

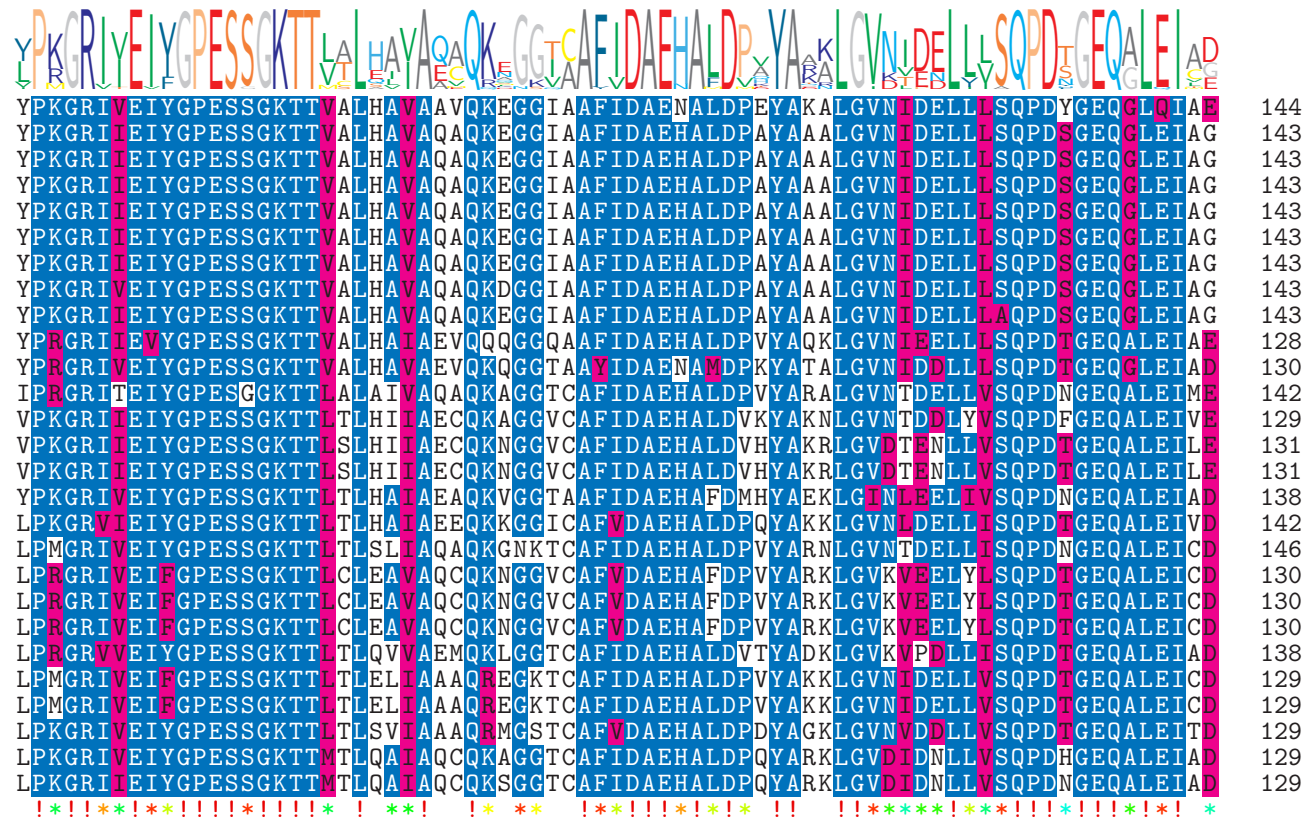
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NTDB id 453 Spy49 1753c ACI62003.1	MAKKLKKNEEITKKFGDERKKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSSGSLALDIALGAGG	68
NTDB id 115 BSU 16940 NP 389576.2	MSDRQAALDMALKQIEKQFGKGSIMKLGEK.TDTRISTVPSGSLALDTALGIGG	53
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NTDB id 1313 DR RS12030 WP 010888966.1	MSKD.ATKEISAPTDAKERSKATETAMSQIEKAFGKGSIMKLGAE.SKLDVQVVSTGSLSLDLALGVGG	67
NTDB id 1245 Cj1673c YP 002345041.1	MDDNKRKSLDAALKSLDKTFGKGTLRLGDK.EVEQIDSIGTGSVGLDLALGIGG	54
NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDEDKKKATSLATKQIDKVFKGKALVRLGDK.QVEKIDAISTGSLGLDLALGIGG	56
NTDB id 1211 C694 RS00795 WP 000963128.1	MAIDEDKKKATSLATKQIDKVFKGKALVRLGDK.QVEKIDSISTGSLGLDLALGIGG	56
NTDB id 1334 RAOC RS04870 WP 004918236.1	M.AKTET..TSPNDKKKKALAAVLEKLDKTYGKGTVMKLGDTVDHNIIEVIPSGLGLDLALGVGG	63
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NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQFGKGAVMRMGDH.DRQAIPSIISTGSLGLDIALGIGG	54
NTDB id 1022 ACIAD RS06390 WP 004925639.1	MDDNKKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGG	54
NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDENKKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGG	54
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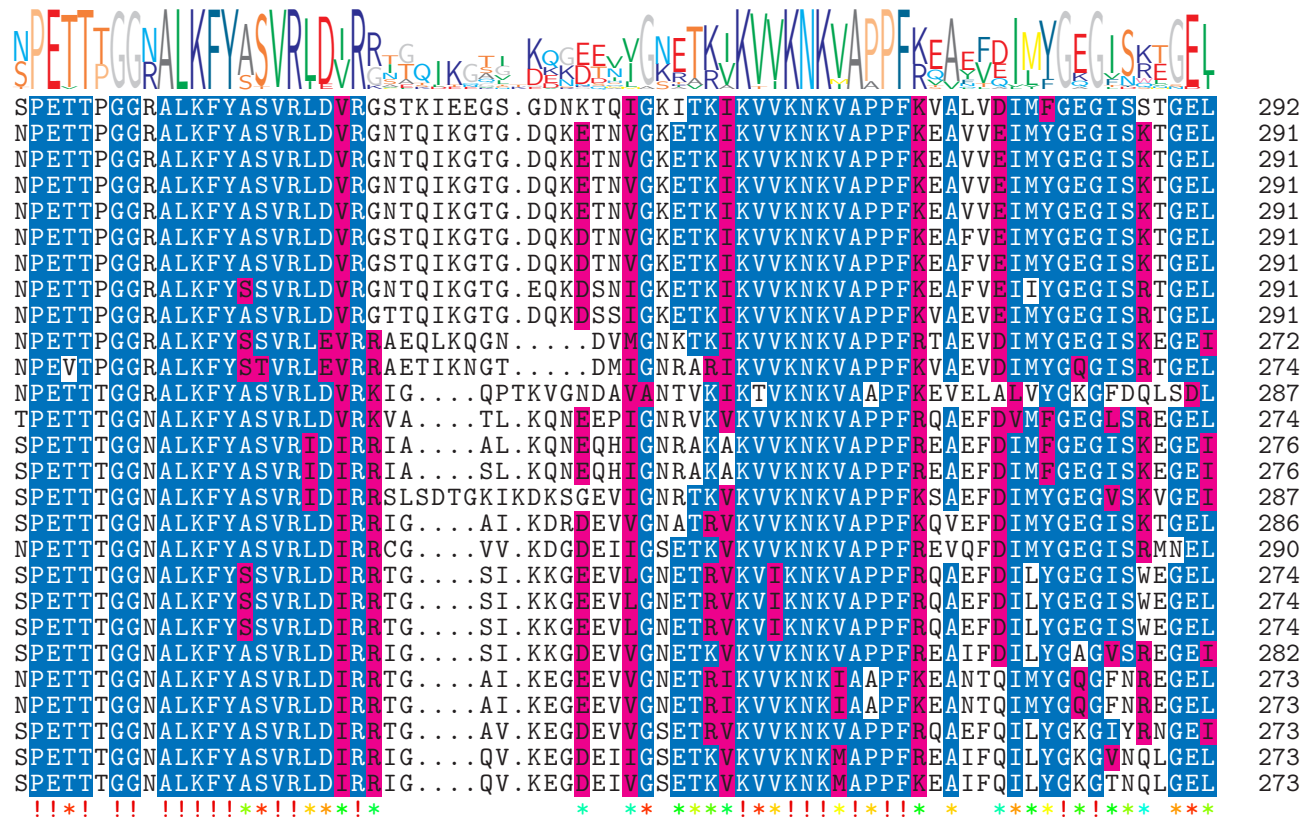
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consensus

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NTDB id 499 SM12261 RS08315 WP 078228268.1	LKIASDLDITIKKAGAWYSYKDEKIGQGS	ENAKKYLA	DNPEIF	FDEIDHQVRVQFGL	IDGEEIAES...KKDE...	359	
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consensus	* **	*** ! !****!	***!!*****	* **	*****	***	*

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consensus		*	

- ⓧ non conserved
- ✖ similar
- ⓧ ≥ 50% conserved