

logo



Sequence	Position 10
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MLCQNCKINDST	12
MLCQNCKINDST	12
MLCQNCNINEAT	12
MLCQNCNINEAT	12
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....MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
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...MTIYSRKMQAIFHRAQLEAERFESPFLETWHVLLAMVEVTGSAVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDL	76
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consensus

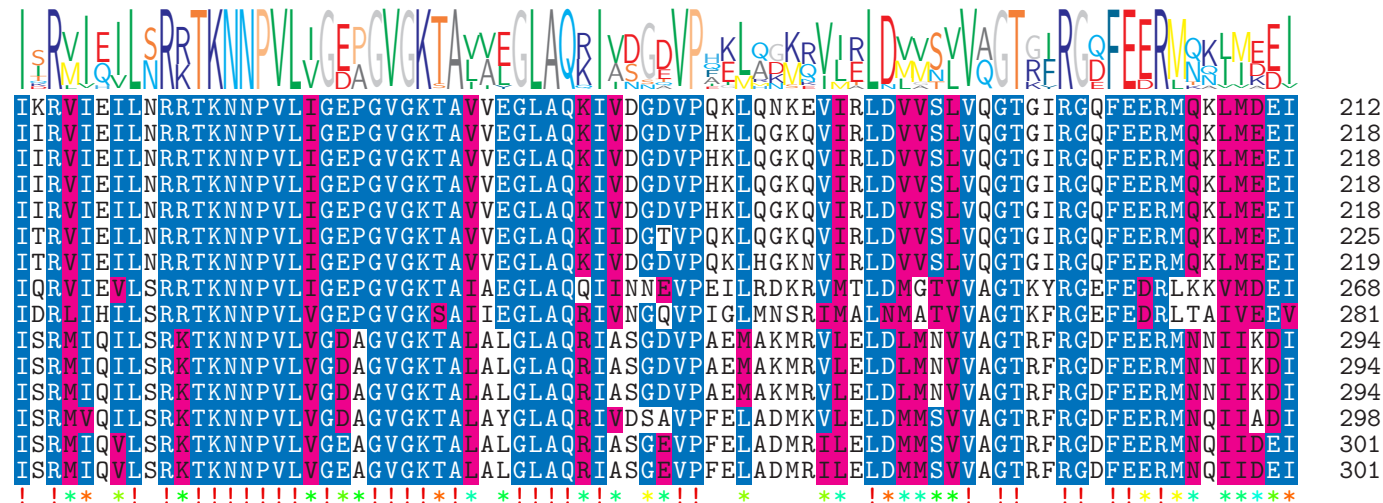
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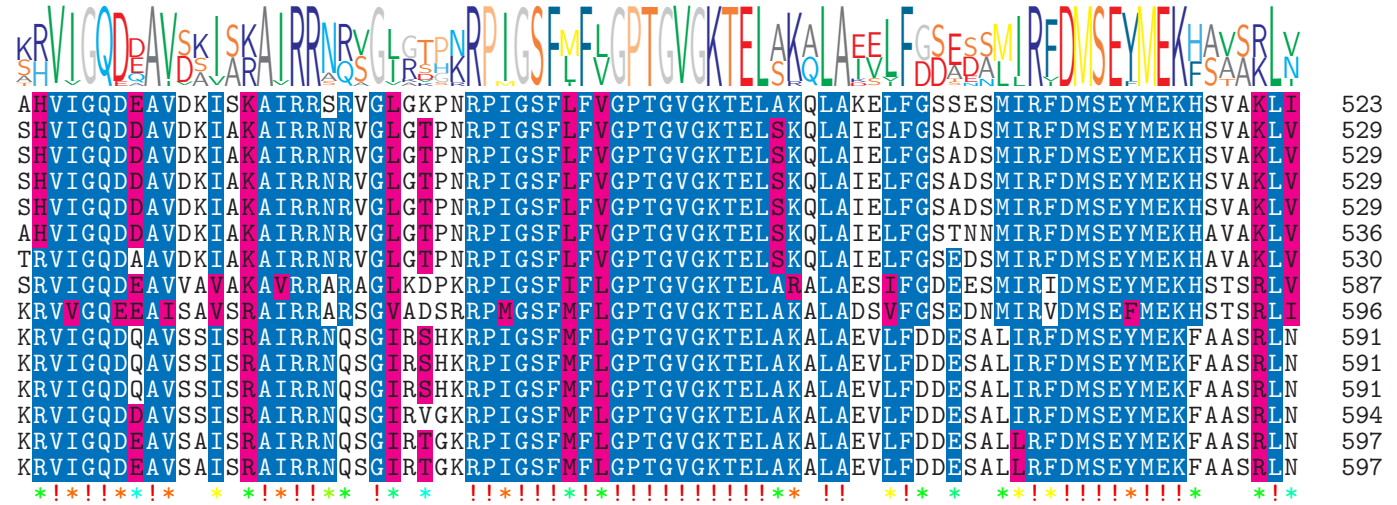
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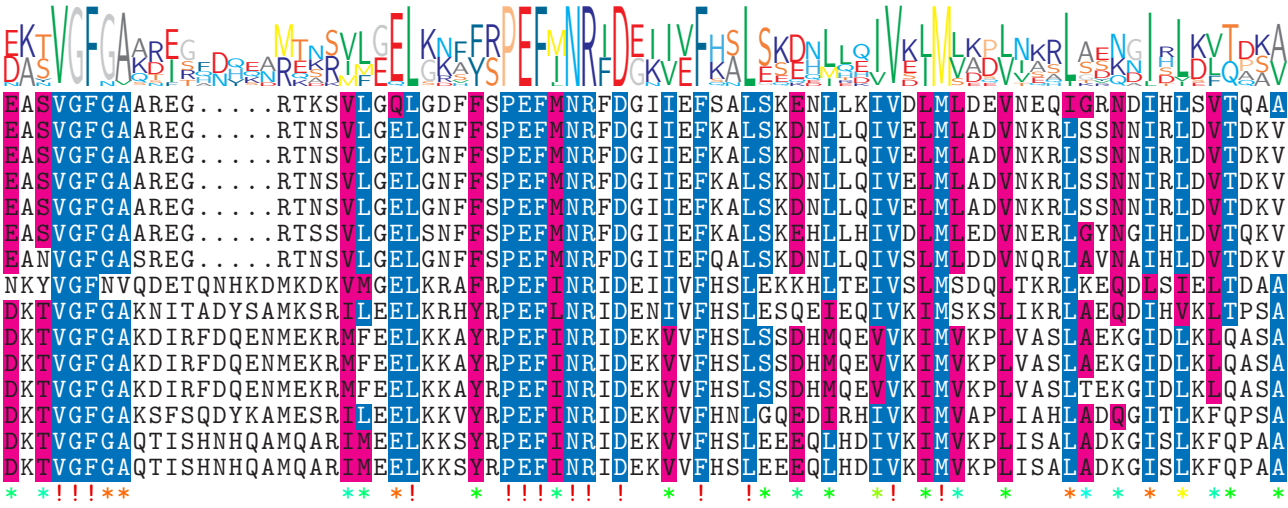
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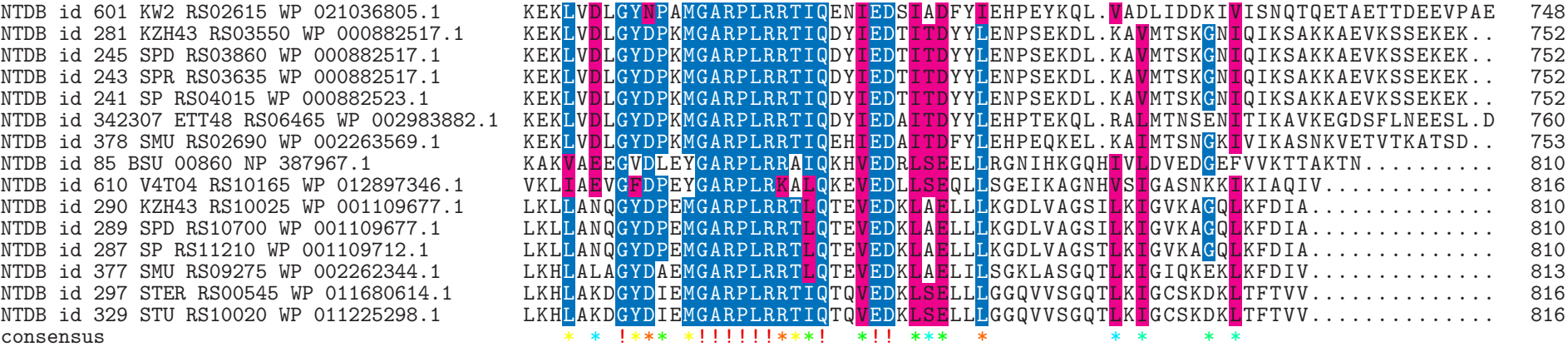
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consensus



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logo



X non conserved
X similar
X ≥ 50% conserved