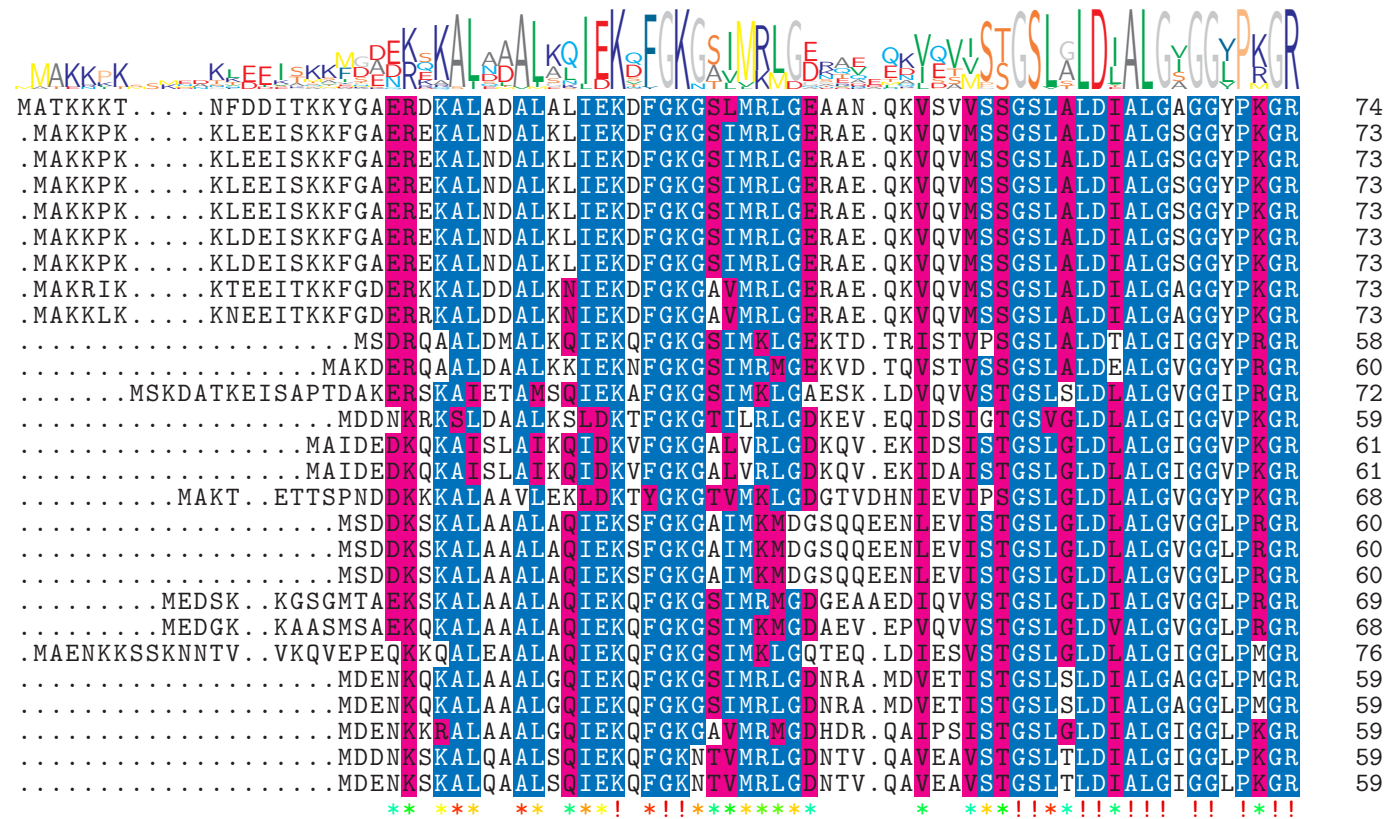


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consensus

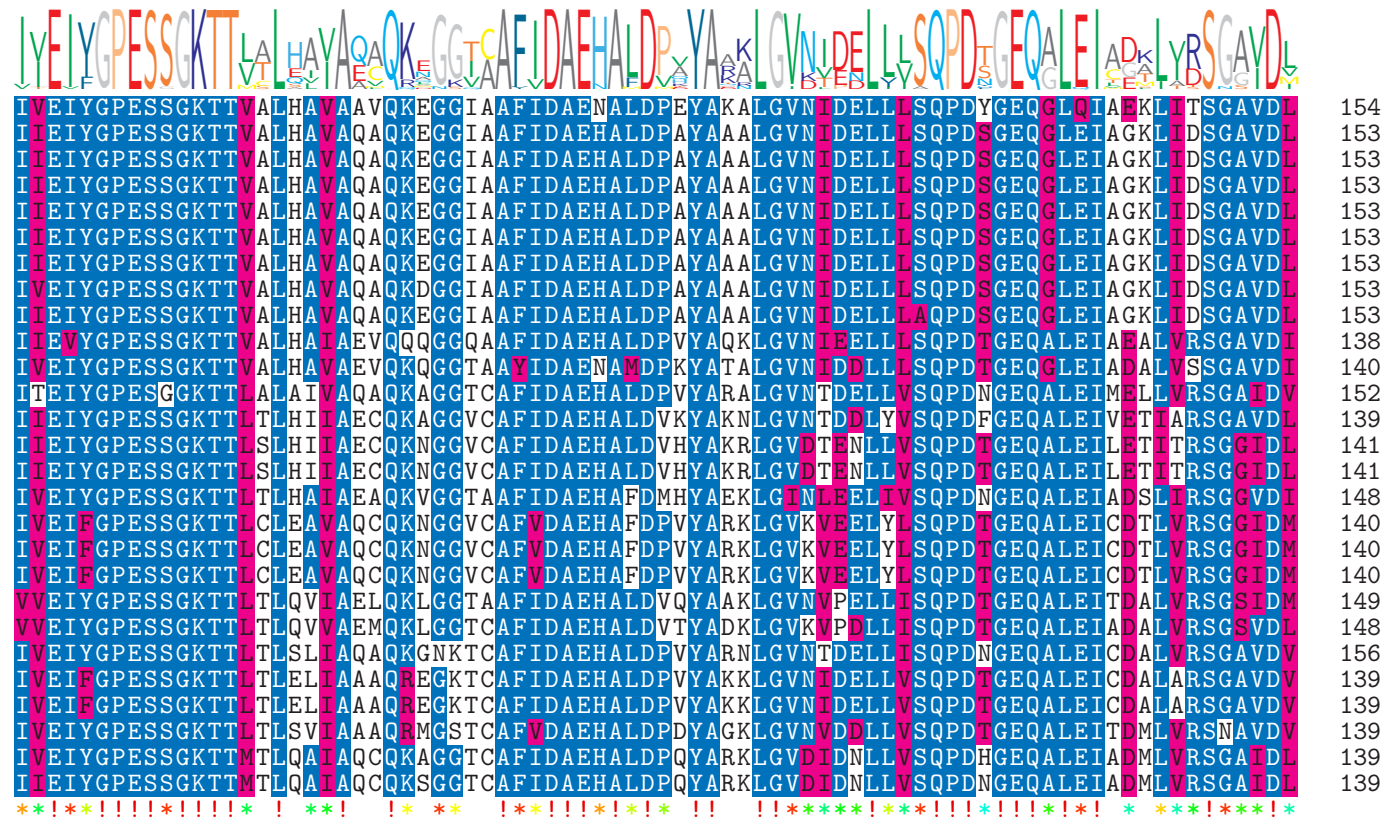
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consensus

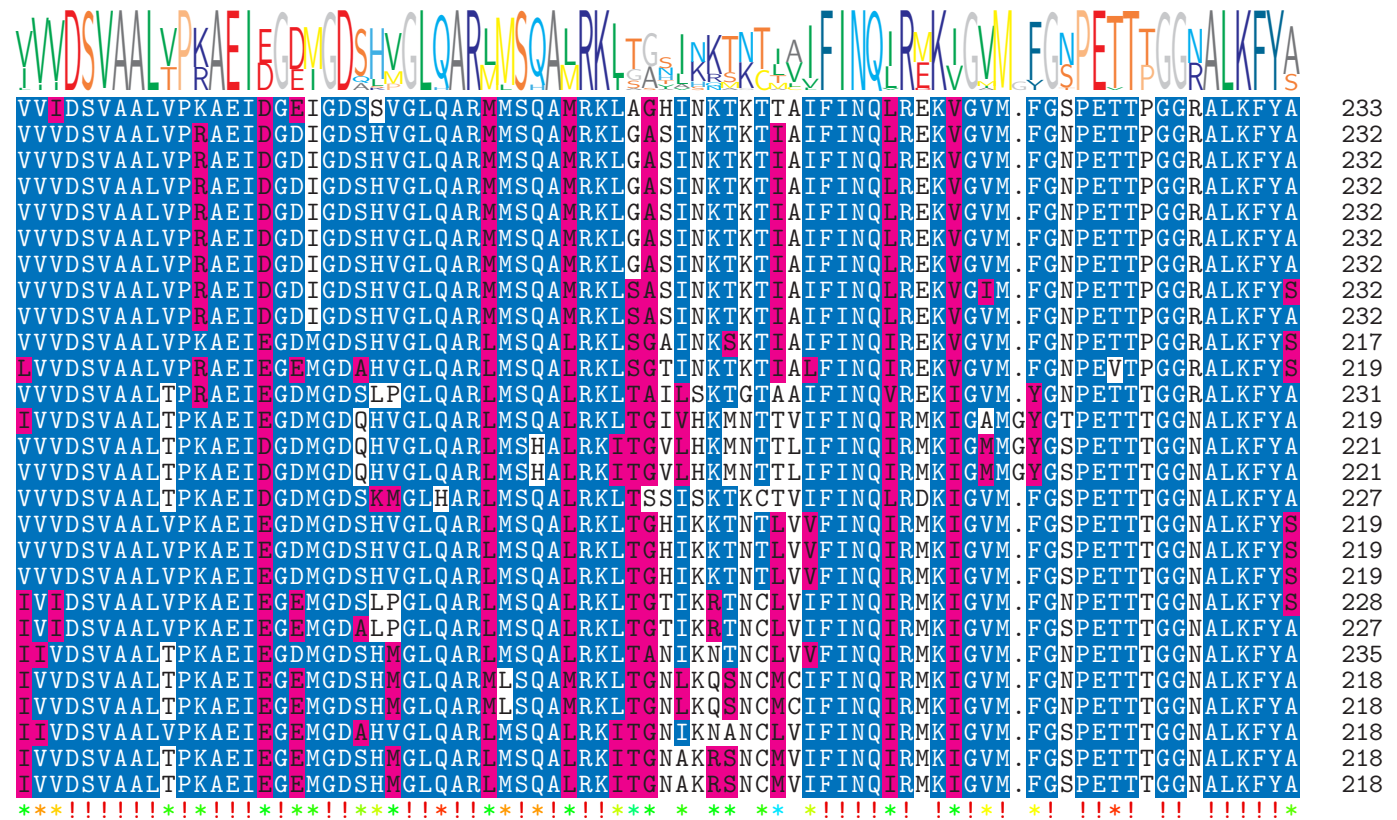
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consensus

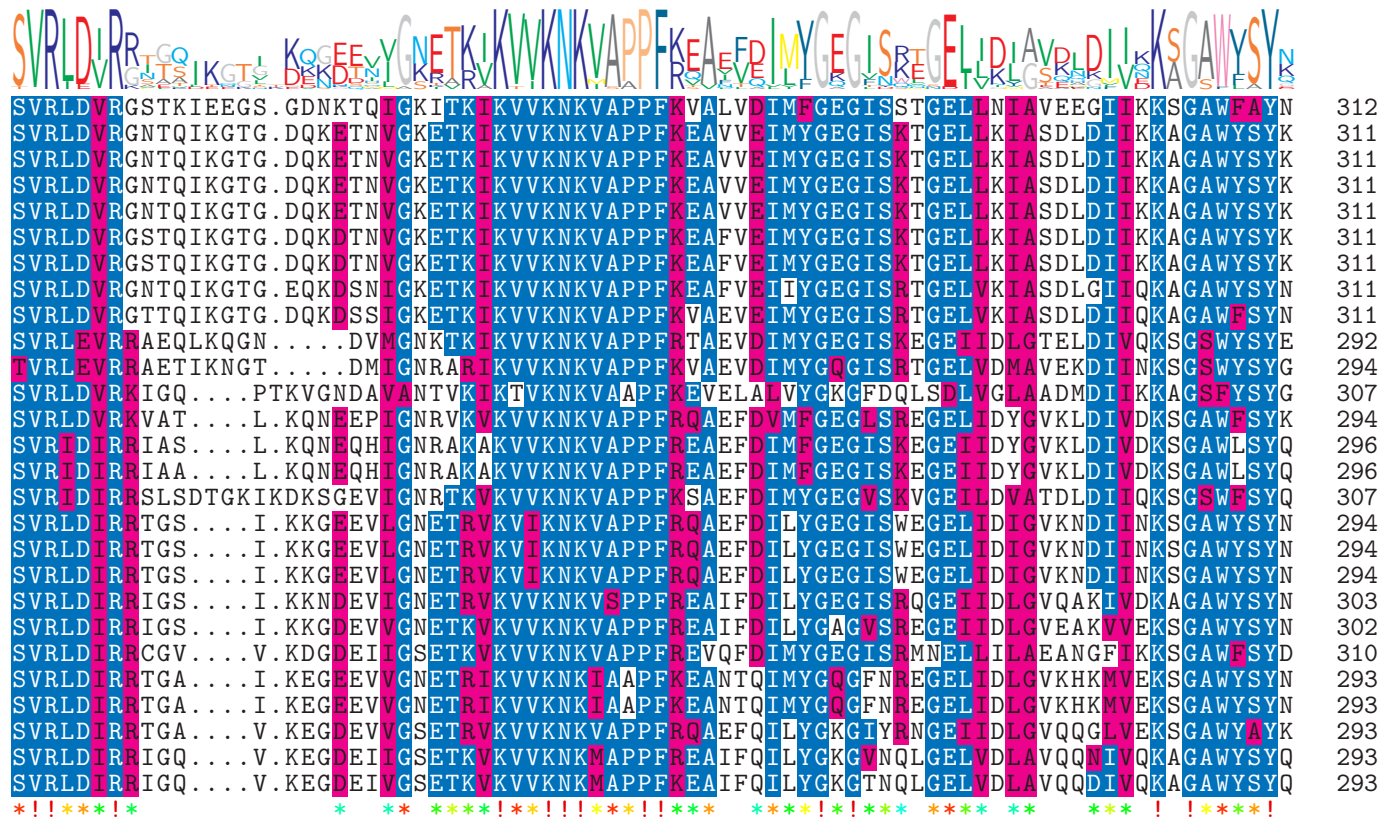
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NTDB id 1022 ACIAD RS06390 WP 004925639.1
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consensus

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consensus

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Sequence	Score
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GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE E TDDLALDLNNGIEIE	377
EERLGGGRENAAKQFLKENKDIMLMIQEQIREHYGLDNGGVVQQAEETQEEL . EFE . E	348
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GERIGGGRDNCREFLRENAELAREIENKVRHGLVTPMGA VT . . LAEE VE ED . . .	352
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GDKIGGQKANACKYLKENPEIAKTLDKKLREMLLNPNENMQ LIAETSSAAD . . DVEFGAVPEEF . . .	354
GSKIGGQKANAAKYLEENPEIGQIEEQIRAKLLVTSANT KATPVAEDLA . . DADI	347
GNIKIGGQKNNVIRYLEENPQISTEIEAVIREQLLTKASDQ TAA . HD . ETEE . . EPDLL . . ES . . .	349
GNKIGGQKNNVIRIFYENNQIAEIEIRNIREQLLTGTNG AVQ . I . EDEE . . EPDLL . . LES . . .	349

logo

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

-  non conserved
-  similar
-  ≥ 50% conserved