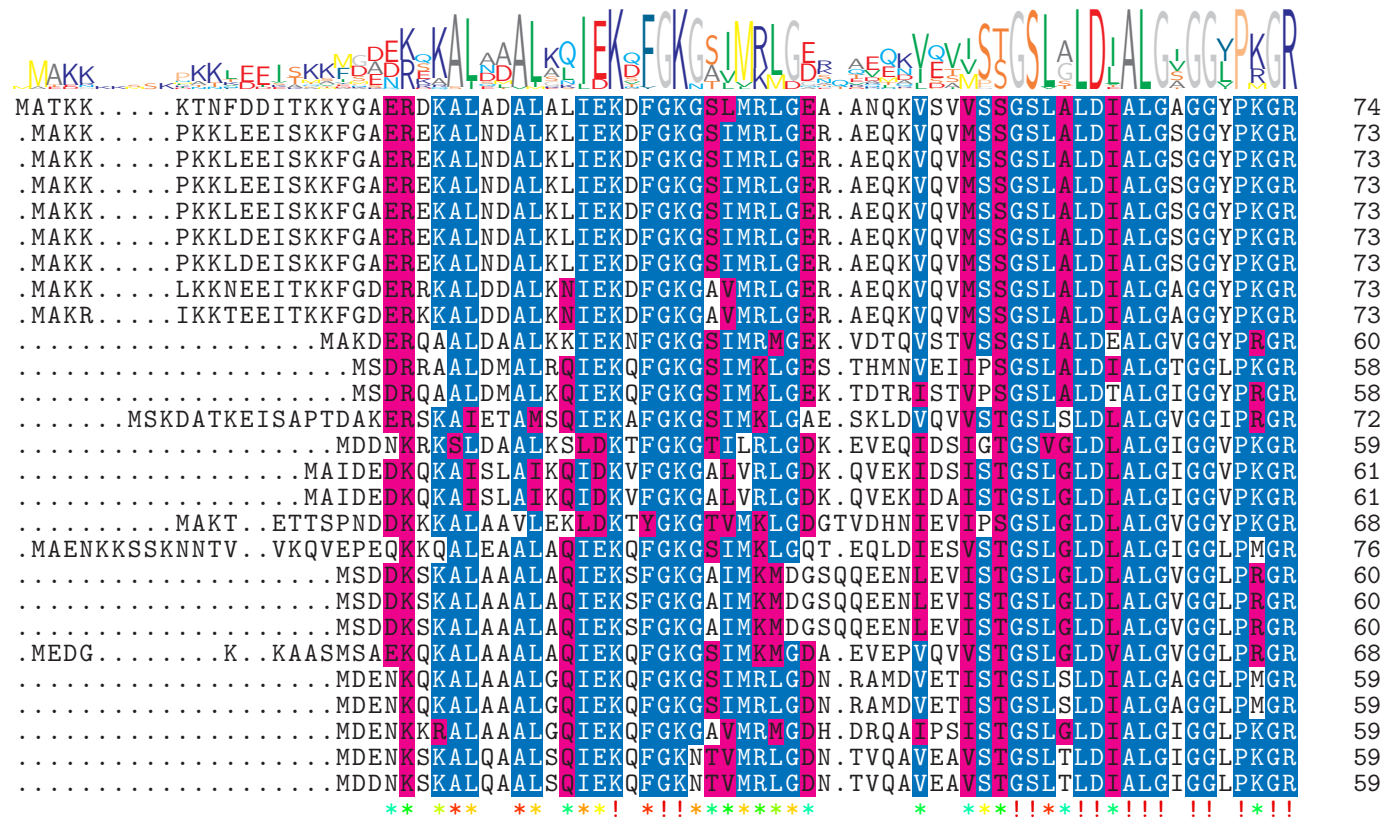


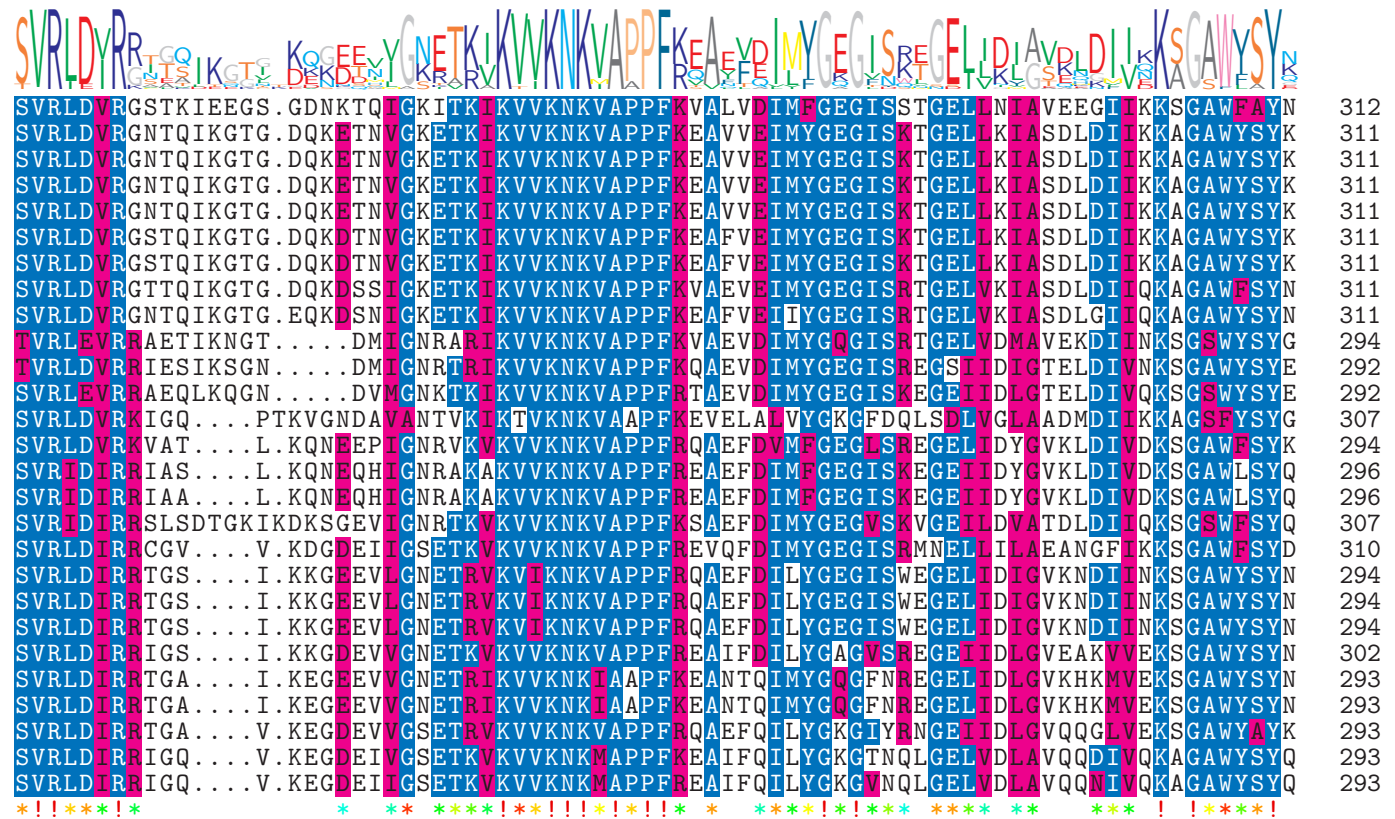
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
consensus

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consensus

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DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE . . . PKKEEA . VNEEVPLDLGDELEIE	385
DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE . . . PKKEEA . VNEEVPLDLGDELEIE	385
DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE . . . PKKEEA . VNEEVPLDLGDELEIE	385
DEKIGQGSENAKKYLAHPEIFDEIDKQVRSKFLIDGEEVSEQDTENKKDE . . . PKKEEA . VNEEVPLDLGDELEIE	385
DEKIGQGSENAKKYLDADNPEIFDEIDHQRVVFGLIDGEEIAES . . . KKDE . . . AAQVDS . VNEEVTLDLGDLEIE	381
DEKIGQGSENAKKYLDADNPEVFAEIDHQRVVFGLIDGEEIAES . . . KKDE . . . VAQVDS . VNEEVTLDLGDLEIE	381
GEKIGQGSENAKRYLADHPQLFEIDHRKVRVKFGLLEESEESAMAVASE E TDDLALDNGIEIE	377
GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK ETDEK . . ADKNLVLELDDTIELE	382
SERIGQGRENAKNYLADHEDVEDEVRLKVRVAAAYGISDVPEEDLPTTEDEQINI . . LPD DSTEE	355
GERLGGGRENAKQFMKEHKDIADITTEKIRVASNLVTAVPAP . . TSEEQKKEA . . . A EEQELFEIN	353
EERLGGGRENAKQFLKENDIMLMIEQIREHYGLDNGVQ . . . Q QA E ETQEELEFE	347
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GAKIGGKDNVVRVWLKENPEISDEIDAKIRALNGVEMH ITEGTQDET DGERPEE	348
GAKIGGKDNVVRVWLKENPEISDEIDAKIRALNGVEMH ITEGTQDET DGERPEE	348
GAKIGGKDNVVRVWLKENPEISDEIDAKIRALNGVEMH ITEGTQDET DGERPEE	348
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consensus			

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