

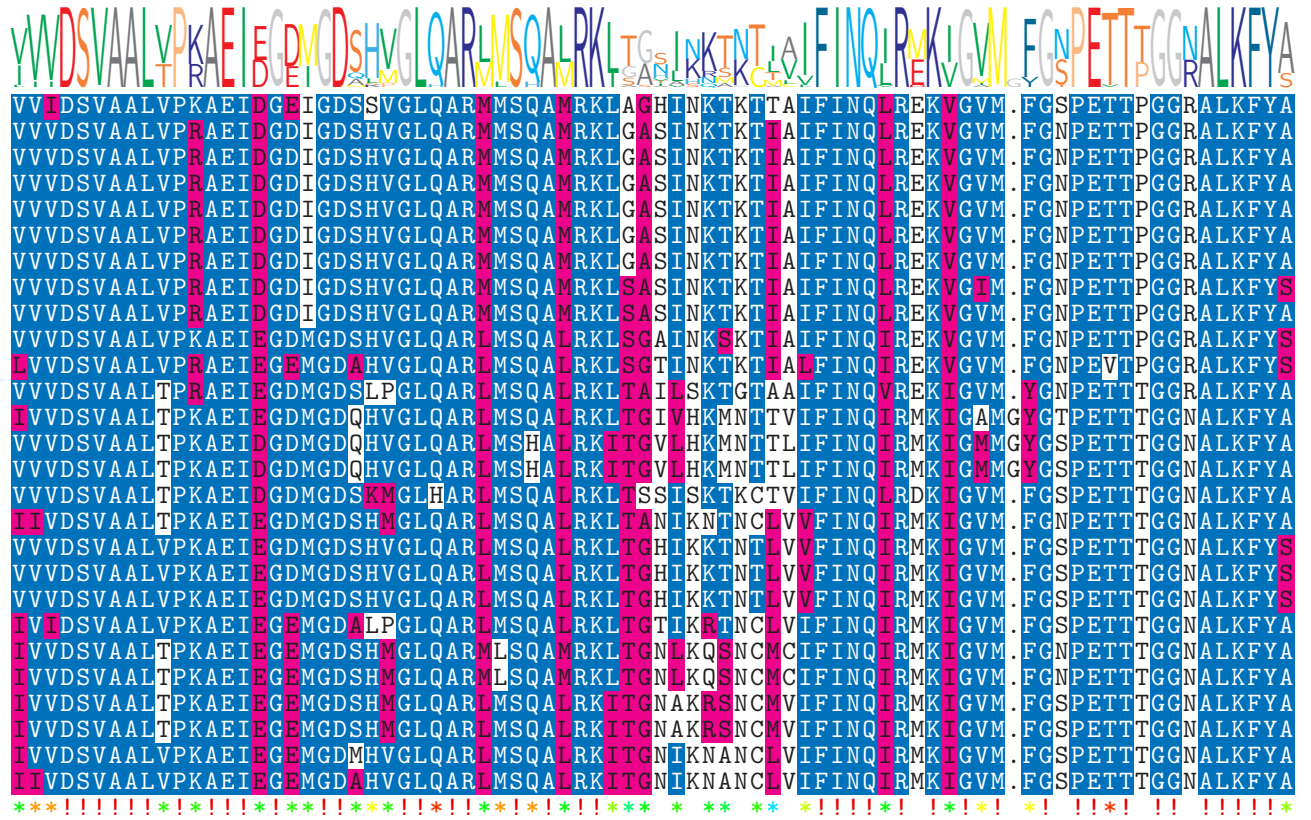
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NTDB id 600 KW2 RS01765 WP 011675413.1	MATKK...KTNFDDITKKYGAERDKALADALALIEKDFGKGSIMRLGEA.ANQKVS	VSSGSLALDI	ALGAGGYPKGR	74
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKK...PKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKV	QVMSSGSLALDI	ALGSGGYPKGR	73
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKK...PKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKV	QVMSSGSLALDI	ALGSGGYPKGR	73
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NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR...IKKTEEITKKFGDERRKALDDALKNIEKDFGKGAVMRLGER.AEQKV	QVMSSGSLALDI	ALGAGGYPKGR	73
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NTDB id 625 LCA RS02525 WP 011374196.1MAKDERQAALDAALKKIEKNFGKGSIMRMGEK.VDTQVSTVSSGSLAL	DEALGVGGYPRGR	60	
NTDB id 1313 DR RS12030 WP 010888966.1MSKDATKEISAPTDAKERSKATETAMSQIEKAFGKGSIMKLGAE.SKLD	VQVVSTGSLSLDLALGVGGIPGR	72	
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NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....K..KAASMSAEKQKALAAALAQIEKQFGKGSIMKMGDA.EVEP	VQVVSTGSLGLDVALGVGGLPRGR	68	
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NTDB id 1142 GTF74 RS11550 WP 000344154.1MDENKQKALAAALGQIEKQFGKGSIMRLGDN.RAMDVETISTGSLSLD	IALGAGGLPMGR	59	
NTDB id 1022 ACIAD RS06390 WP 004925639.1MDNKSALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDI	ALGIGGLPKGR	59	
NTDB id 1074 ABD1 RS10195 WP 000344167.1MDENKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDI	ALGIGGLPKGR	59	
NTDB id 1082868 ACLH0J RS27745 WP 003172169.1MDNKKKALAAALGQIERQFGKGAVMRMGDH.DRQAIPAISTGSLGLDI	ALGIGGLPKGR	59	
NTDB id 1205 PSJM300 05545 AFN77183.1MDENKKRALAAALGQIEKQFGKGAVMRMGDH.DRQAIPSI	ISTGSLGLDI	ALGIGGLPKGR	59
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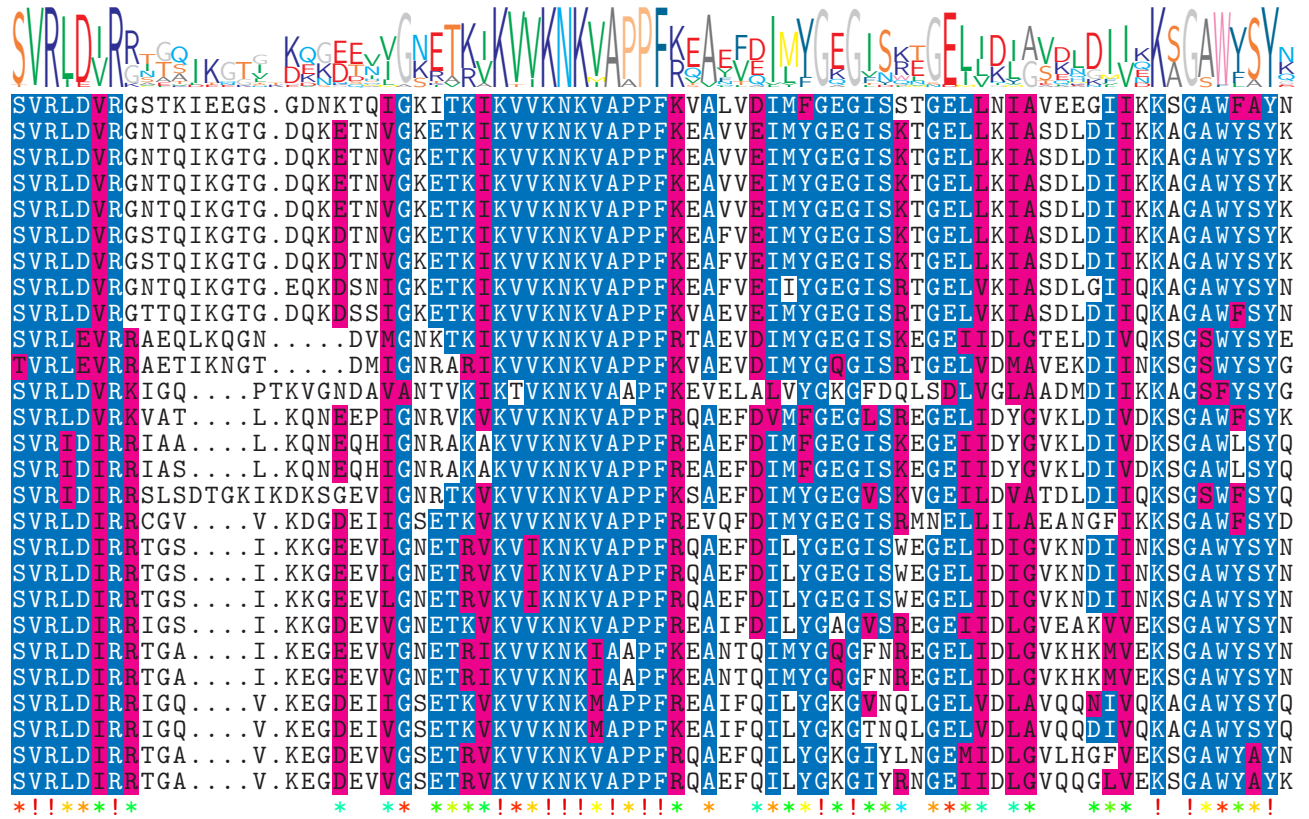
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
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NTDB id 600 KW2 RS01765 WP 011675413.1
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NTDB id 600 KW2 RS01765 WP 011675413.1	...	387
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consensus

 non conserved
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 ≥ 50% conserved