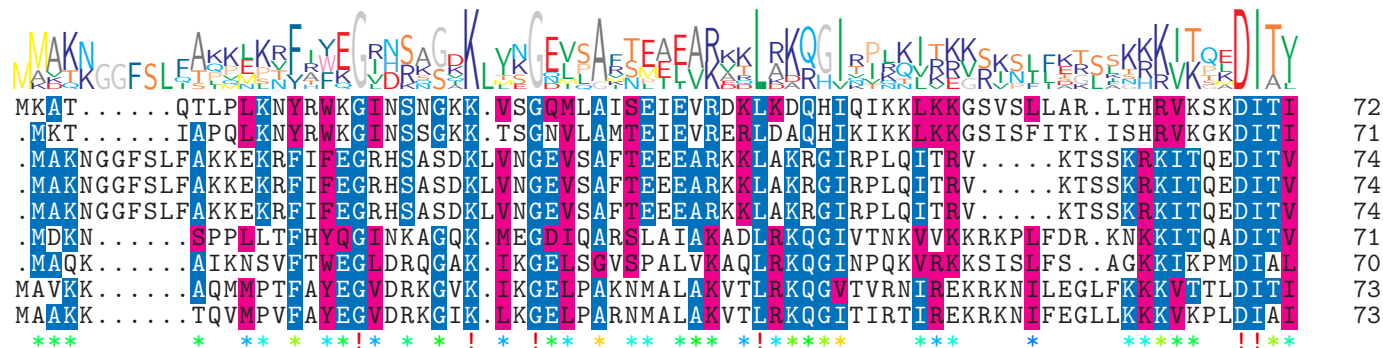


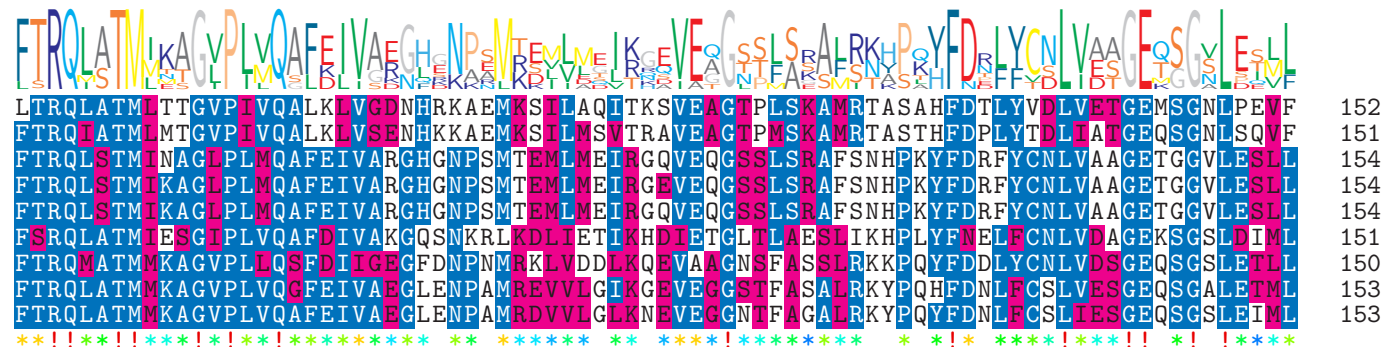
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consensus



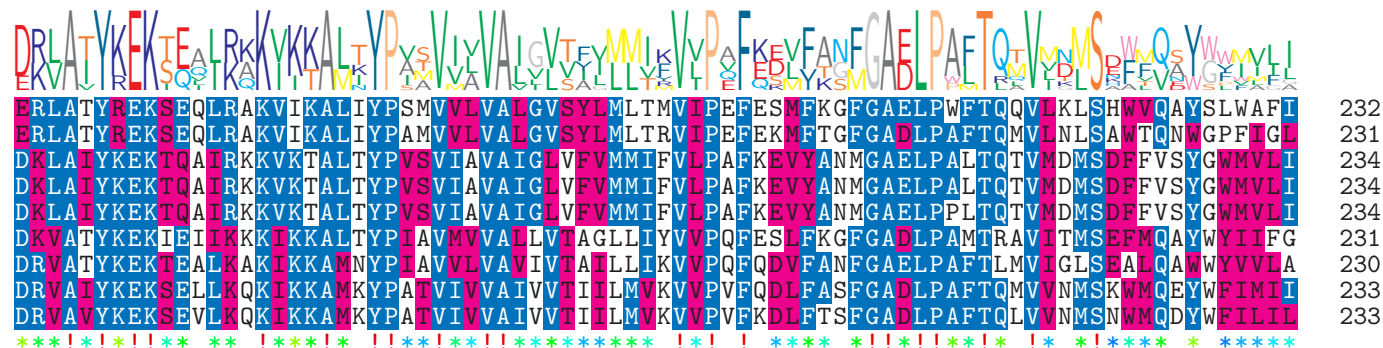
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consensus

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Sequence	Position 1	Position 2	Position 3	Position 4	Position 5	Position 6	Position 7	Position 8	Position 9	Position 10	Score
VVAMYLPIFNLMSVLG.	V	A	M	L	P	I	F	N	L	M	408
VTAMYLPIFNLMSVLG.	V	T	A	M	L	P	I	F	N	L	407
LVAMYLPIFNLGNVVA*	L	V	A	M	L	P	I	F	N	L	410
LVAMYLPIFNLGNVVA.	L	V	A	M	L	P	I	F	N	L	410
LVAMYLPIFNLGNVVA.	L	V	A	M	L	P	I	F	N	L	410
VVAMYLPIFKLGSAV..	V	V	A	M	L	P	I	F	K	L	406
ITAMYLPIFQMGSVV..	I	T	A	M	L	P	I	F	Q	M	405
VIAMYLPIFQMGSVV..	V	I	A	M	L	P	I	F	Q	M	408
VIAMYLPIFQMGSVV..	V	I	A	M	L	P	I	F	Q	M	408

* * * * * ! * * * * * * * * * *

-  non conserved
-  similar
-  > 50% conserved