

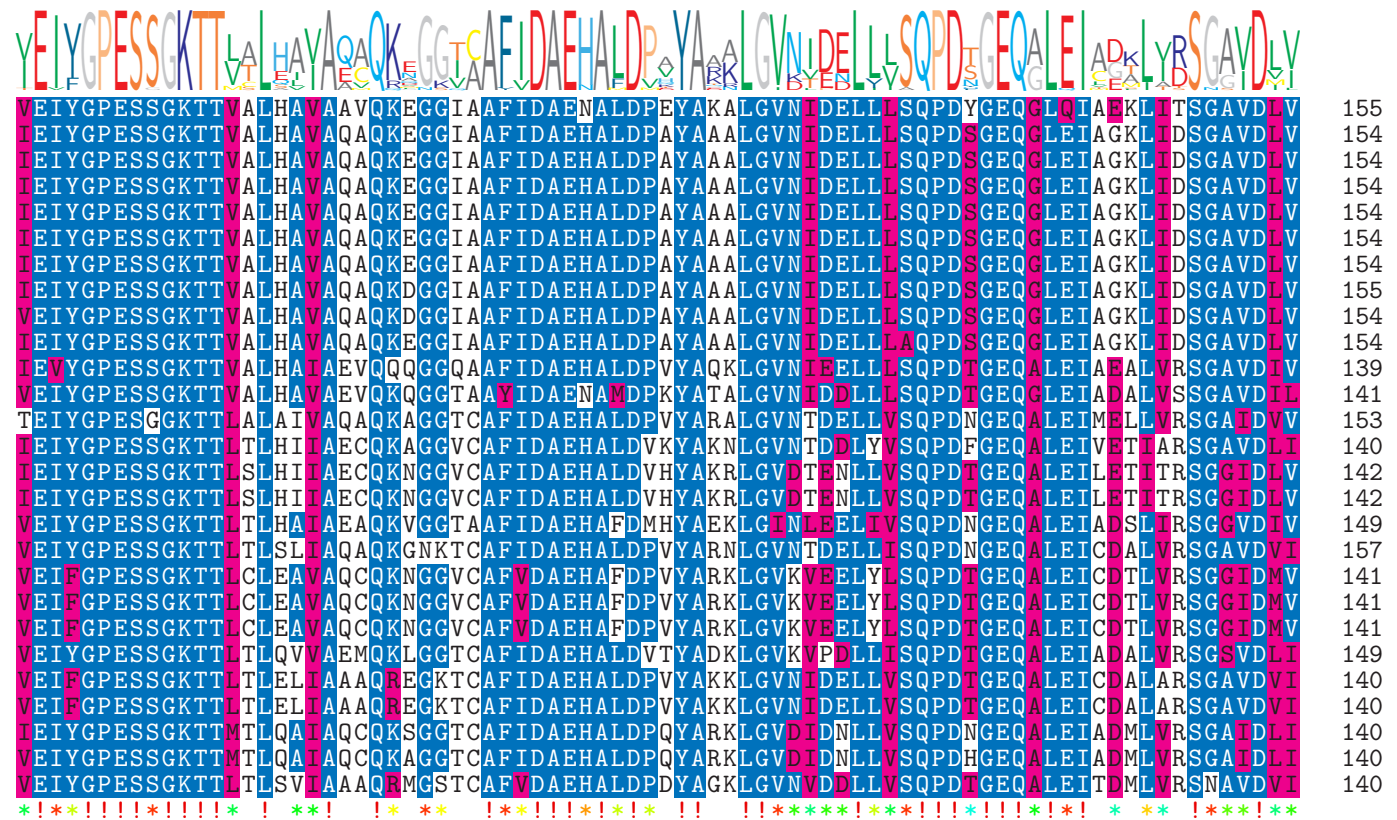
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NTDB id 194 SPR RS08825 WP 001085462.1	.MAK...KPKKLEEISKKFGAEREKALNDALKLIEKDFGKGSLMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGRI	74
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NTDB id 408 SMU RS09530 WP 002262392.1	.MAK...RIKKTEEITKKFGERKKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSSGSLALDIALGAGGYPKGRI	74
NTDB id 453 Spy49 1753c ACI62003.1	.MAK...KLKKNEEITKKFGERRRKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSSGSLALDIALGAGGYPKGRI	74
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NTDB id 625 LCA RS02525 WP 011374196.1	MAKDERQAALDAALKKIEKNFGKGSLMRLGEK.VDTQVSTVSSGSLALDEALGVGGYPKGRI	61
NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTDAKERSKATETAMSQIEKAFGKGSLMRLGAE.SKLDVQVVSTGSLSLDLALGVGGIPKGRI	73
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDENKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGGLPKGRI	60
NTDB id 1022 ACIAD RS06390 WP 004925639.1	MDDNKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGGLPKGRI	60
NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQFGKGAVMRMGDHD.RQAIPSIISTGSLGLDIALGIGGLPKGRI	60
consensus	*** ** *	

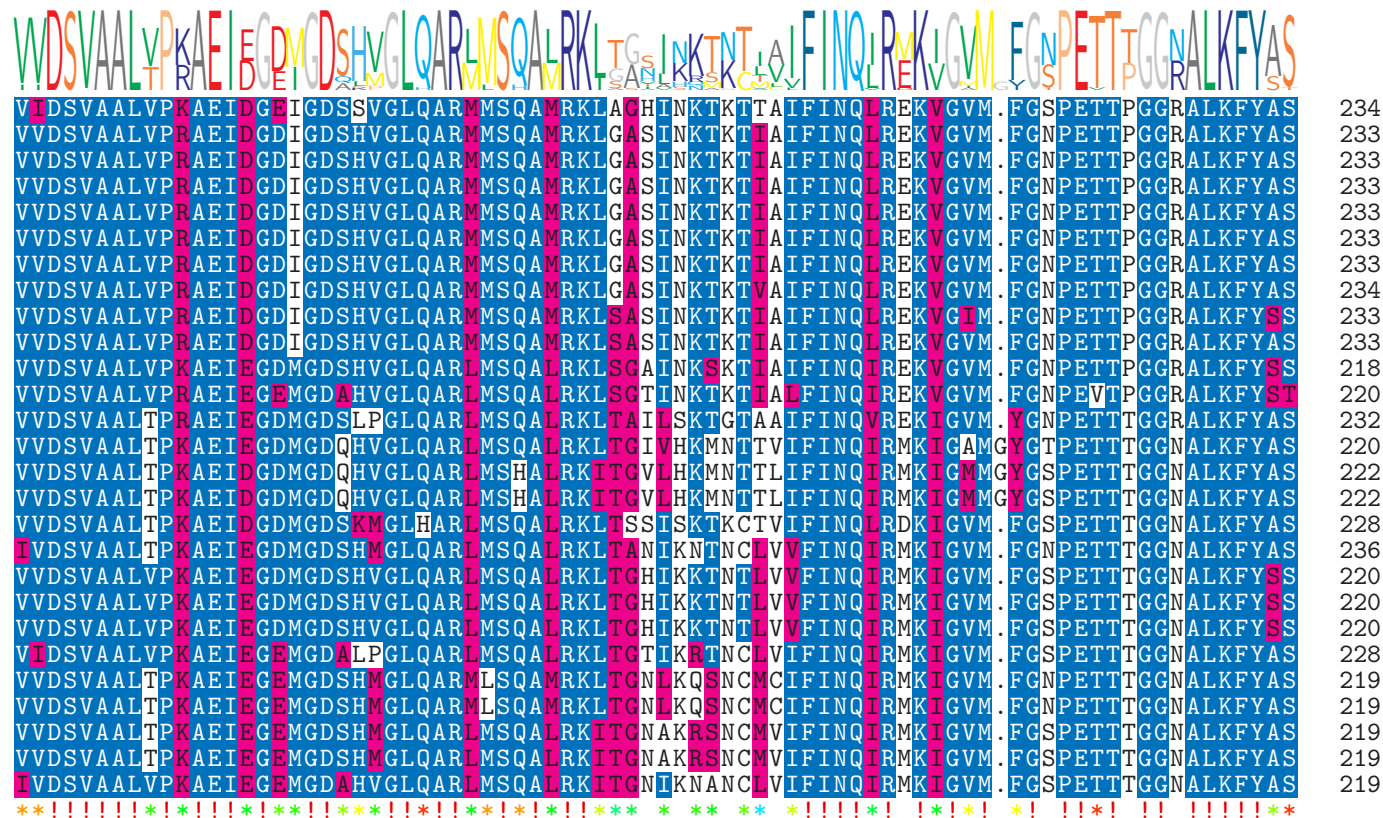
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consensus



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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved