

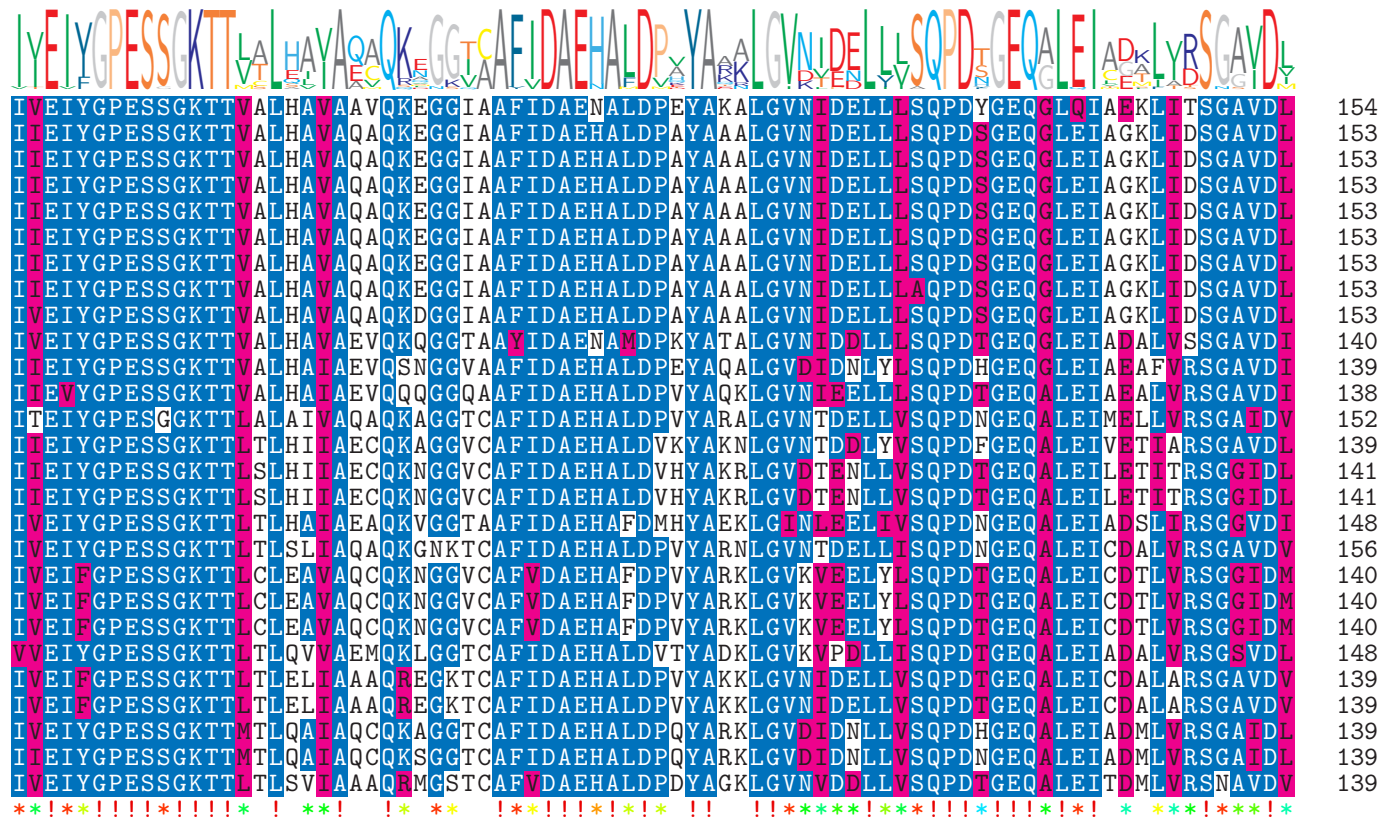
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consensus		*** **	***	***!	***!	***!	***!	***!	***!		

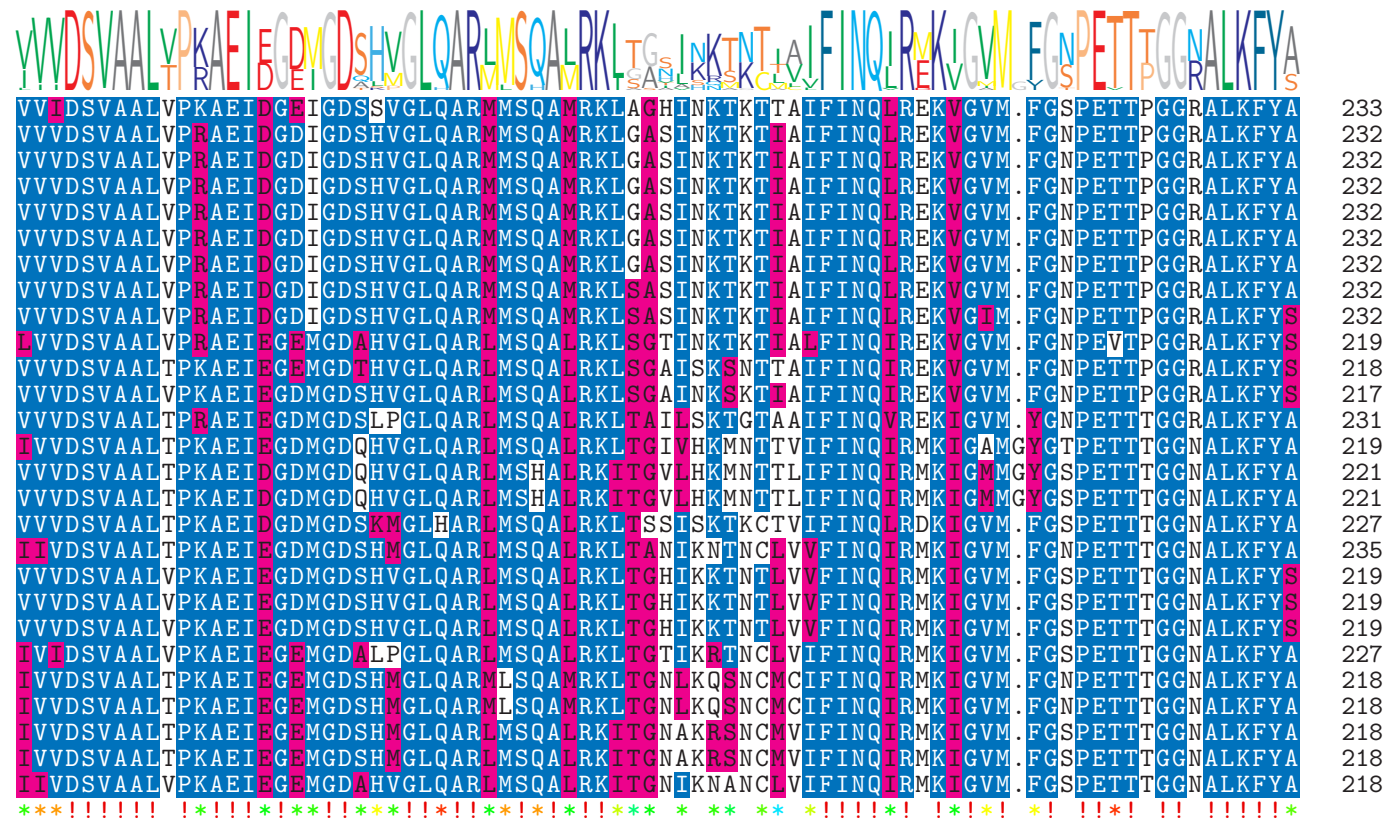
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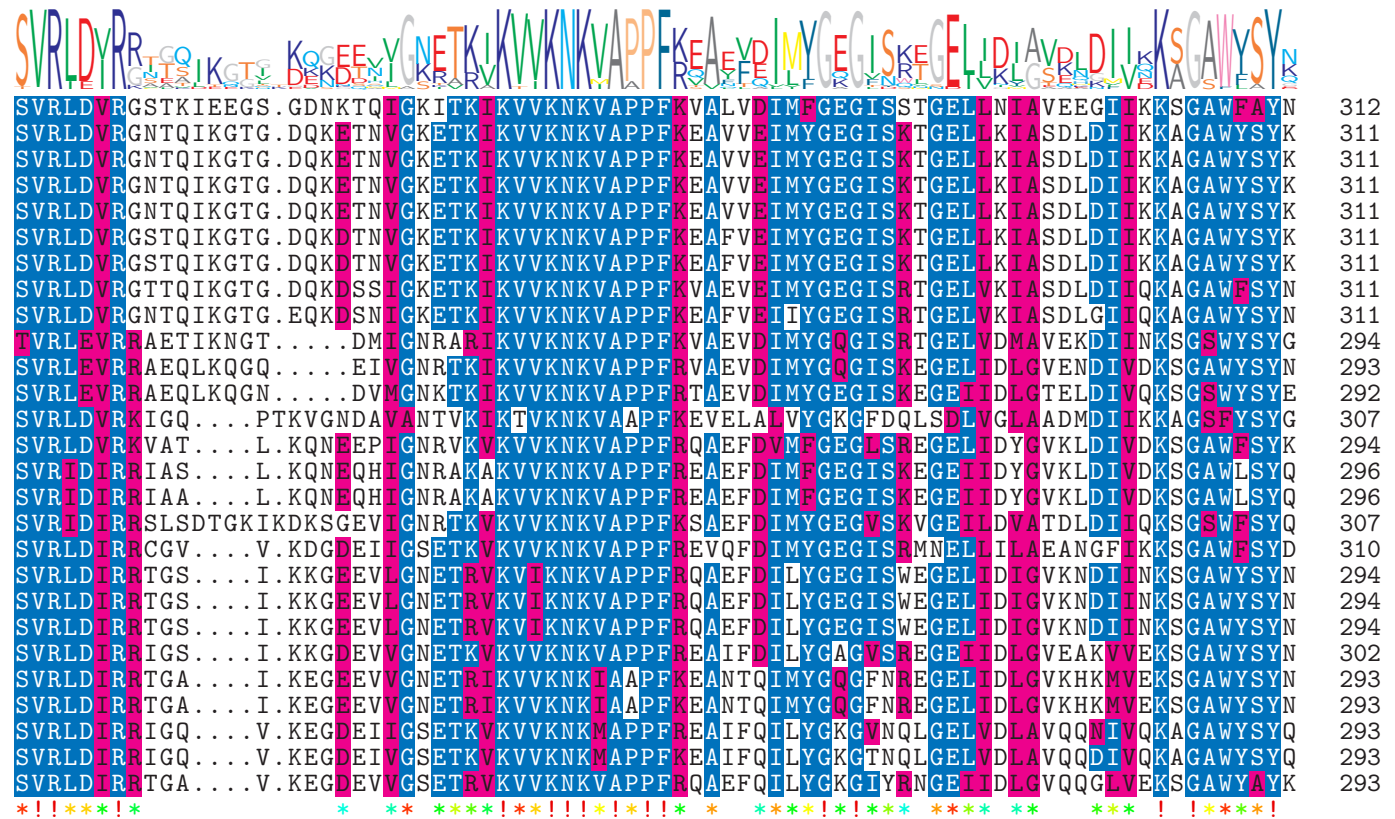
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consensus

Sequence	Score
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DEKIGGGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EIF ⁺ DE ⁺ ID ⁺ DK ⁺ QV ⁺ RS ⁺ KF ⁺ CL ⁺ ID ⁺ GEE ⁺ VSE ⁺ QD ⁺ TEN ⁺ KK ⁺ DE ⁺PK ⁺ KEEA ⁺ ..VNEEV ⁺ PLDLG ⁺ DELEIE	385
DEKIGGGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EIF ⁺ DE ⁺ ID ⁺ DK ⁺ QV ⁺ RS ⁺ KF ⁺ CL ⁺ ID ⁺ GEE ⁺ VSE ⁺ QD ⁺ TEN ⁺ KK ⁺ DE ⁺PK ⁺ KEEA ⁺ ..VNEEV ⁺ PLDLG ⁺ DELEIE	385
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SERIGGG ⁺ REN ⁺ AK ⁺ NYL ⁺ AD ⁺ HED ⁺ VE ⁺ DEV ⁺ RL ⁺ KV ⁺ RA ⁺ AY ⁺ CL ⁺ IS ⁺ DV ⁺ PE ⁺ ED ⁺ LPT ⁺ TE ⁺ DE ⁺ QI ⁺ NIL ⁺ PDD ⁺ STEE ⁺	355
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DKKLGGG ⁺ RENS ⁺ KA ⁺ FL ⁺ KEN ⁺ PE ⁺ IA ⁺ DE ⁺ IT ⁺ KA ⁺ IQ ⁺ NS ⁺ MC ⁺ IE ⁺ GMI ⁺SG ⁺ SEDD ⁺ ..EG ⁺ EE ⁺	343
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logo

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consensus

 non conserved
 similar
 ≥ 50% conserved