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NTDB id 1307	DR RS10055 WP 010888596.1	MTSGIDITDITRLAADKGSADTILLAGLPQFKISGDIYAQGFEPISASDTRKLMYSMMNERQQRIFEFEKRELDDFSFALG	80
NTDB id 1185	GTF74 RS11940 WP 000350195.1	MDIAELLEFSVKHNASDLHLSAGVPPMVRIDGEVRKLGVPFATHSDVHRLIFEIMNDAQRSEYEKLEVDDFSFLP	76
NTDB id 1176	A1552VC RS01160 WP 000350195.1	MDIAELLEFSVKHNASDLHLSAGVPPMVRIDGEVRKLGVPFATHSDVHRLIFEIMNDAQRSEYEKLEVDDFSFLP	76
NTDB id 1081	FDQ49 RS05335 WP 006581387.1	MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALAHKDVHRLVYDIMNDKQRQDYEEKLETDFSFEVP	76
NTDB id 1060	ABD1 RS04380 WP 000355489.1	MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALAHKDVHRLVYDIMNDKQRQDYEEKLETDFSFEVP	76
NTDB id 1040	HON27 RS13360 WP 000355489.1	MDITELLAFSVKNGASDLHLSAGMPPMIRVDGEVRRINLPALAHKDVHRLVYDIMNDKQRQDYEEKLETDFSFEVP	76
NTDB id 1202	PSJM300 01600 AFN76400.1	MDITELLAFSAKQGASDLHLSGLPPMIRVDGVRRINLPAMDHKQVHALIYDIMNDKQRKDFFEFLETDFSFEVP	76
NTDB id 1194	PAKAF RS01995 WP 003084552.1	MDITELLAFSAKQGASDLHLSAGLPPMIRVDGVRRINLPLEHKQVHALIYDIMNDKQRKDFFEFLETDFSFEVP	76
NTDB id 210309	Ga0123462 RS08445 WP 100265893.1	MTENISIRQLLQVMVDKASDLYLMAGTTPPGYRIRNGVIHRLGDKEFNPIITVERLAGELMSEKQKADFASTMEMNLSSAFP	80
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NTDB id 1203	PSJM300 01605 AFN76401.1	MEFEKLLRLMVEKGCSDLFITAGVPPSMKVNGKILPVNKTTPMSPEMTRETIVHGMVNEQQRREFTENHECNFAISAR	76
NTDB id 1019	ACIAD RS04205 WP 004922051.1	MDFNDLLNLMAKHKASDLFITAGVEPSLKINGETTPVSKTKLDGDMIAQLIDSITTTQKQRQEFITHECNFAILNR	76
consensus		*** *!* **** *!*!*****!*****!***** *! * * *************** *! *****	

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NTDB id 1176	A1552VC RS01160 WP 000350195.1	N.VGRFRVNAFHQARGCSAVFRTIPTVTIPTLEQLDAPEIFSKIANYEKGLVLVTGPTGSGKSTTLAAMVNYVNAHNNKHI	155
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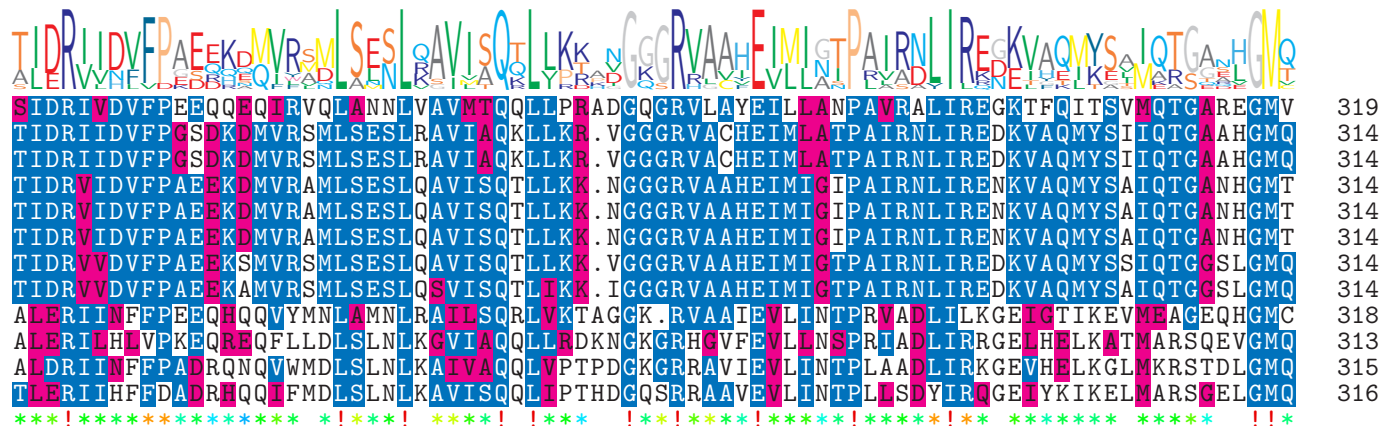
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consensus		***!!!!*!*! *****!!!!*!*! *! !!*****!*****!*****!*****!*****!*****	

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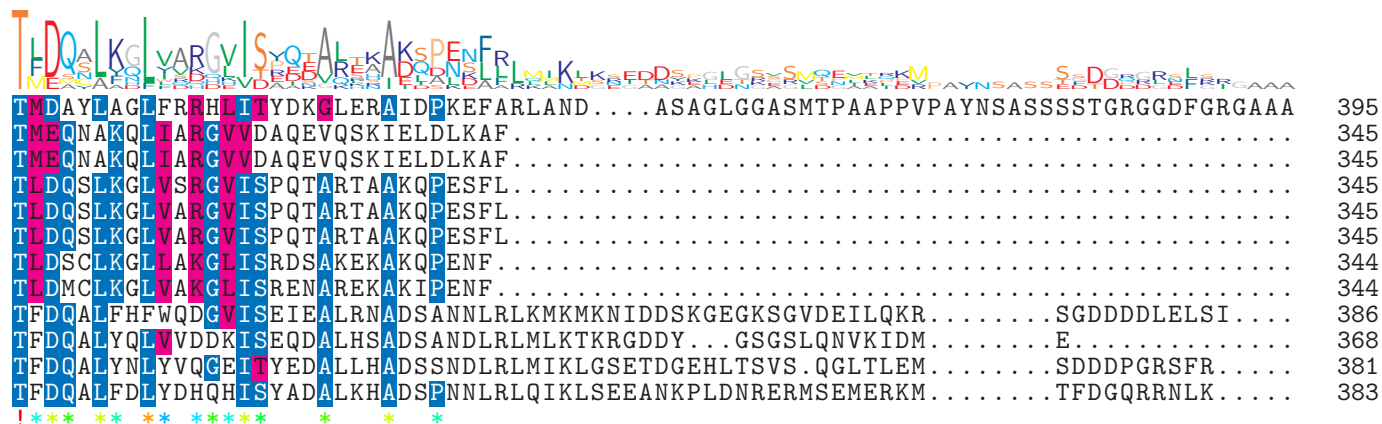
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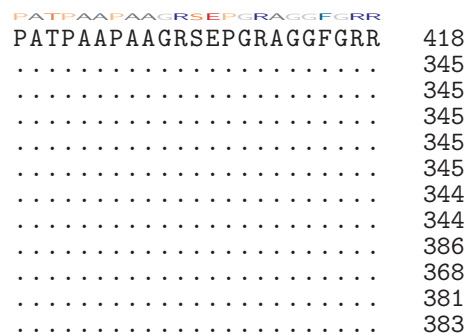
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NTDB id 1203 PSJM300 01605 AFN76401.1
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consensus

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- X non conserved
- X similar
- X >50% conserved