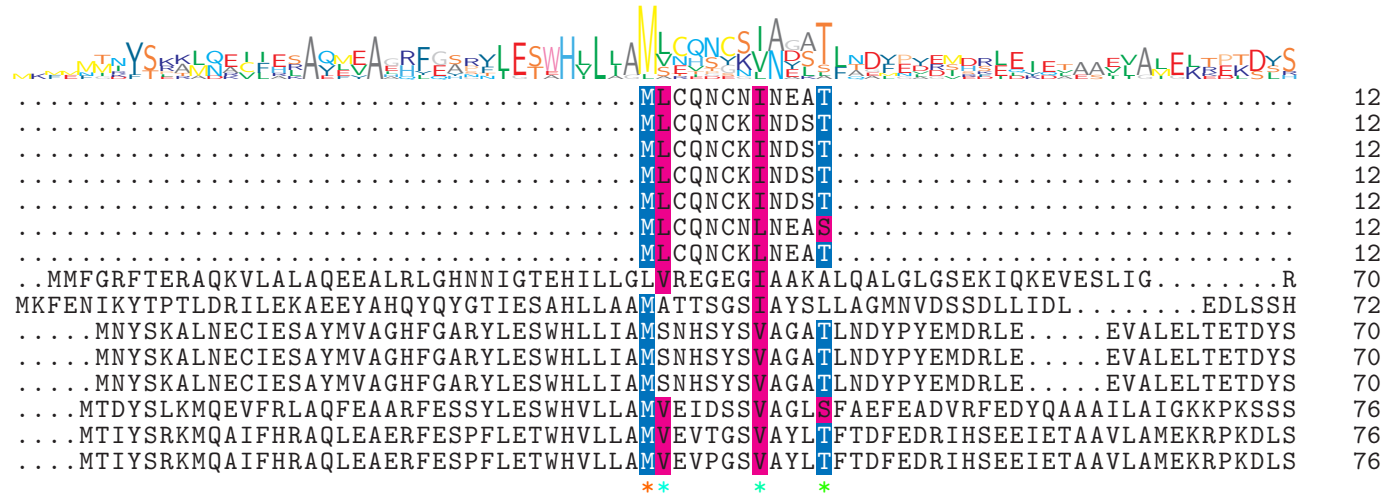


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consensus

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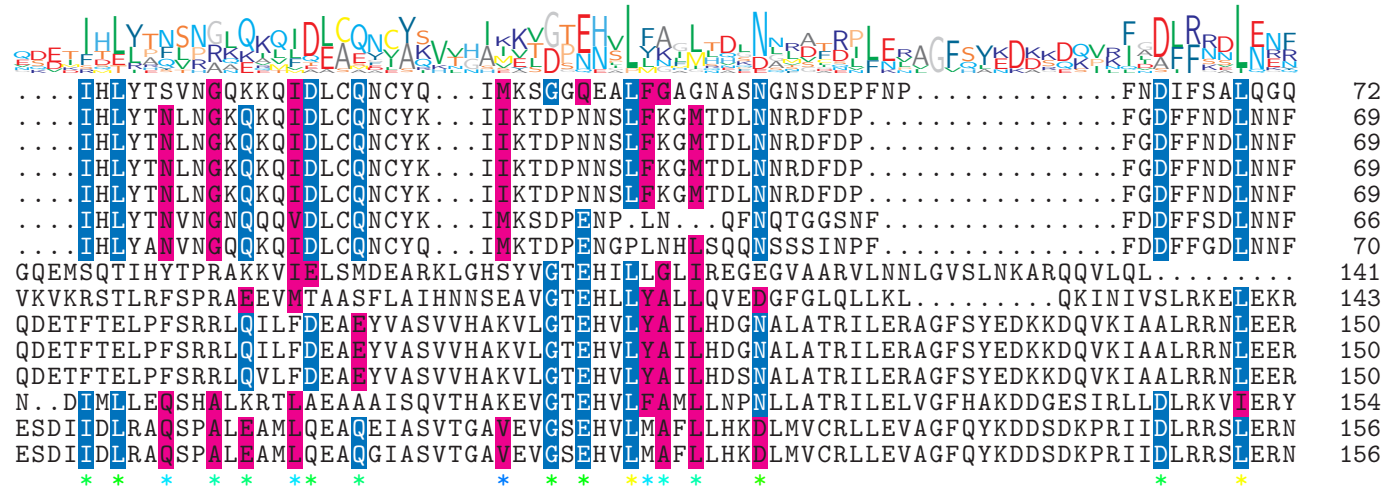


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consensus

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consensus

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RPSSN..TP....PIPTQTSGGGYGGNGGY.GSQNRGSAQTTPPSQEKGLLEEFGINVTEIARRGIDPVIGRDEEII	141
RSS.NGDLP....NTPPTQEGCNRGNGGNTQGPRPGGPRQAPQPPQGLLEEFGINLTDIARRGIDPVIGRDEEII	140
RAFNNQDLP....NTPPTQAGCGNGNGGNNR...RPGGPQQQASQKPNGLLEEFGINLTDIARRGIDPVIGRDEEII	142
LGSNET.....GSSAAGTNSMANPTLDSLARDLTALAKEDSLDPVIGRSKEIQR	191
TGLKVPESKKAVTPM SKRMAK.....VAENSTTPTLDSVSSDLTEBARLCKLDPMIGRAEIDR	204
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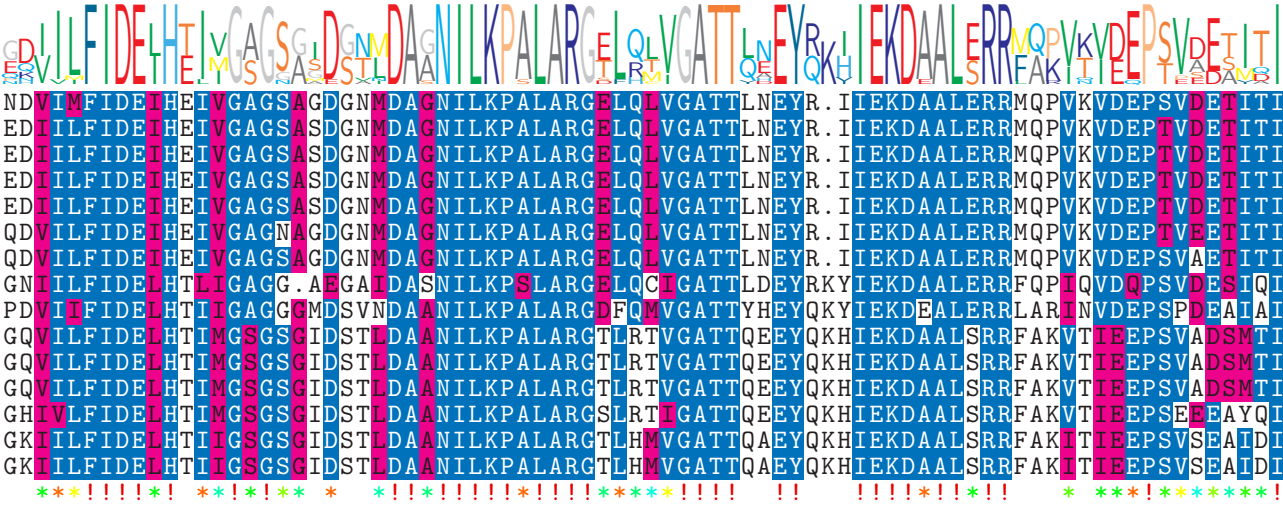
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consensus

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consensus



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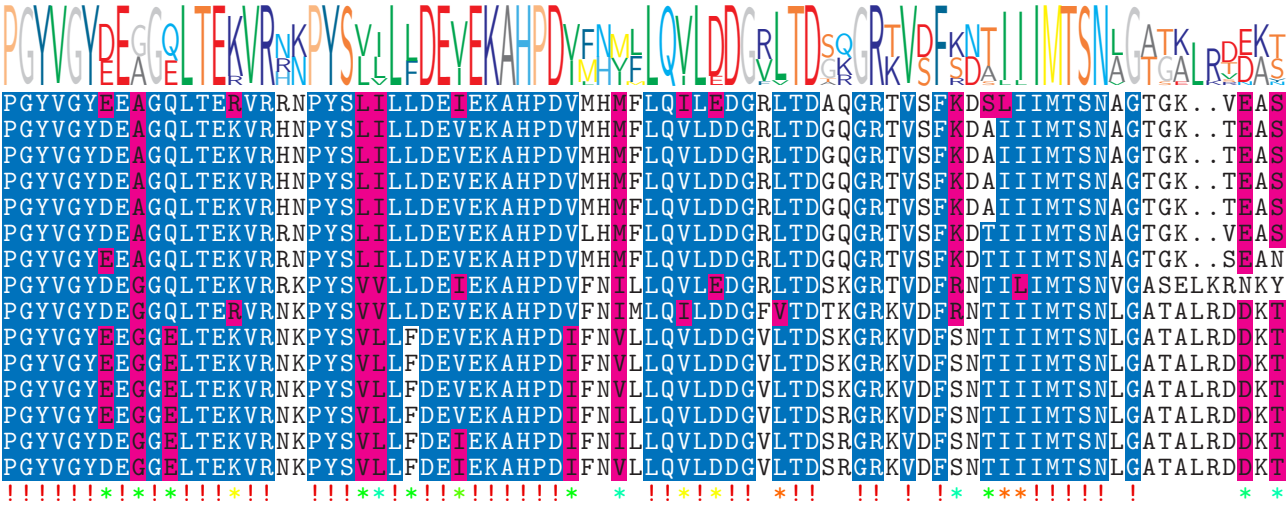
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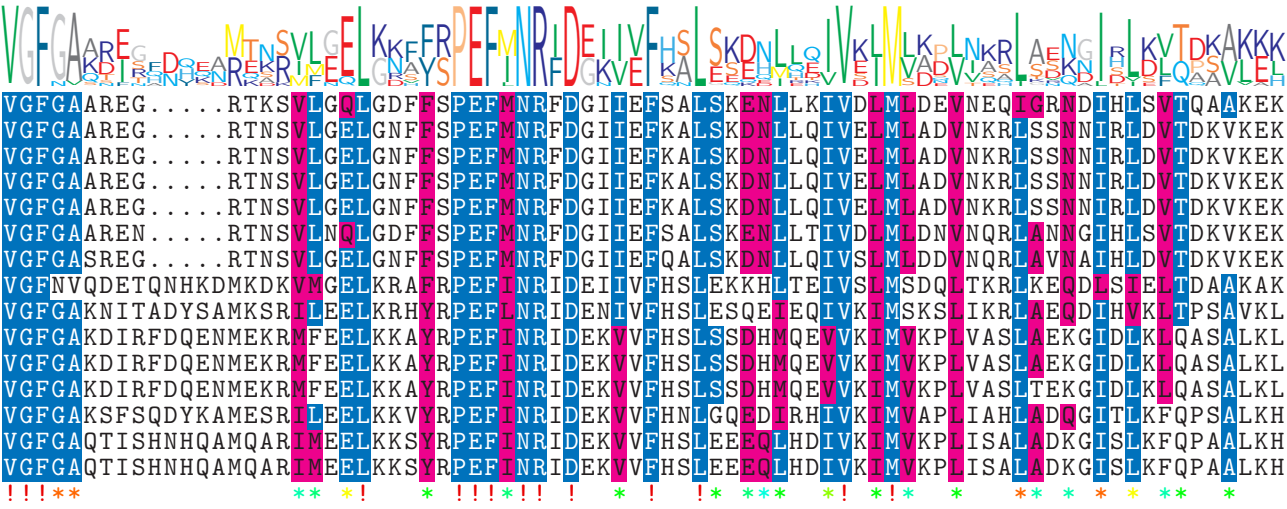
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consensus



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consensus



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consensus	* * !** * * !!!!!!!!!** * * * ! *	

- non conserved
- similar
- ≥ 50% conserved