

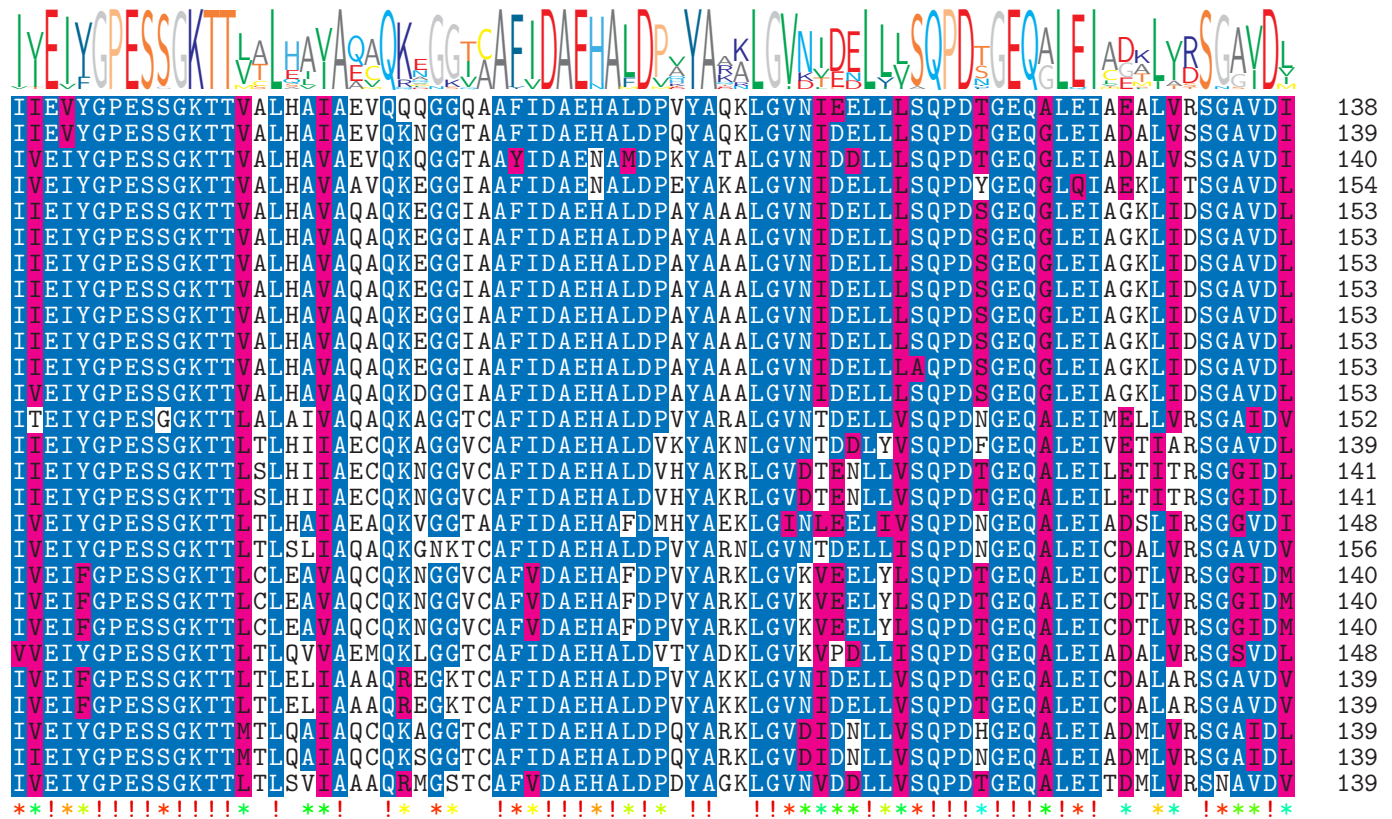
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NTDB id 453 Spy49 1753c ACI62003.1	.MAK.....	KLKKNEEITKKFG	DERR	KALDDALKN	IEKD	FGKGAVMRLGER	.AEQKV	QVMSS	SGSLALD	TALGAGGYPKGR	73
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NTDB id 1245 Cj1673c YP 002345041.1	MD	DNKRKSLDAALKS	LDKT	FGKGTILRLGDK	.EVEQ	IDSIG	TGSVGLD	LALGI	GGVPRGR	59	
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NTDB id 1123 OK783 RS03845 WP 003688695.1	MS	DDKSKALAAALAQ	IEKS	FGKGAIMKMDGS	QREEN	LEVIST	TGSLGLD	LALGV	GGLPRGR	60	
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....	K..KAASMSAEKQ	KALAAALAQ	IEKQ	FGKGSIMKMGDA	.EVEP	VQVST	TGSLGLD	VALGV	GGLPRGR	68
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	MD	ENKSKALQAALSQ	IEKQ	FGKNTVMRLGDN	.TVQ	AVEAV	STGSLTLD	I	ALGI	GGLPKGR	59
NTDB id 1205 PSJM300 05545 AFN77183.1	MD	ENKKRALAAALGQ	IEKQ	FGKGAVMRMGDH	.DRQA	IPSIS	TGSLGLD	I	ALGI	GGLPKGR	59
consensus		*** **	*	***	***	*****	*	*****	*****	***	

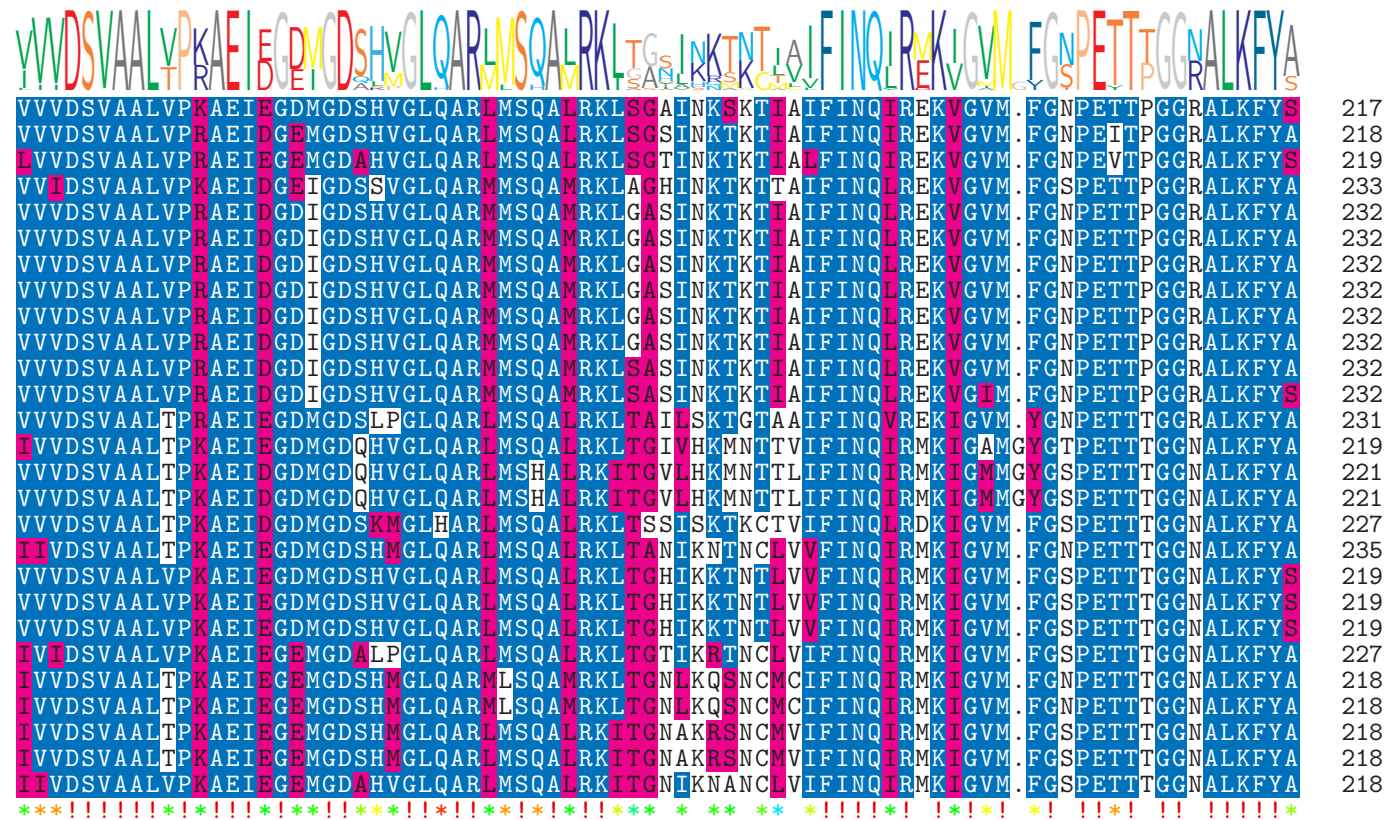
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NTDB id 1313 DR RS12030 WP 010888966.1
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consensus

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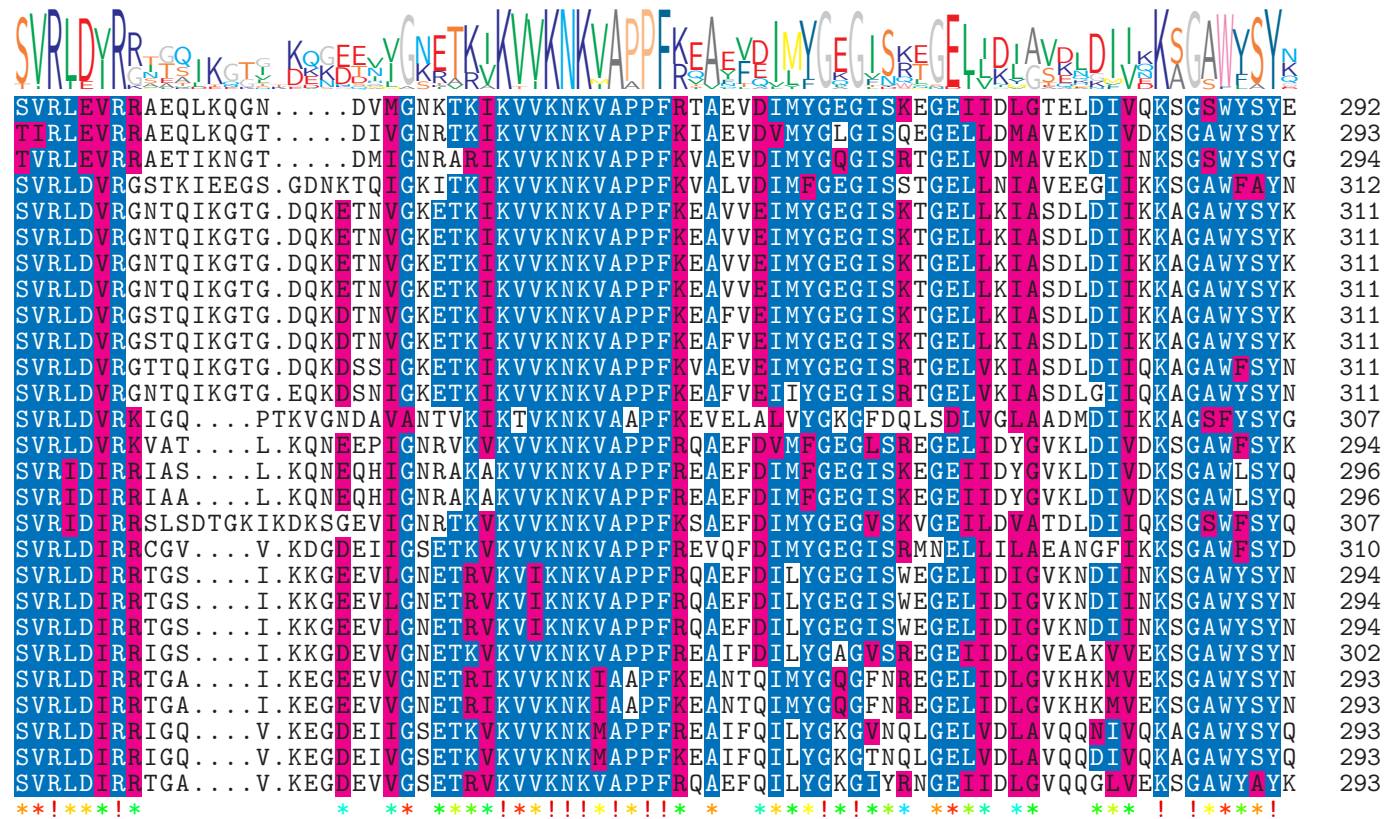
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consensus



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consensus

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NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGSENAkkYLAEHPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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consensus	***!!!***** ** * *** ***** ** **	

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consensus			

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