

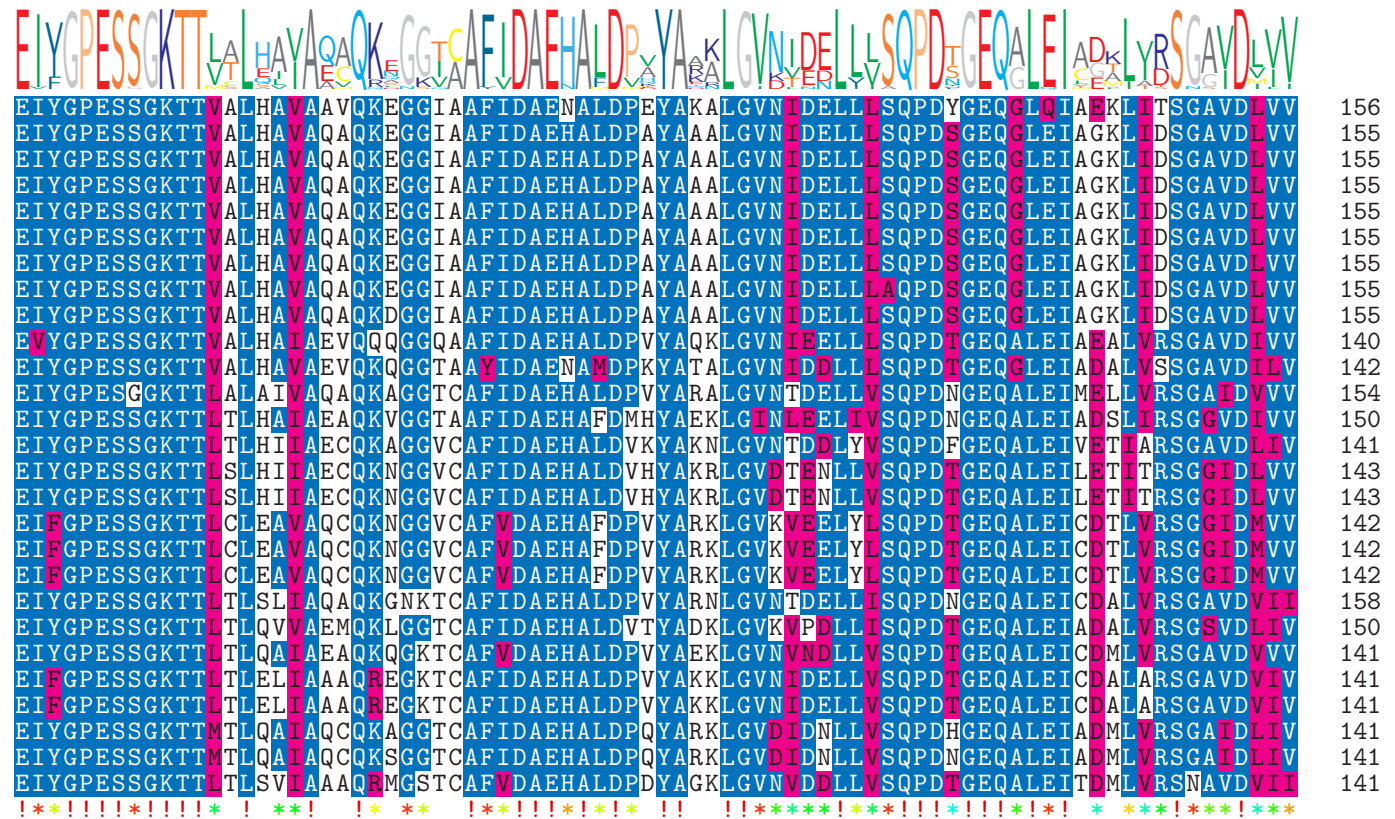
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NTDB id 600 KW2 RS01765 WP 011675413.1	MATKKK...TNFDDITKKYGAERDKALADALALIEKD FGKGSIMRLGER.ANQKVSVVSSGSLALDIALGAGGYPKGRIV	76
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NTDB id 625 LCA RS02525 WP 011374196.1	MAKDERQAALDAALKKIEKNFGKGSIMRMGEK.VDTQVSTVSSGSLALDEALGVGGYPRGRIV	62
NTDB id 1313 DR RS12030 WP 010888966.1	.MSKD....ATKEISAPTDAKERSKATETAMSIIEKAFGKGSIMKLGAE.SKLDVQVVSTGSLSLDLALGVGGIPRGRIT	74
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NTDB id 1245 Cj1673c YP 002345041.1	MDNKRKSLDAALKSLDKTTFGKGTILRLGDK.EVEQIDSIGTGSVGLDLALGIGGVPKGRIT	61
NTDB id 1211 C694 RS00795 WP 000963128.1	MAIDEDKQKATISLAIKQIDKVFVGKGALVRLGDK.QVEKIDSISTGSLGLDLALGIGGVPKGRIT	63
NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDEDKQKATISLAIKQIDKVFVGKGALVRLGDK.QVEKIDAISTGSLGLDLALGIGGVPKGRIT	63
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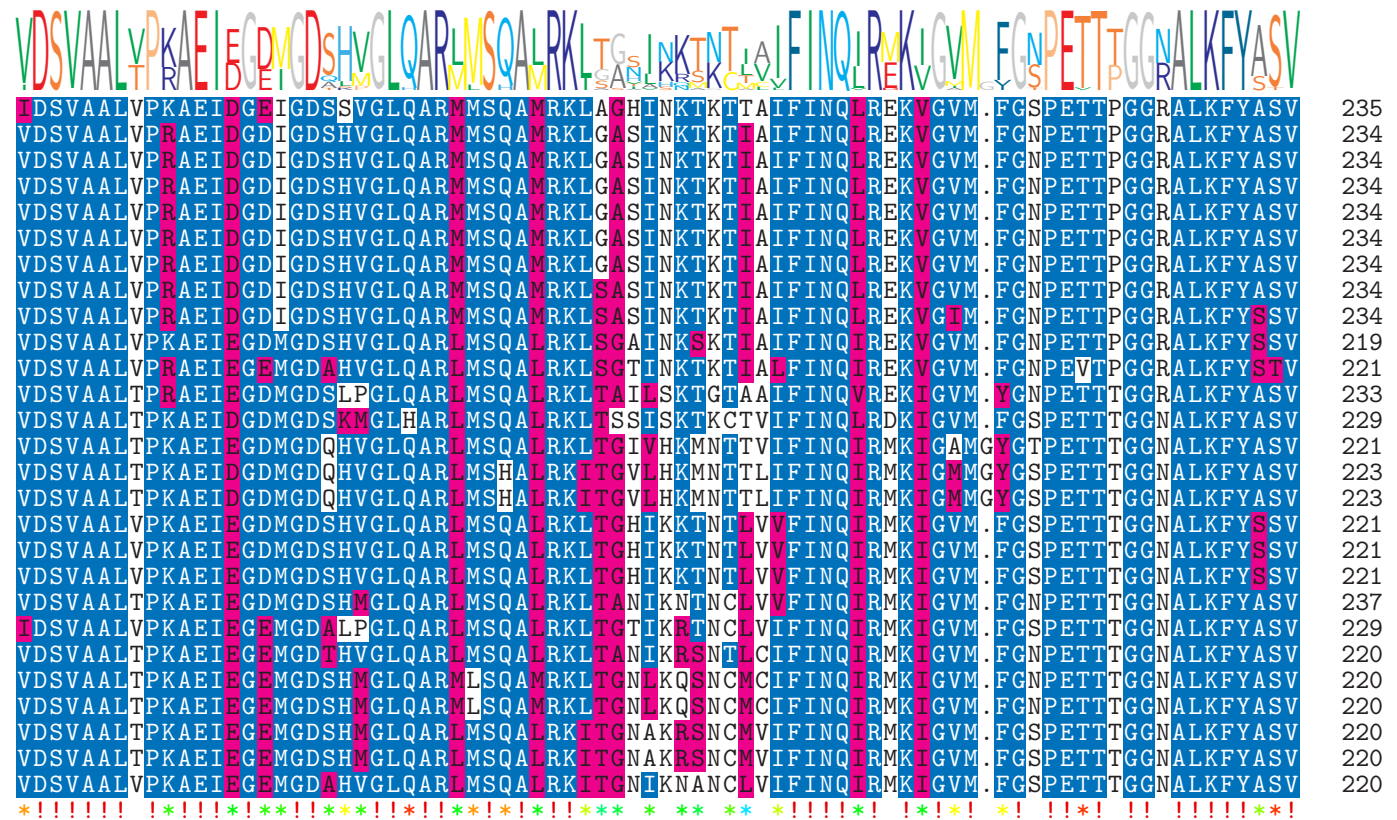
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NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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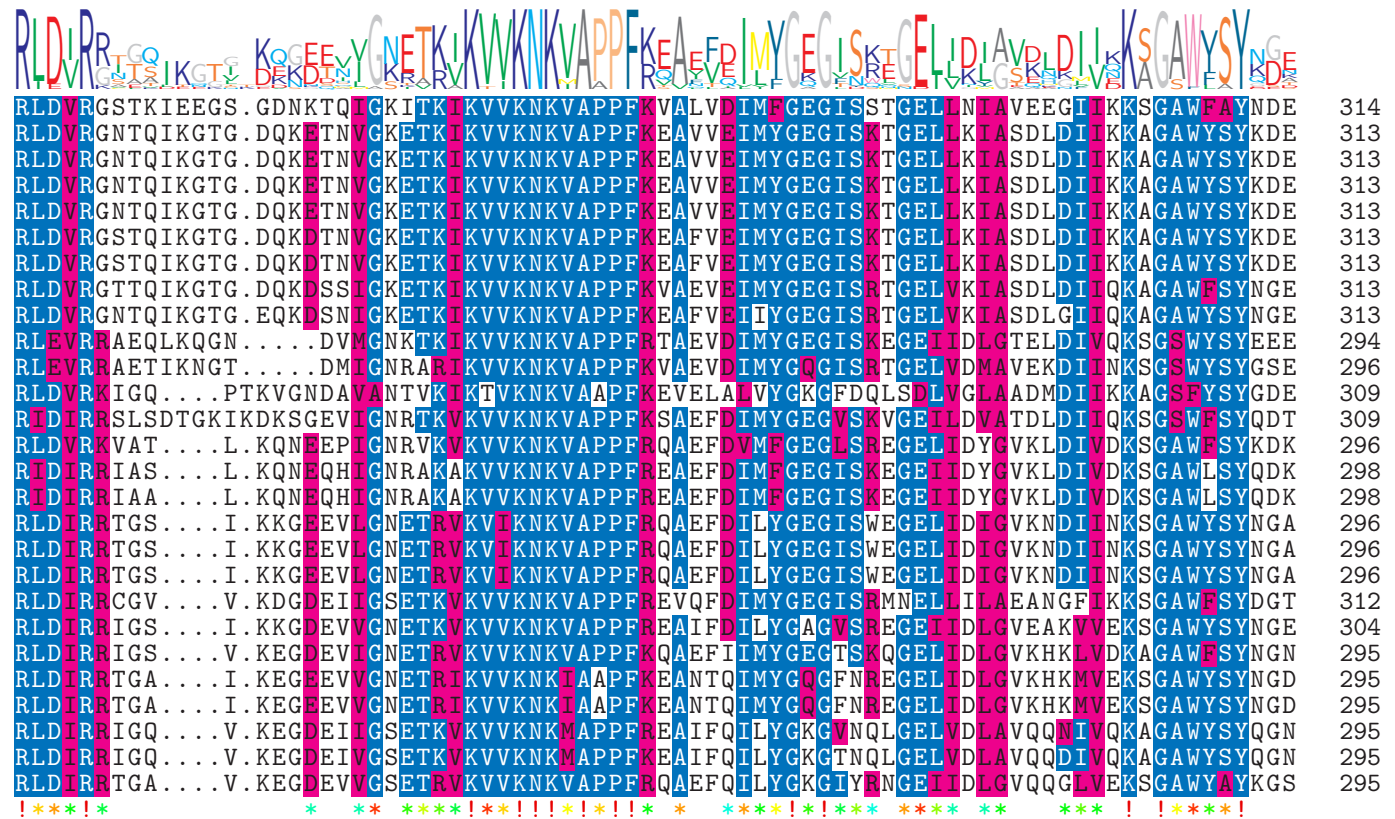
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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KIGQGSENAKKYLAADNPEIFDEIDHQVRVQFGLIDGEEIAES....KKDE....AAQVDS..VNEEVPLDLGDELEIEIE	383
KIGQGSENAKKYLAADNPEVFAEIDHQVRVQFGLIDGEEIAES....KKDE....VAQVDS..VNEEVPLDLGDELEIEIE	383
KIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLDNGIEIED.	378
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RIGQGRENAKNYLADHEDVEDEVRLKVRVAAAYCISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
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KLGGGRENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDEPLEEME.....	347
KIGQGKDNVRVWLKENPEISDEIDAKIRALNCVEMHI....TEGTQDETD.....GERPEE.....	348
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logo

NTDB id 600 KW2 RS01765 WP 011675413.1	.	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	E	388
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consensus

 non conserved
 similar
 ≥ 50% conserved