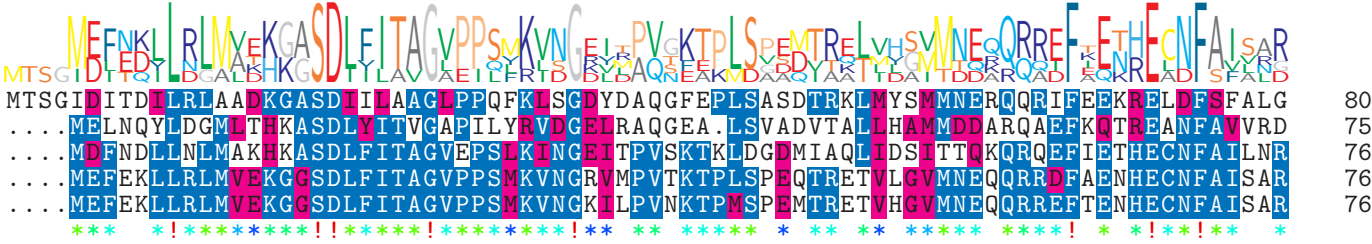


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NTDB id 1203 PSJM300 01605 AFN76401.1
consensus



logo

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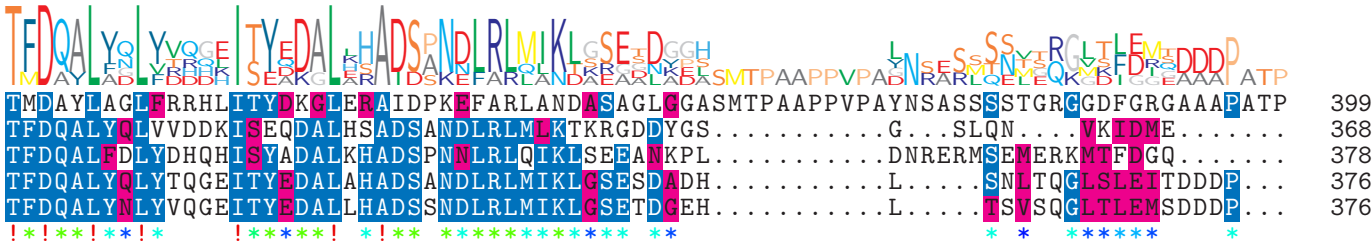
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consensus



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NTDB id 1019 ACIAD RS04205 WP 004922051.1	R R N.....L K ..	383
NTDB id 225372 B7W86 RS02130 WP 003110016.1	... A ..G R R..... F R ..	382
NTDB id 1203 PSJM300 01605 AFN76401.1	... G .. R S..... F R ..	381
consensus	* * **	

Sequence logos for the 12 amino acids of the 12mer peptide. The logos show the relative frequency of each amino acid at each position. The sequences are: A A P G A R S R E P G R A G G L F R R, A A P A A G R S E P G R A G G F G R R, R R R, L K, F R, F R. Asterisks are placed below the first, second, and last two positions.

- X non conserved
- X similar
- X $\geq 50\%$ conserved