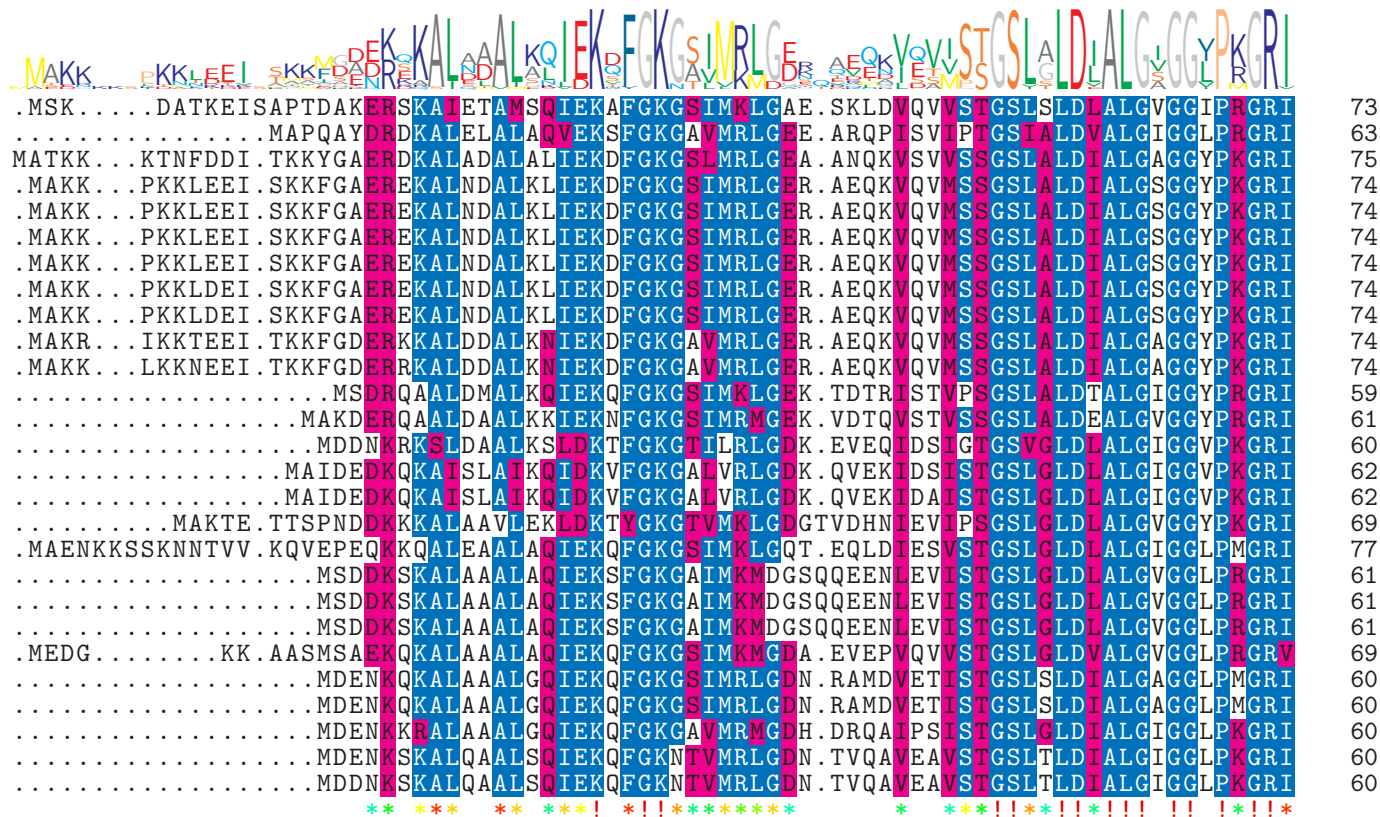


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consensus

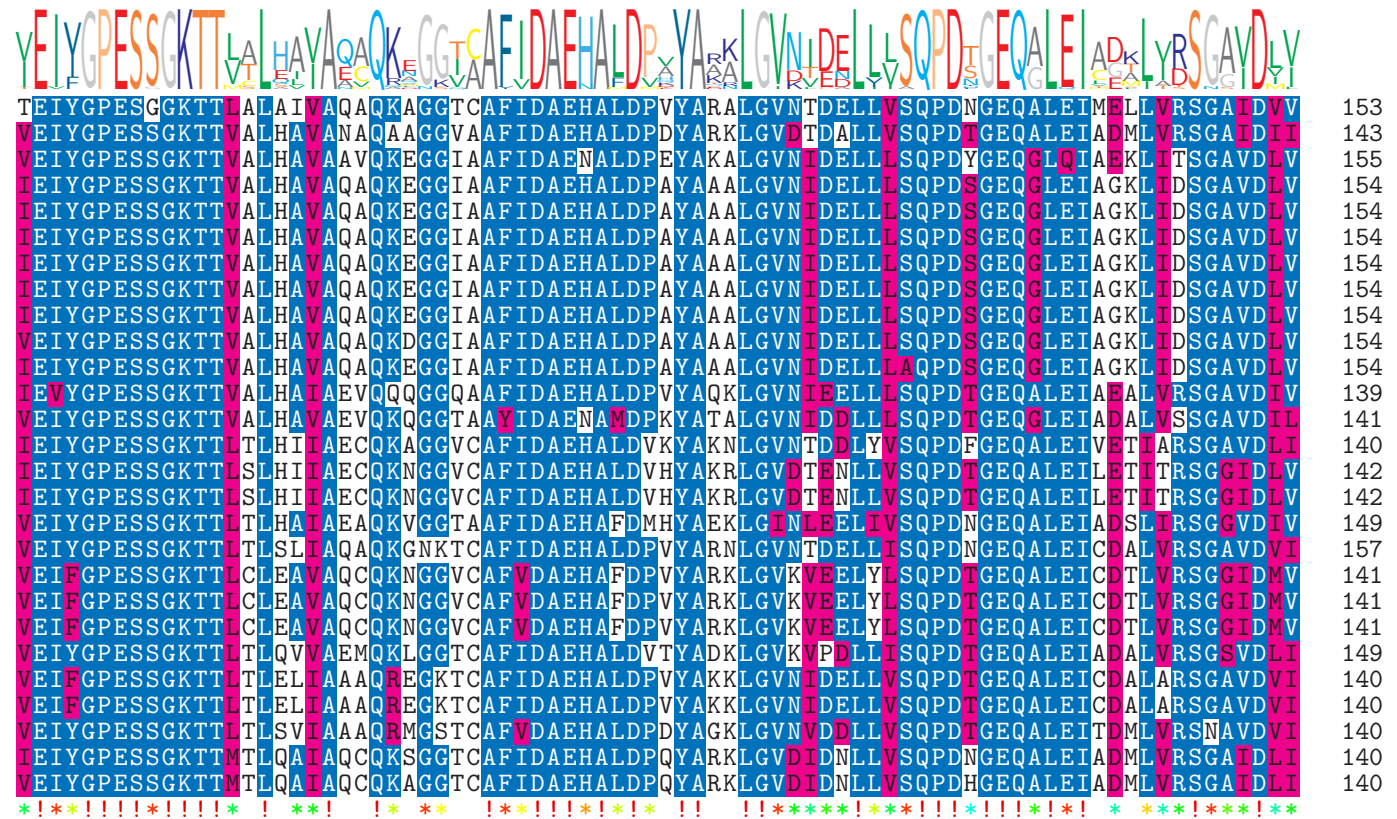
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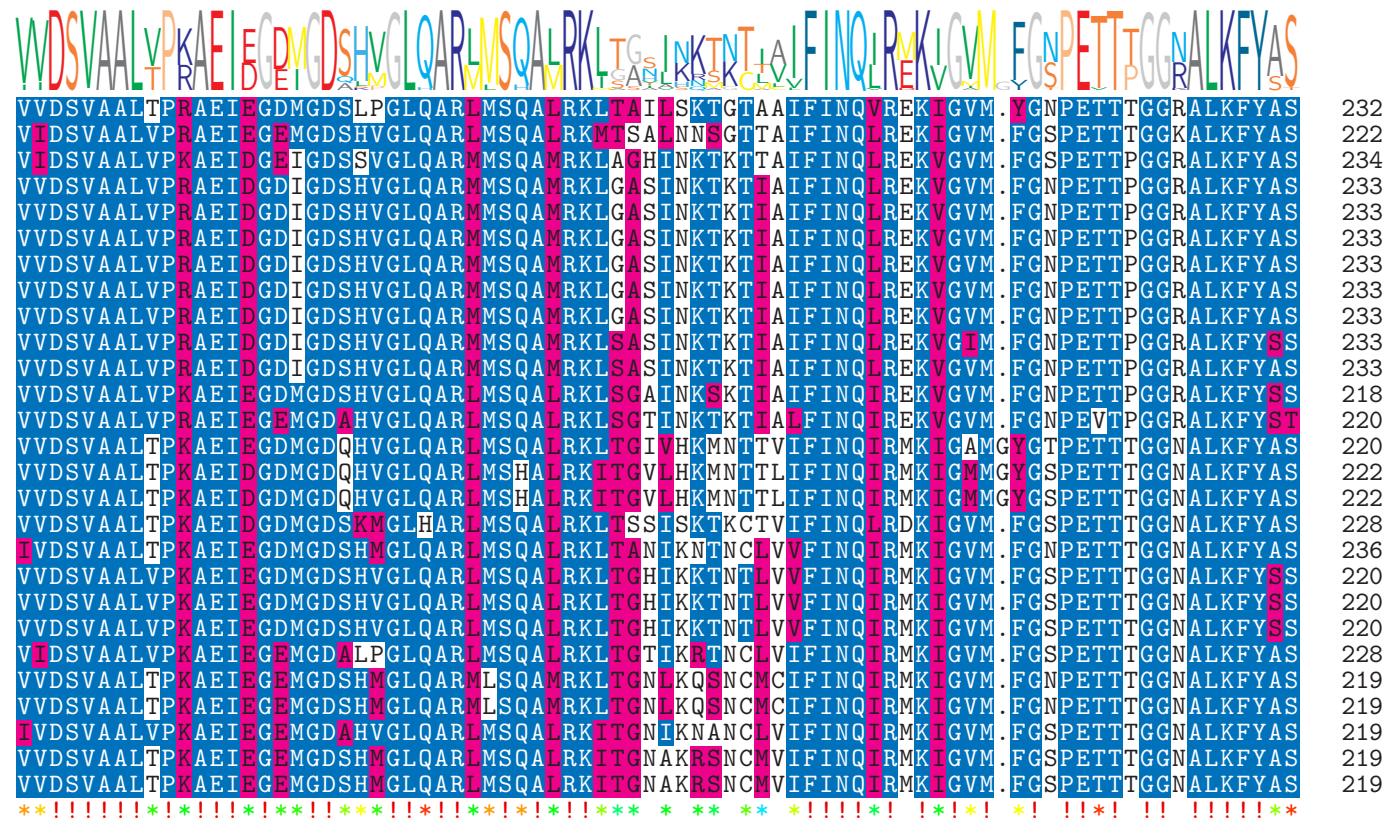
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NTDB id 1022 ACIAD RS06390 WP 004925639.1
consensus

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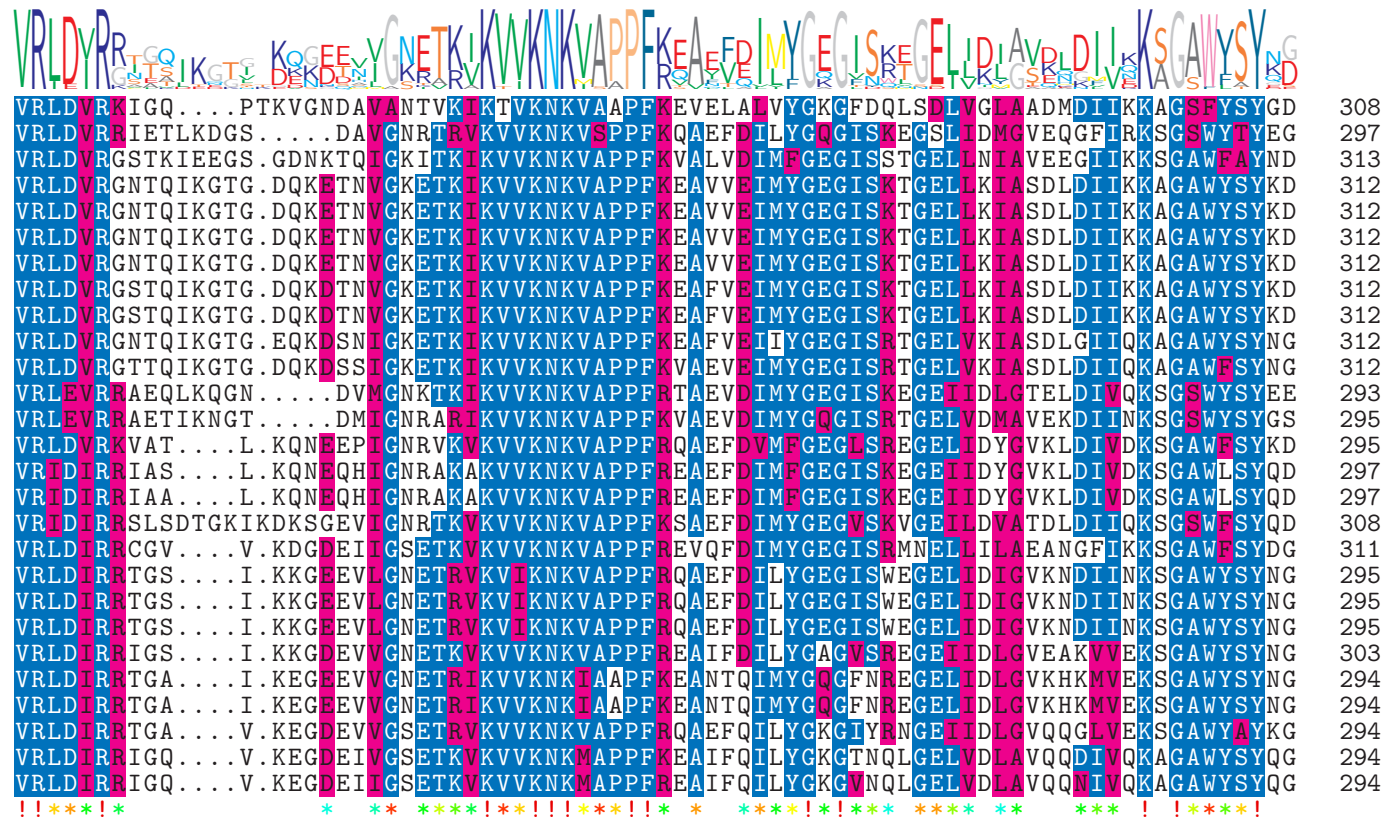
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NTDB id 1022 ACIAD RS06390 WP 004925639.1
consensus

Sequence	Score
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DQLGGGKENARKFLLENTDVRDEIEKKIKELGIGADLTAEDAAEVPAD.F.....	347
EKIGQGAEKAKNYLKEHQDVFDEIDHKVRAAHGLDDSEVAETEETTASKTKAKAKKEEKAVETEEIELELED.....	387
EKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE...PKKEEA..VNEEVPLDLGDELEIEI	386
EKIGQGSSENAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAES...KKDE...AAQVDS..VNEEVTLDLGDELEIEI	382
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ERLGGGRENAKQFLKENKDIMLMIEQIEHYGLDNNGVVQQAAEETQEEL.EFE.E.....	348
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KKLGGGRENKAFLEKENPEIADAITKAIQNSMGIEGMISGSSEDDE..GEE.....	343
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NKIGQGGKNNVIRYFEENNQIAEIEIERNIREQLLTGTGNGAVQ.I.EDEEPPD.....LLES.....	349
NKIGQGGKNNVIRYLEENPQISTEIEAVIREQLLTASDQTAA.HDETEEPPD.....LLES.....	349

logo

NTDB id 1313 DR RS12030 WP 010888966.1	..	363
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NTDB id 600 KW2 RS01765 WP 011675413.1	..	387
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	..	349

consensus

 non conserved
 similar
 ≥ 50% conserved