

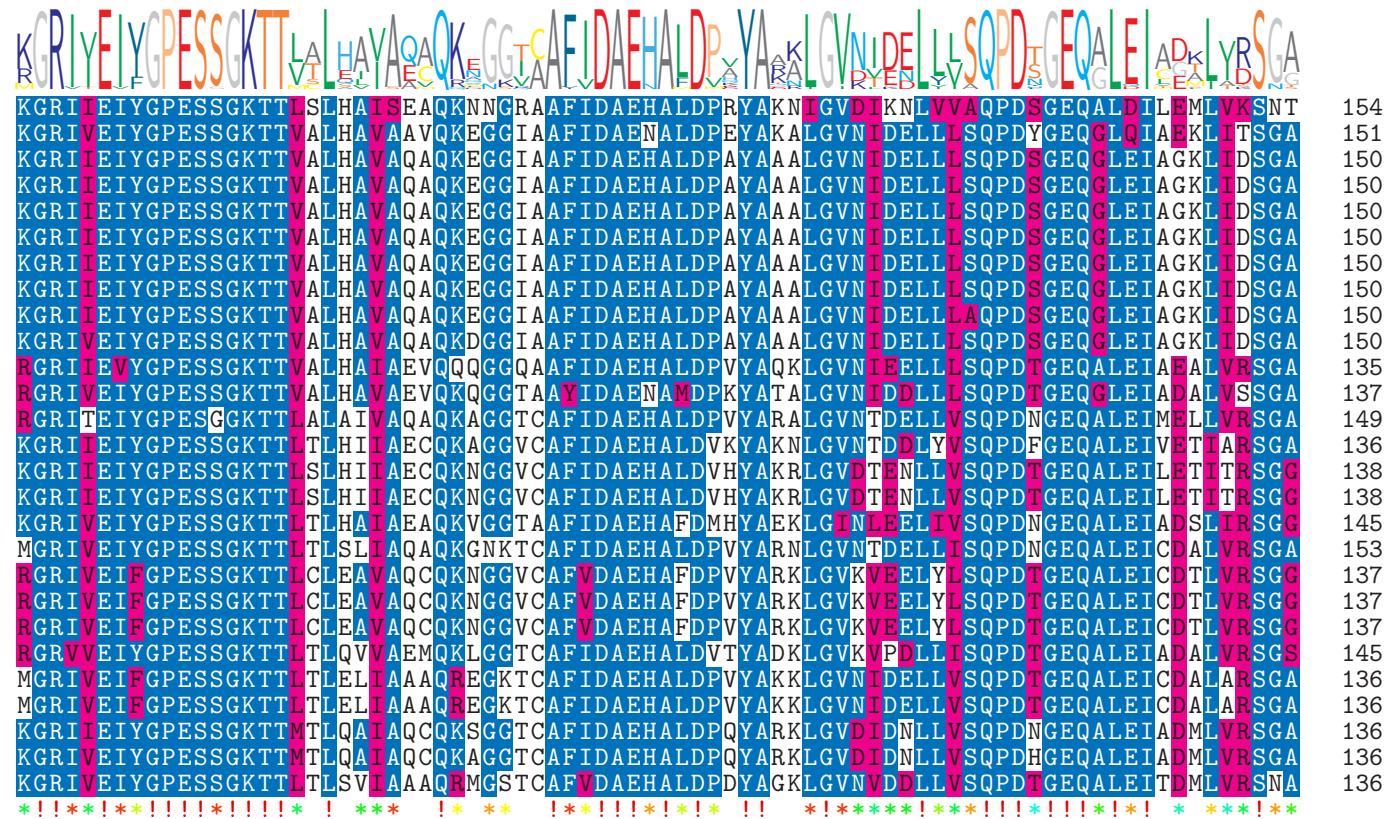
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NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKK....PKKLEEISKKF...GA	EREKALNDALKL	IEKDFGKGSIMRL	GER.AEQKVQ	VMSGSLALDIALGSGGYP	70
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NTDB id 453 Spy49 1753c ACI62003.1	.MAKK....LKKNEEITKKF...GD	ERRKALDDALKN	IEKDFGKGAVMRL	GER.AEQKVQ	VMSGSLALDIALGAGGYP	70
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR....IKKTEEITKKF...GD	ERKKALDDALKN	IEKDFGKGAVMRL	GER.AEQKVQ	VMSGSLALDIALGAGGYP	70
NTDB id 115 BSU 16940 NP 389576.2	MS	DRQAALDMALKQ	IEKQFGKGSIMKL	GEEK.TDTR	ISTVPSGSLALDIALGIGGYP	55
NTDB id 625 LCA RS02525 WP 011374196.1	MA...KD	ERQAALDAALKK	IEKNFGKGSIMRM	GEEK.VDTQ	VSTVSSGSLALDEALGVGGYP	57
NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTD...AK	ERSKATETAMSQ	IEKAFGKGSIMKL	GAE.SKLD	VQVVSTGSLSLDLALGVGGIP	69
NTDB id 1245 Cj1673c YP 002345041.1	M...DDN	KRKSLLDAALKS	LDKTFGKGTILRL	GDK.EVEQ	IDSIGTGSVGLDLALGIGVVP	56
NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAI...DE	DKQKATSLATIKQ	IDKVFVGKALVRL	GDK.QVEK	IDAISTGSLGLDLALGIGVVP	58
NTDB id 1211 C694 RS00795 WP 000963128.1	MAI...DE	DKQKATSLATIKQ	IDKVFVGKALVRL	GDK.QVEK	IDSISTGSLGLDLALGIGVVP	58
NTDB id 1334 RAOC RS04870 WP 004918236.1	MAKT..ETTSP...ND	DKKKALAAVLEK	LDKTYGKGTVMKL	GDTVDHN	IEVIPSGSLGLDLALGVGGYP	65
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKSSKNNTV..VKQVE...PE	QKKQALEAALAQ	IEKQFGKGSIMKL	GQT.EQLD	IESVSTGSLGLDLALGIGGLP	73
NTDB id 1130 NGFG RS03960 WP 003688695.1	M...SD	DKSKALAAALAQ	IEKSFGKGAIMKMD	GSQQUEEN	LEVISTGSLGLDLALGVGGLP	57
NTDB id 1124 NGFG RS03960 WP 003688695.1	M...SD	DKSKALAAALAQ	IEKSFGKGAIMKMD	GSQQUEEN	LEVISTGSLGLDLALGVGGLP	57
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NTDB id 1165 A1552VC RS01565 WP 000344154.1	M...DEN	KQKALAAALGQ	IEKQFGKGSIMRL	GDN.RAMD	VETISTGSLSLDIALGAGGLP	56
NTDB id 1142 GTF74 RS11550 WP 000344154.1	M...DEN	KQKALAAALGQ	IEKQFGKGSIMRL	GDN.RAMD	VETISTGSLSLDIALGAGGLP	56
NTDB id 1074 ABD1 RS10195 WP 000344167.1	M...DEN	KSKALQAALSQ	IEKQFGKNTVMRL	GDN.TVQA	VEAVSTGSLTLDIALGIGGLP	56
NTDB id 1022 ACIAD RS06390 WP 004925639.1	M...DDN	KSKALQAALSQ	IEKQFGKNTVMRL	GDN.TVQA	VEAVSTGSLTLDIALGIGGLP	56
NTDB id 1205 PSJM300 05545 AFN77183.1	M...DEN	KKRALAAALGQ	IEKQFGKGAVMRM	GMDH.DRQA	IPSISTGSLGLDIALGIGGLP	56
consensus			*** **	*** **	*** **	

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NTDB id 159 SP RS09750 WP 001085462.1
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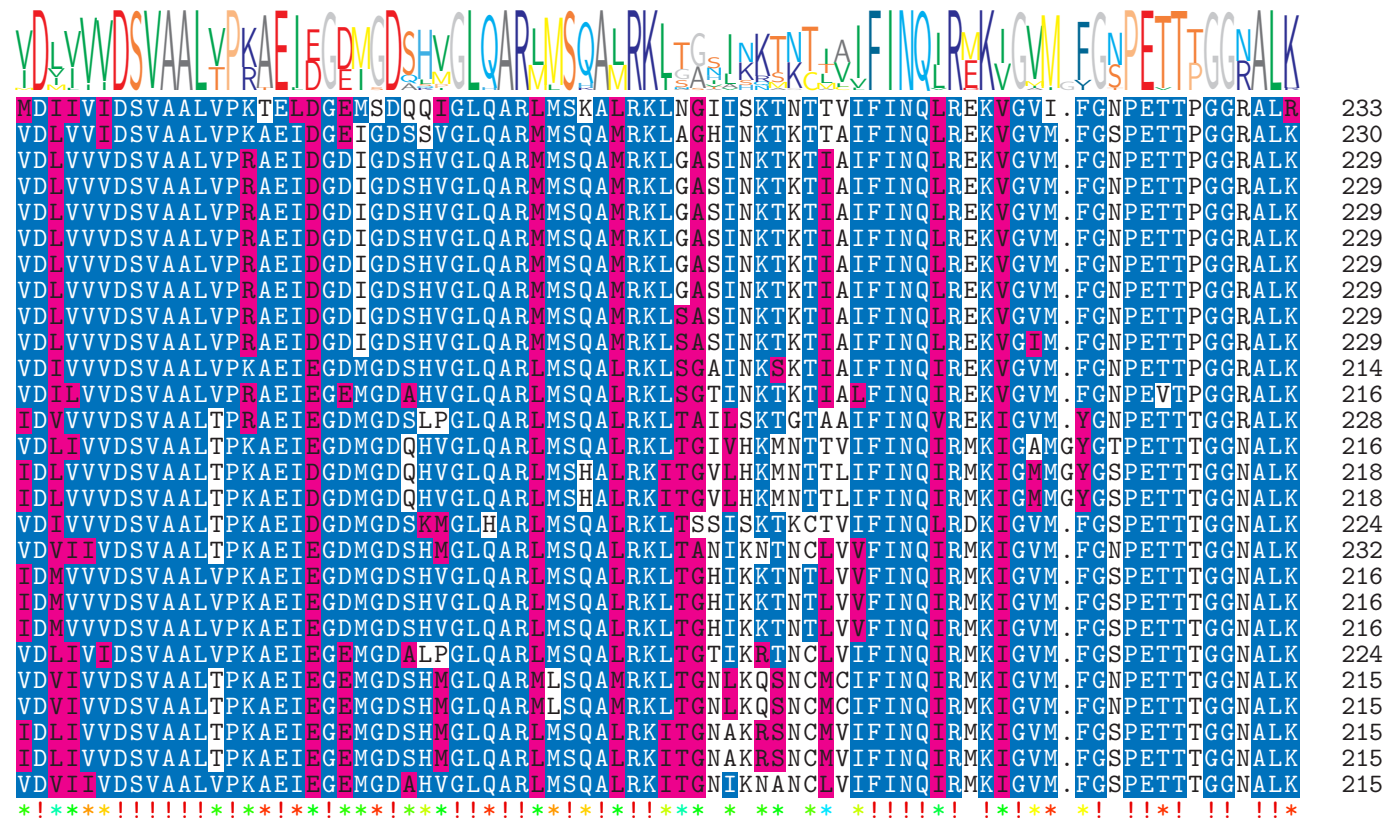
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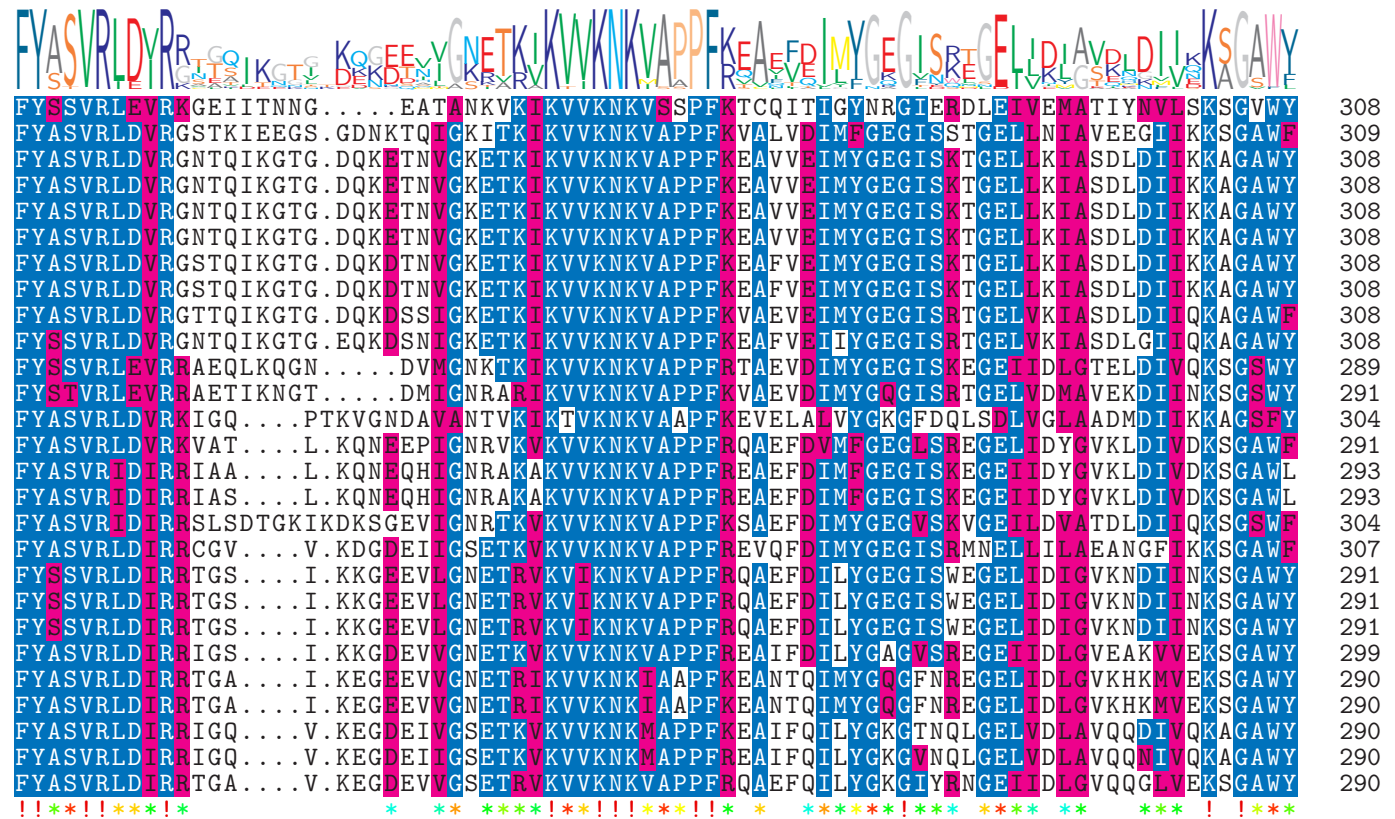
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consensus

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NTDB id 1123 OK783 RS03845 WP 003688695.1	SYNGAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI....TEGTQDET.....GERPEE.....	348
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	SYNGDKIGQGGKANACKYLLKENPEIAKTLDKKLREMLLNPNM....QLIAETSSAADDVE..FGAVPEEF.....	354
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consensus			

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