

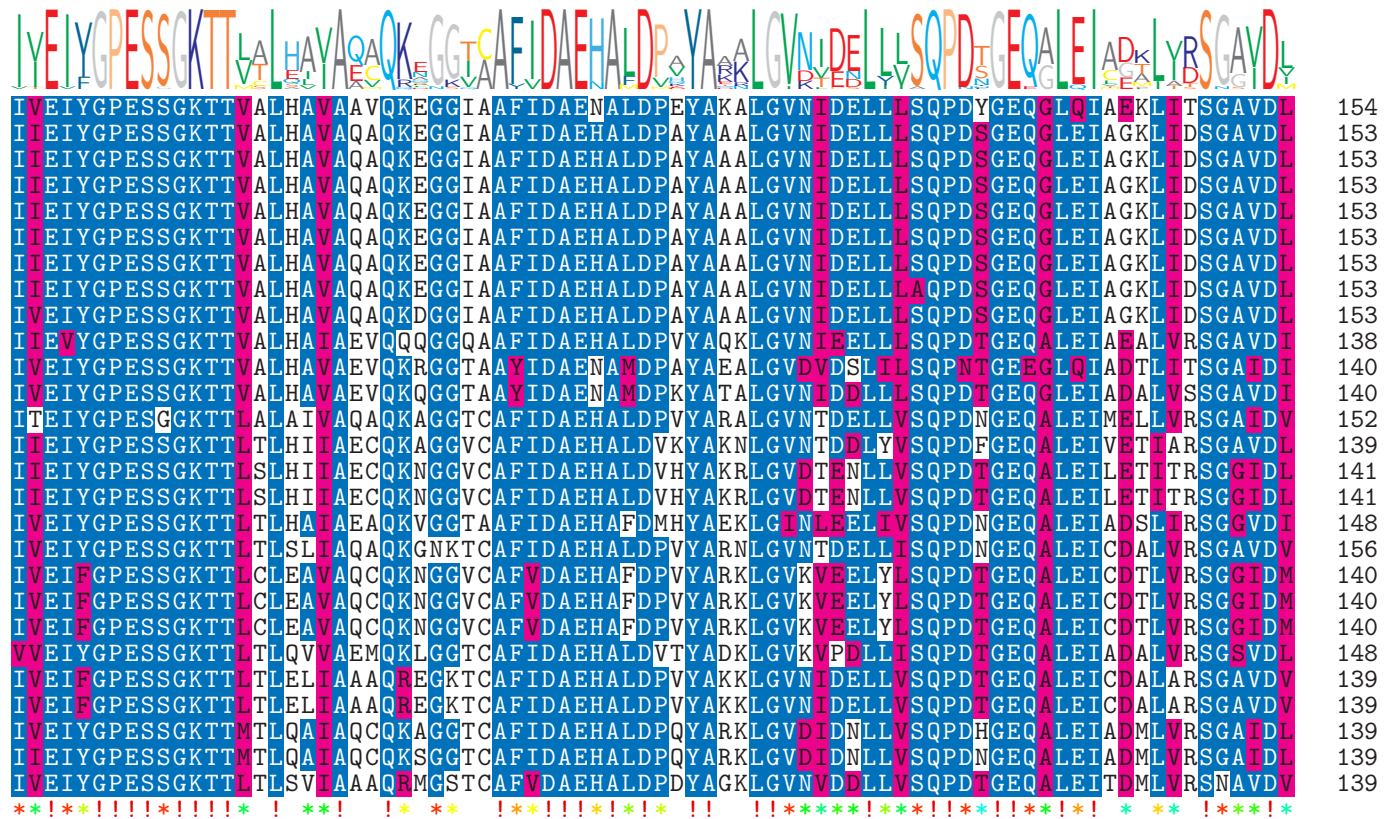
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NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR.....IKKTEEITKKFG	DERRKALDDALKN	IEKD	FGKGAVMRLGER	.AEQKVQVMSSGSLALD	I	ALGAGGYPKGR	73
NTDB id 115 BSU 16940 NP 389576.2	MSDRQAALDMALKQ	IEKQFGKGSIMKLGEK	.TDTRISTVP	SGSLALD	TALGIGGYPRGR			58
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NTDB id 625 LCA RS02525 WP 011374196.1	MAKDERQAALDAALKK	IEKNFGKGSIMRMGEK	.VDTQVSTVSSGSLALDEALGVGGYPRGR					60
NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTDAKERSKAT	IETAMSQIEKA	FGKGSIMKLGA	E.SKLDVQVVSTGSLSLDLALGVGGIPRGR				72
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDEDKQKATISLATIKQ	IDKVFVGKGALVRLGDK	.QVEKIDISTGSLGLDLALGIGGVPRGR					61
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKSSKNNTV..VKQVEP	EQKKQALEAALAQIEKQ	FGKGSIMKLQGT	.EQLDIESVSTGSLGLDLALGIGGLPMGR				76
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NTDB id 1124 NGFG RS03960 WP 003688695.1	MSDDKSKALAAALAQIEKS	FGKGAIMKMDGSQ	QEEENLEVISTGSLGLDLALGVGGLPRGR					60
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NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQ	FGKGAVMRMGDH	.DRQAIPSISTGSLGLDI	ALGIGGLPKGR				59
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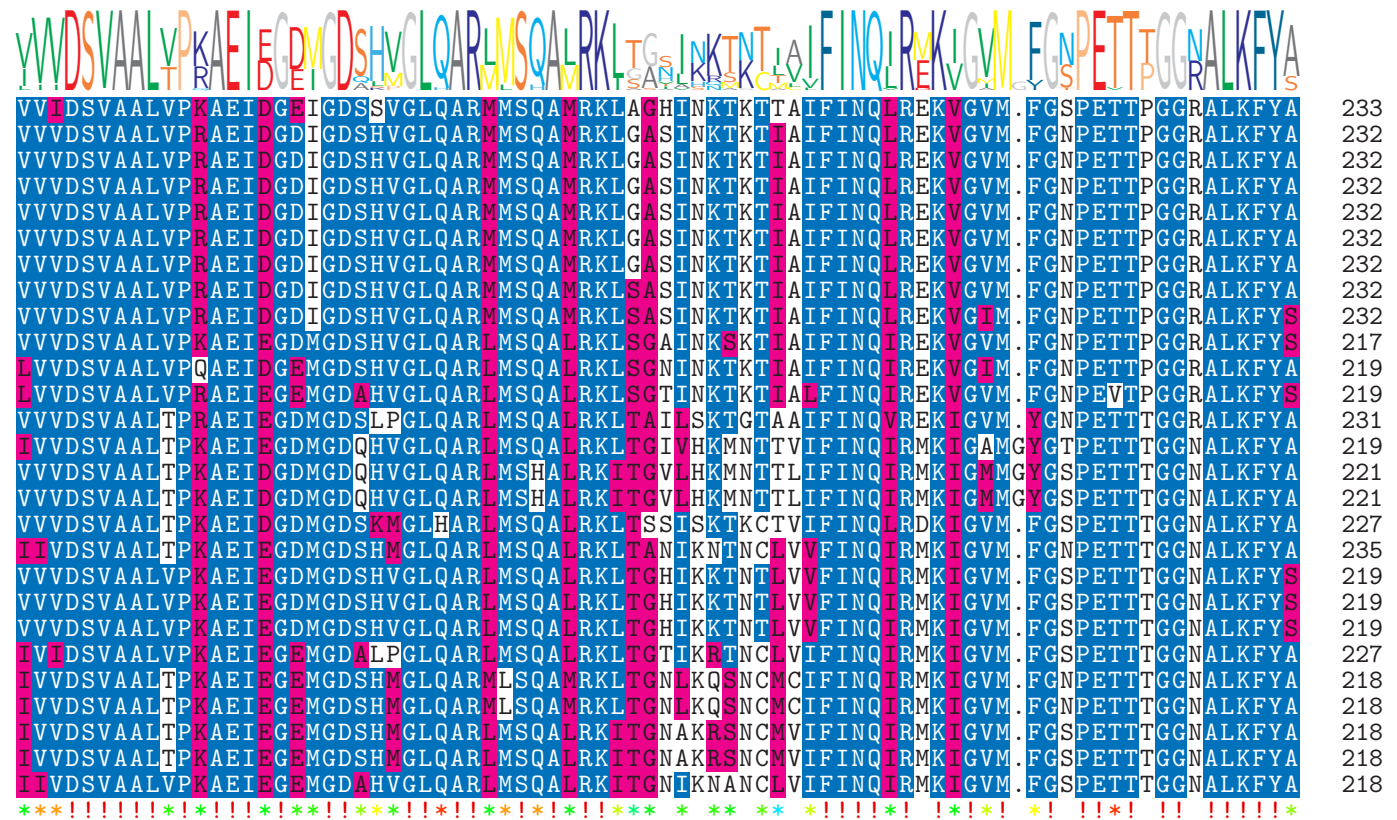
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NTDB id 1205 PSJM300 05545 AFN77183.1
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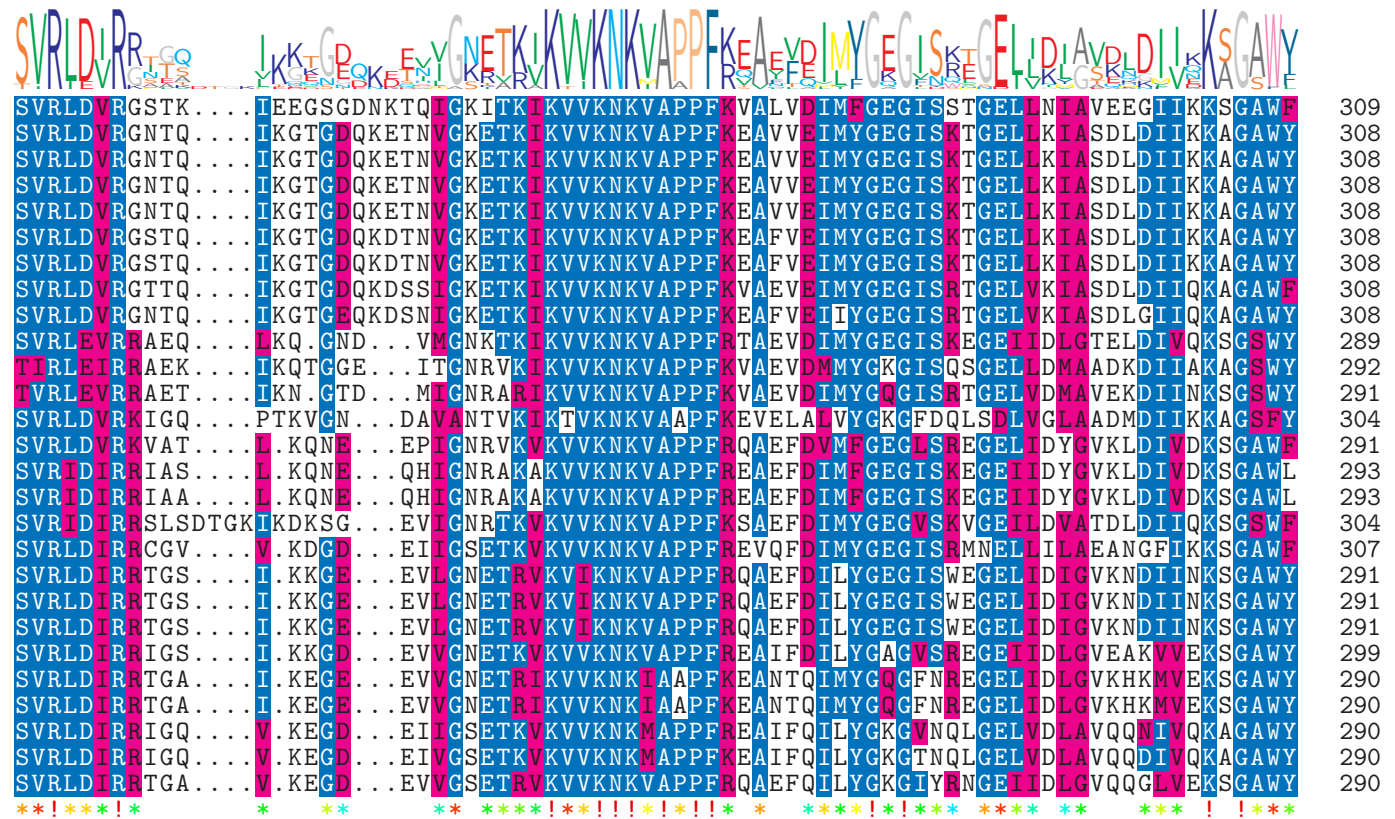
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consensus

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NTDB id 1205 PSJM300 05545 AFN77183.1	AYKGS	KIGQGR	KANAACYLEENPEI	GQIEIEQQIRAKLL	VTSAN....TKATPVAED	LADADI.....	347
consensus	*!	**!!!*****	* * *****	*****	*****	* *	* *

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved