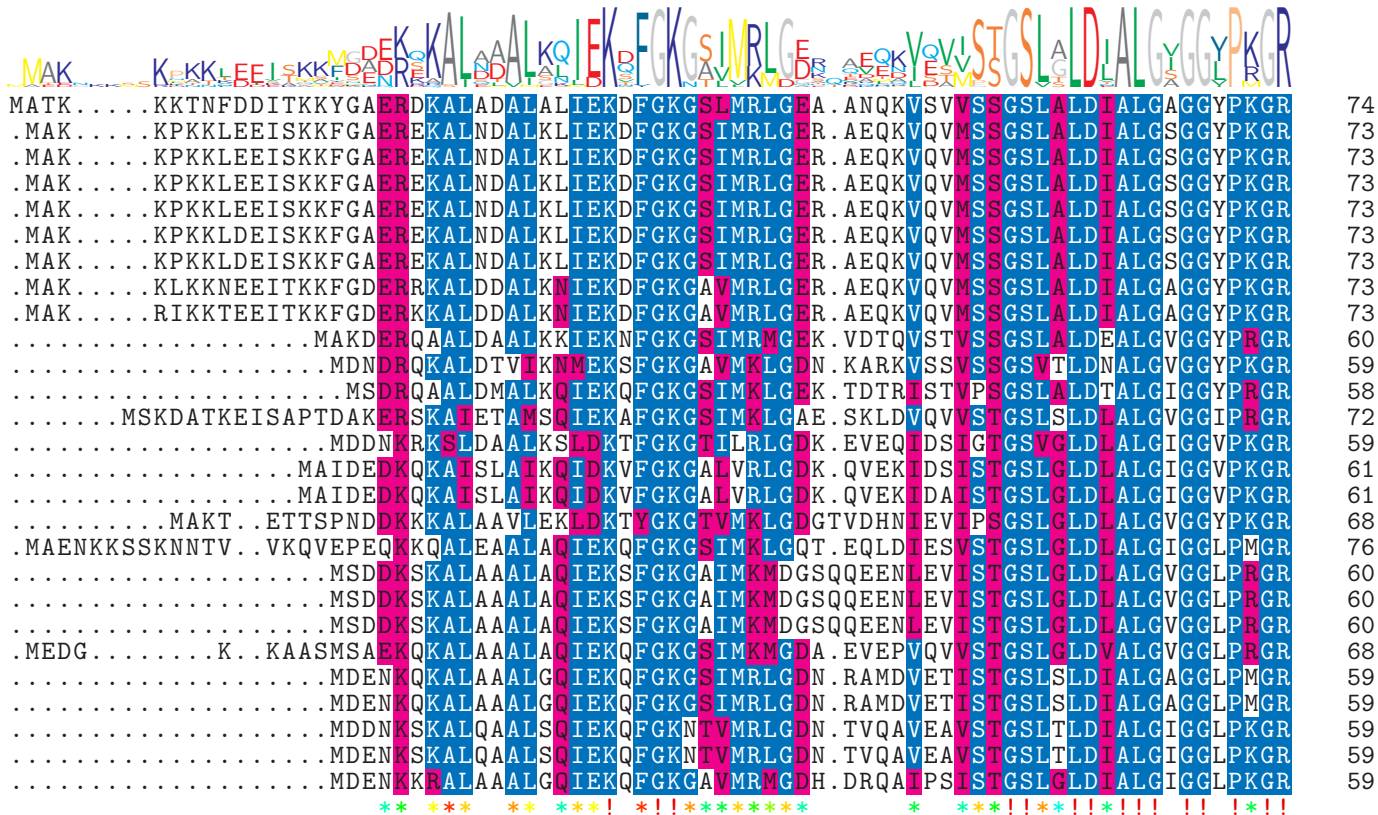


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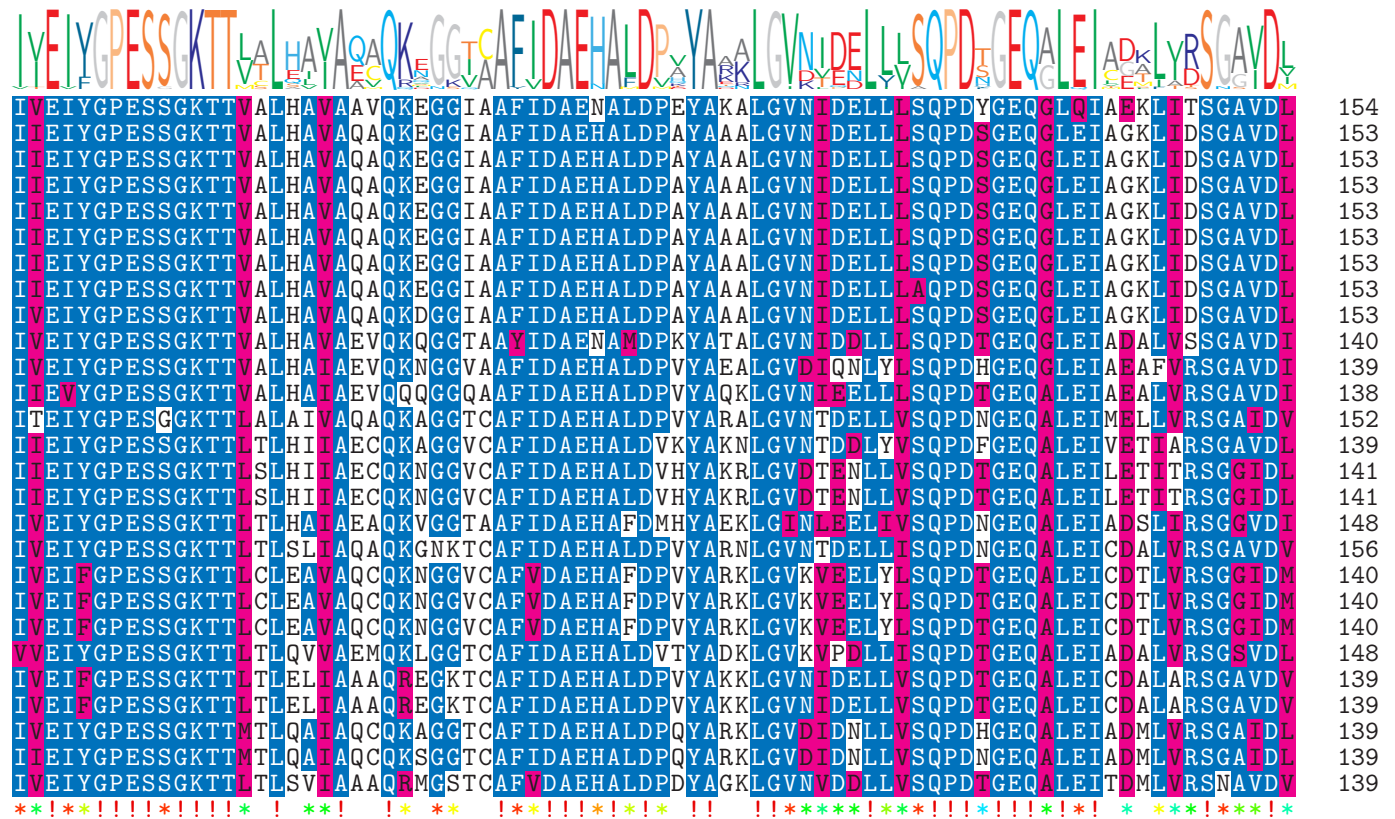
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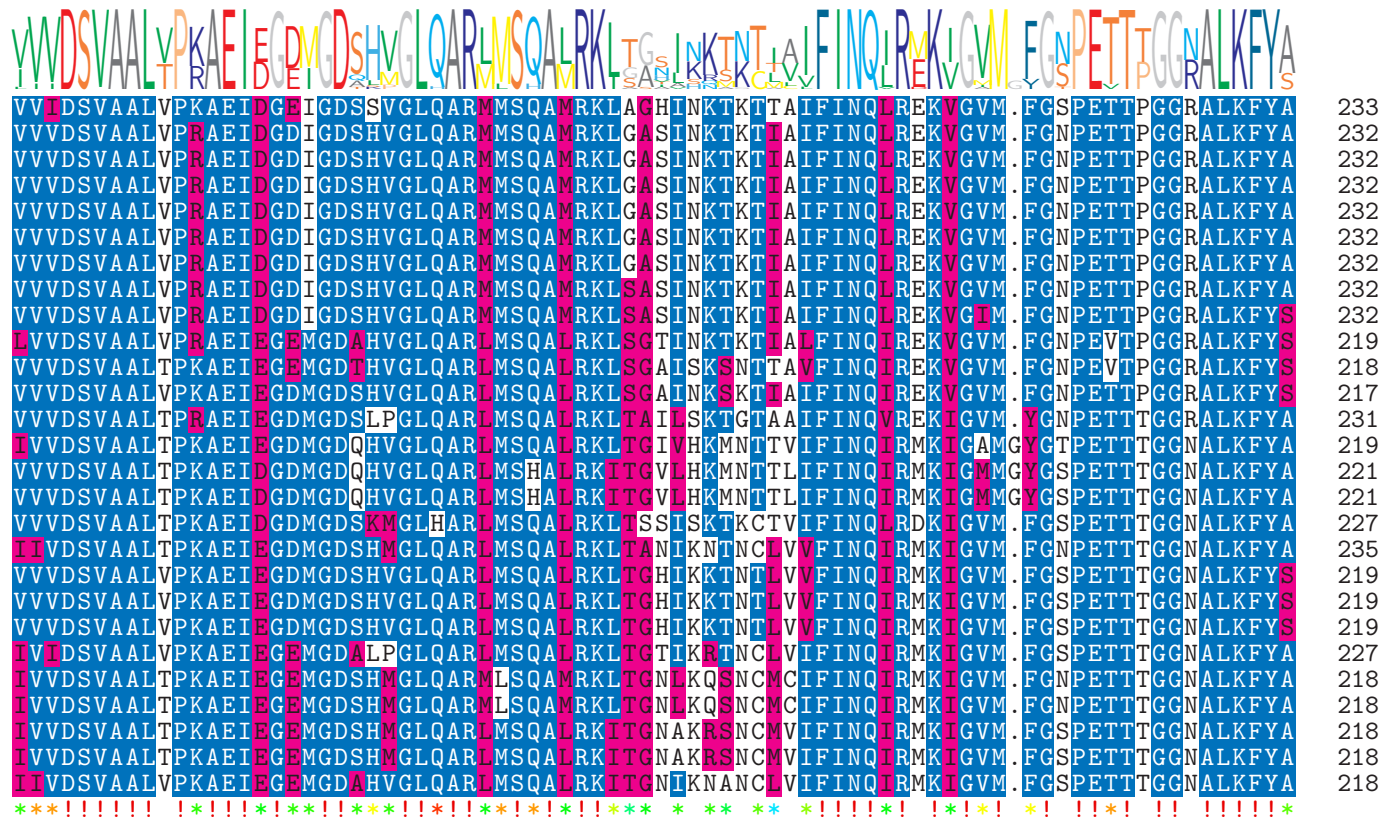
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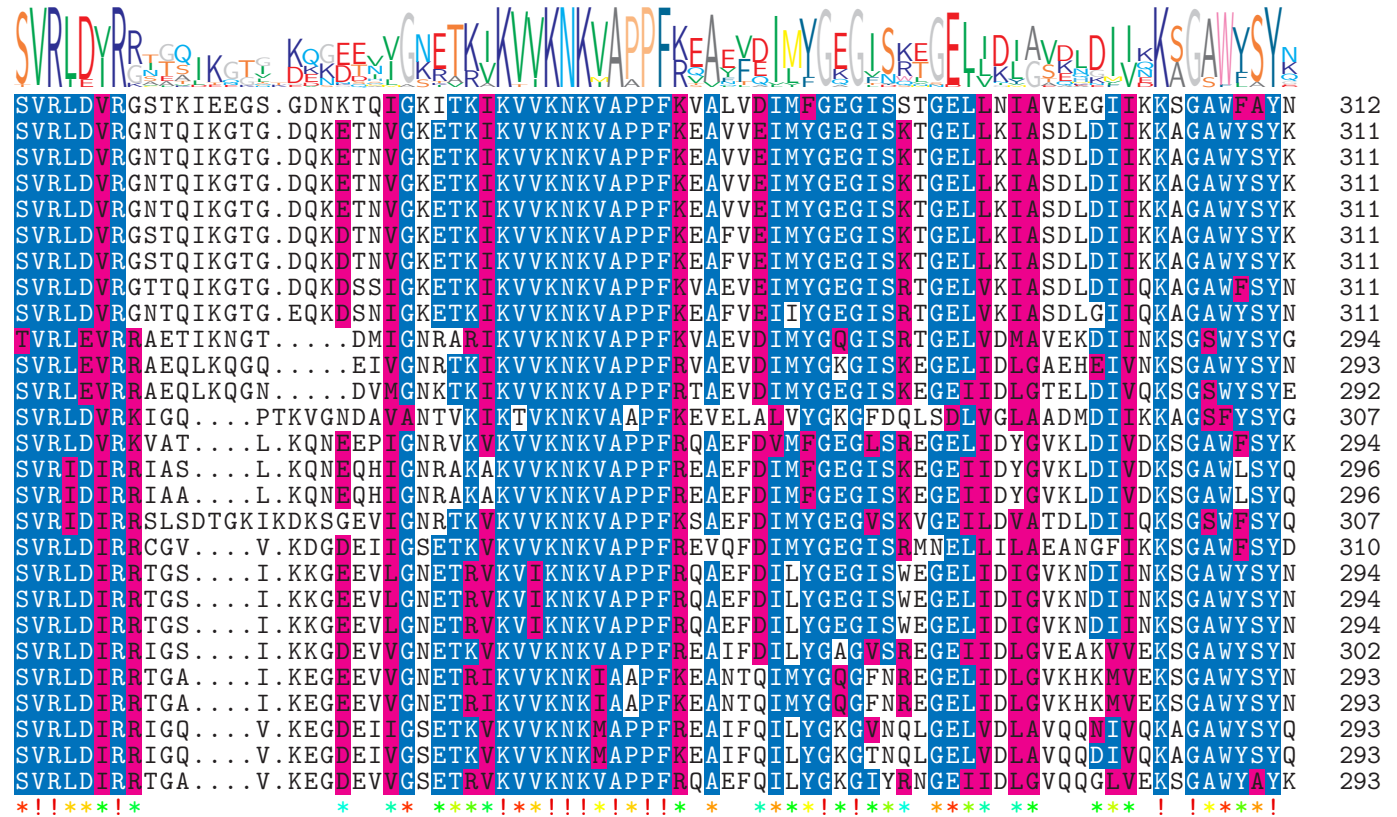
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GN KIGQGGKNNVIRYFEENNRIABEIERNIREQLLTGTN GAVQ . I . ED EE . EPDLL . LES 349

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consensus			

- non conserved
- similar
- ≥ 50% conserved