

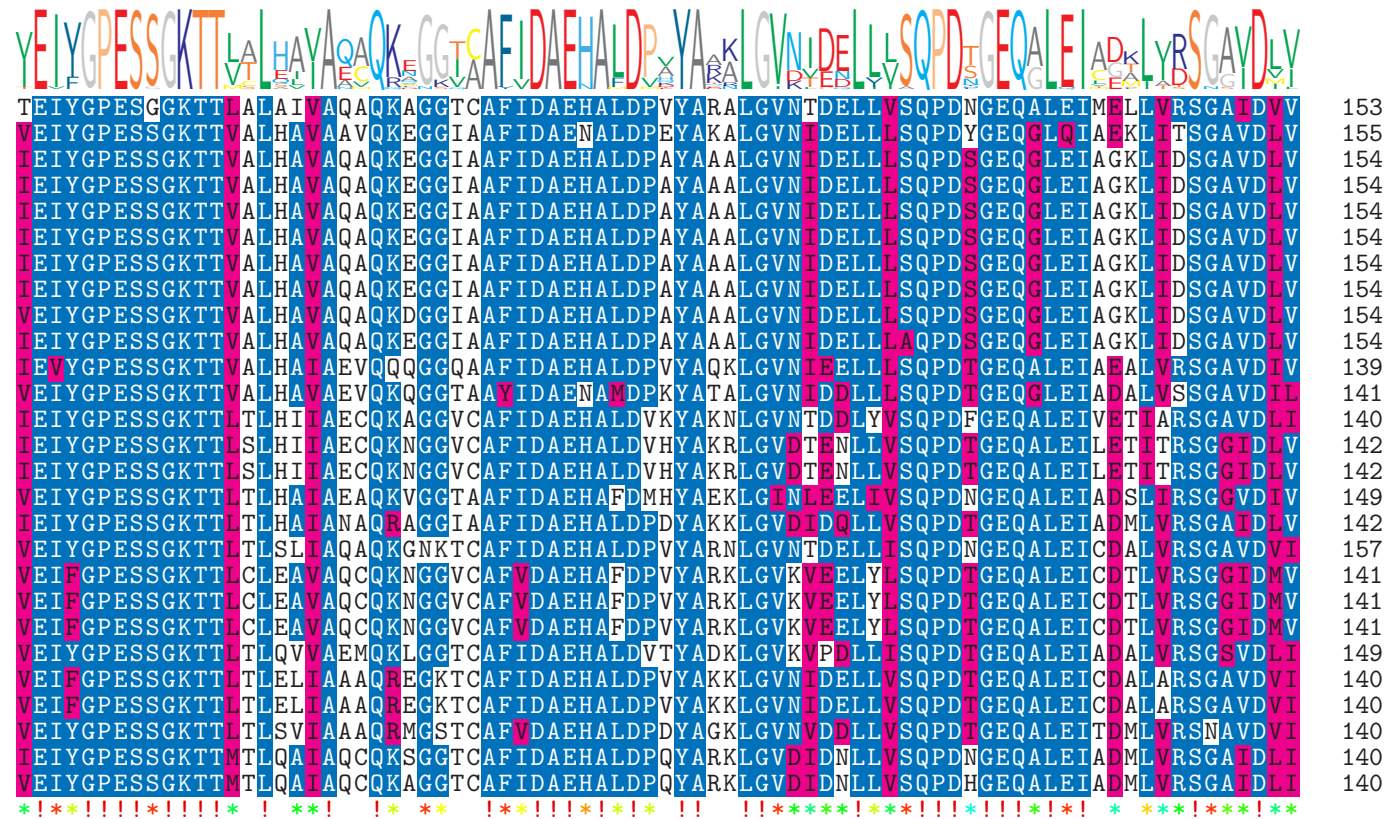
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NTDB id 625 LCA RS02525 WP 011374196.1	MAKD	ERQAALDAALKK	IEKNFGKGSIMRMGEK	.VDTQVSTVSSGSLALDE	ALGVGGYPRGRI	61				
NTDB id 1245 Cj1673c YP 002345041.1	MDDNKRKSLDAALKS	LDKTFGKGTILRLGDK	.EVEQIDSIGTGSVGLDL	LALGIGGVPKGRI		60				
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDE	DKQKAISLA	IKQIDKVFGKGALVRLGDK	.QVEKIDAISTGSLGLDL	LALGIGGVPKGRI	62				
NTDB id 1334 RAOC RS04870 WP 004918236.1	MAKTE.TTSPND	DKKKALAAVLEK	LDKTYGKGTVMKLG	DGTVDHNI	EVIPSGSLGLDLALGVGGYPKGRI	69				
NTDB id 329196 EHF32 RS02045 WP 137416577.1	MPSPA	DREKALESALAL	IDRQFGKGSVMRLGSE	.DRAPVEIPTGSTIALD	VALGVGGLPRGRI	62				
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKSSKNNTVV.KQVEPEQ	KKQALEAALAQ	IEKQFGKGSIMKLGQT	.EQLDIESVSTGSLGLDL	LALGIGGLPMGRI	77				
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	MDDN	KSKALQAALSQ	IEKQFGKNTVMRLGDN	.TVQAVEAVSTGSLTLD	IALGIGGLPKGRI	60				
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NTDB id 194 SPR RS08825 WP 001085462.1
NTDB id 159 SP RS09750 WP 001085462.1
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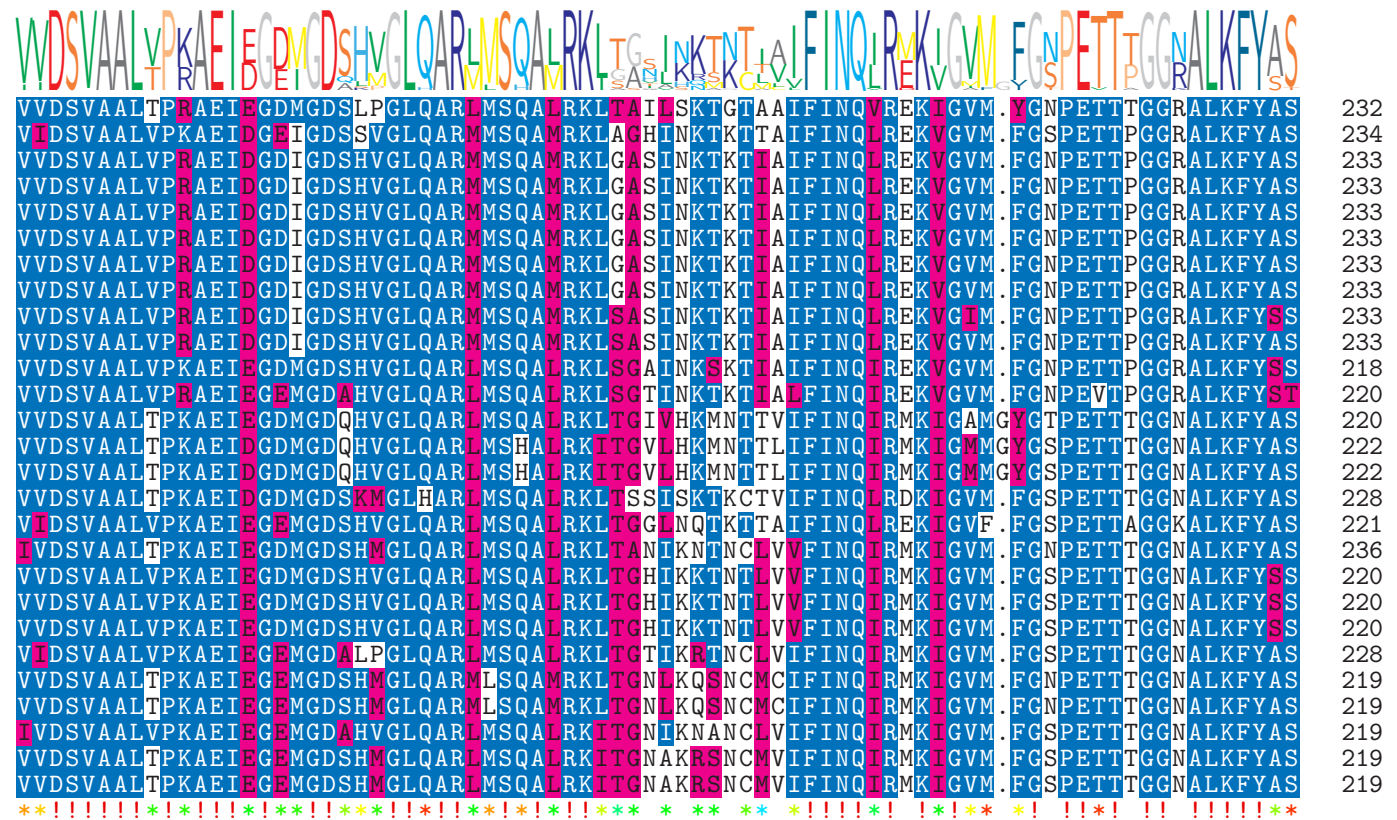
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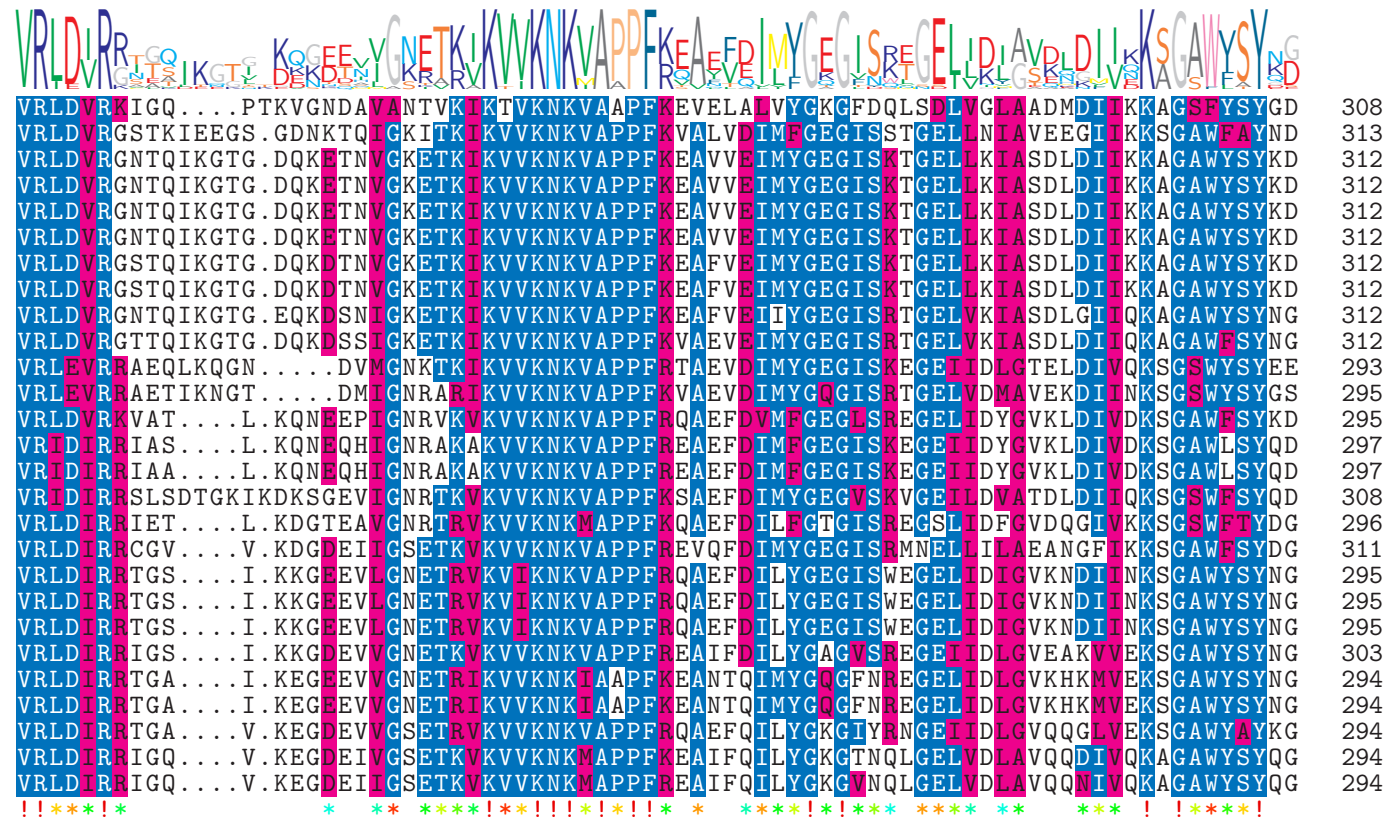
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consensus

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NTDB id 1130 NGFG RS03960 WP 003688695.1	AKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDET.....DG..ERPEE.....	348
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	DKIGQGGKANACKYLKENPEIAKTLDKKLREMLLNPNENMQLIAETSSAADDV.....EFG..AVPEEF.....	354
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NTDB id 1313 DR RS12030 WP 010888966.1	..	363
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consensus

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 ≥ 50% conserved