

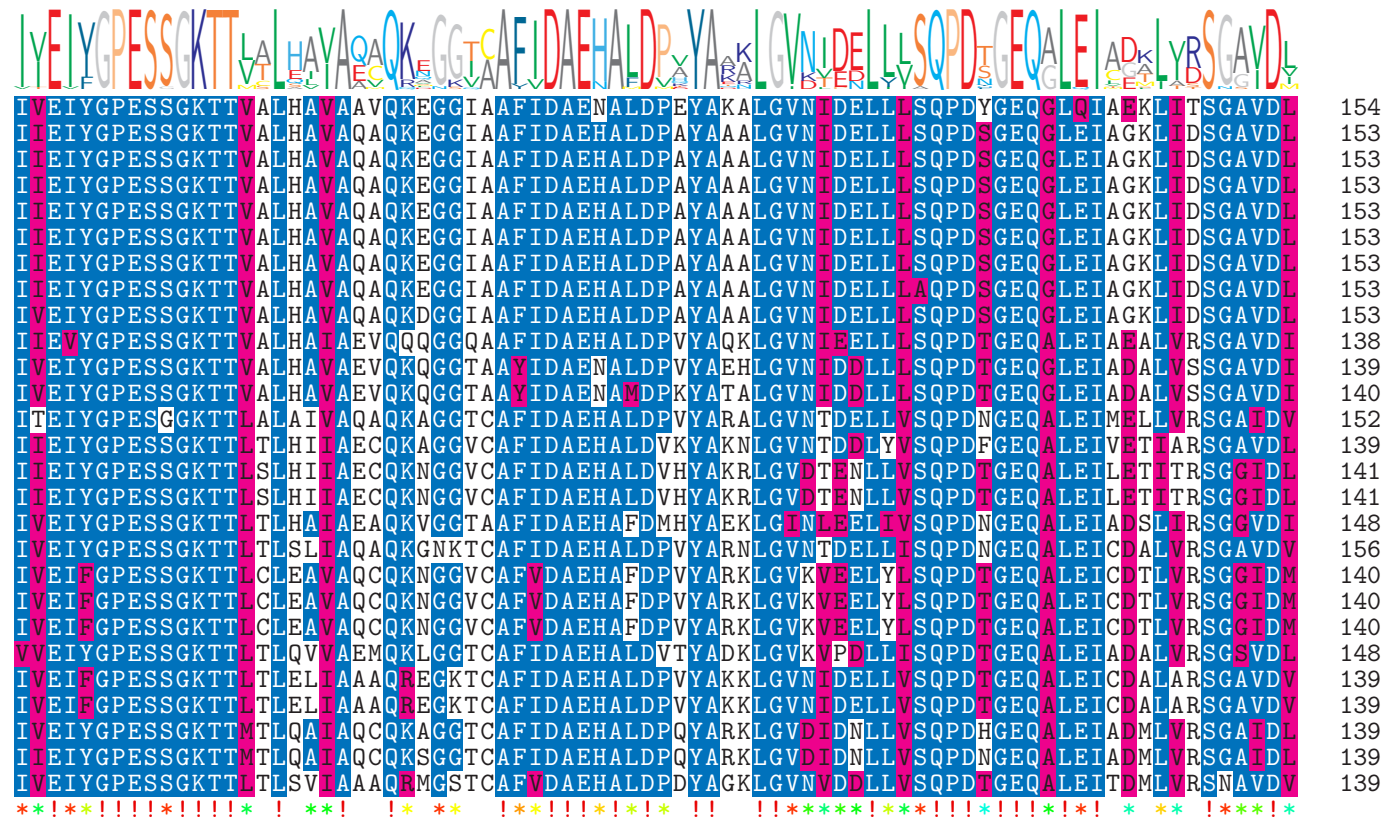
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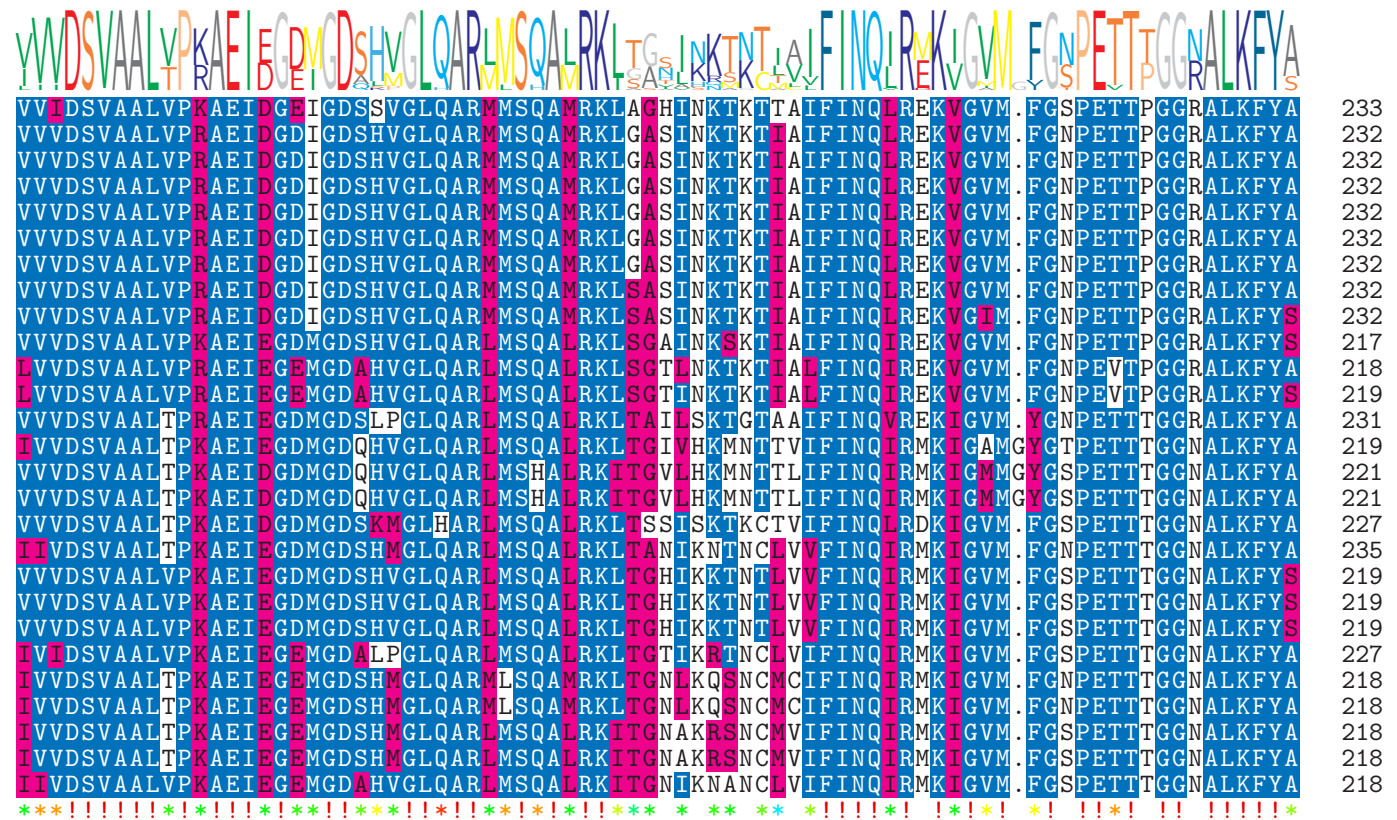
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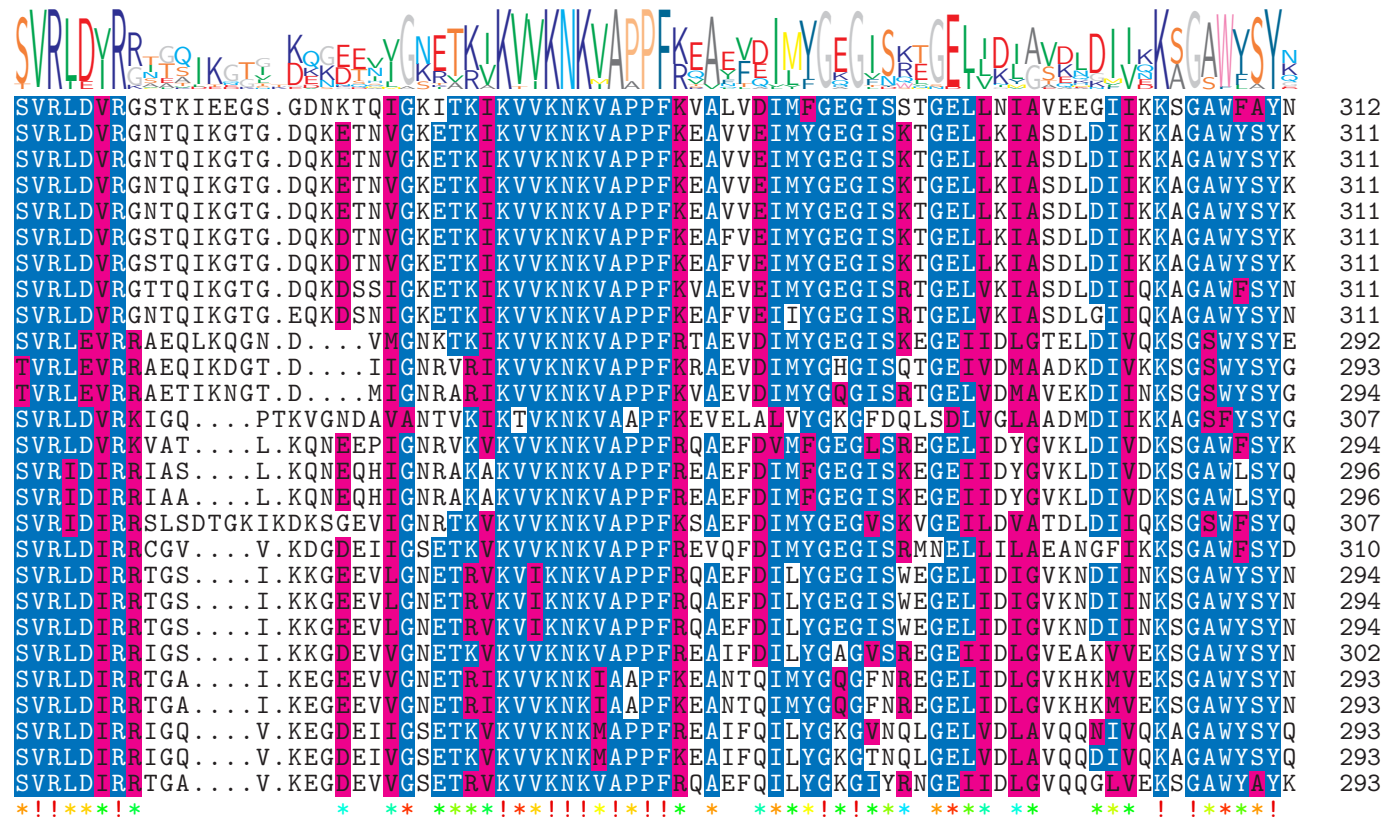
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Sequence	Score
DEKIGGGA ¹ E ² KA ³ KN ⁴ YL ⁵ KE ⁶ HQ ⁷ DV ⁸ FD ⁹ EID ¹⁰ DHK ¹¹ V ¹² RAAH ¹³ GL ¹⁴ LD ¹⁵ DSEVAETE ¹⁶ EET ¹⁷ TASK ¹⁸ TKAKAKKEEK ¹⁹ AVE ²⁰ ..TEE ²¹ IELE ²² LED ²³ ...387	387
DEKIGGGS ¹ EN ² AK ³ KYL ⁴ AE ⁵ HP ⁶ EIF ⁷ DEID ⁸ DKQ ⁹ VR ¹⁰ SKF ¹¹ GL ¹² IDGEEVSEQD ¹³ TENKKDE ¹⁴PKKEEA ¹⁵ ..V ¹⁶ ..NEEVPLDLGDELE ¹⁷ ...383	383
DEKIGGGS ¹ EN ² AK ³ KYL ⁴ AE ⁵ HP ⁶ EIF ⁷ DEID ⁸ DKQ ⁹ VR ¹⁰ SKF ¹¹ GL ¹² IDGEEVSEQD ¹³ TENKKDE ¹⁴PKKEEA ¹⁵ ..V ¹⁶ ..NEEVPLDLGDELE ¹⁷ ...383	383
DEKIGGGS ¹ EN ² AK ³ KYL ⁴ AE ⁵ HP ⁶ EIF ⁷ DEID ⁸ DKQ ⁹ VR ¹⁰ SKF ¹¹ GL ¹² IDGEEVSEQD ¹³ TENKKDE ¹⁴PKKEEA ¹⁵ ..V ¹⁶ ..NEEVPLDLGDELE ¹⁷ ...383	383
DEKIGGGS ¹ EN ² AK ³ KYL ⁴ AD ⁵ NPE ⁶ VFA ⁷ EID ⁸ HQ ⁹ VR ¹⁰ VQF ¹¹ GL ¹² IDGEEAAE ¹³SKKDE ¹⁴VAQVDS ¹⁵ ..V ¹⁶ ..NEEVTLDLGDELE ¹⁷ ...379	379
DEKIGGGS ¹ EN ² AK ³ KYL ⁴ AD ⁵ NPE ⁶ IF ⁷ DEID ⁸ HQ ⁹ VR ¹⁰ VQF ¹¹ GL ¹² IDGEEIAE ¹³SKKDE ¹⁴AAQVDS ¹⁵ ..V ¹⁶ ..NEEVTLDLGDELE ¹⁷ ...379	379
GEKIGGGS ¹ EN ² AK ³ RYL ⁴ AD ⁵ HP ⁶ QIF ⁷ DEID ⁸ DKR ⁹ VR ¹⁰ VKF ¹¹ GL ¹² LEESEESAMAVASE ¹³E ¹⁴TDDLALDLN ¹⁵ NGIE ¹⁶ ...375	375
GEKIGGGS ¹ EN ² AK ³ KFL ⁴ AD ⁵ NPE ⁶ IF ⁷ DDI ⁸ DHK ⁹ VR ¹⁰ VQY ¹¹ GL ¹² IEEDNTEEKQSSKEK ¹³ETDEK ¹⁴ ..A ¹⁵ ..DKNLVLELDDTIE ¹⁶ ...380	380
EE ¹ RL ² LG ³ GG ⁴ RE ⁵ NA ⁶ KQ ⁷ FL ⁸ KE ⁹ NKD ¹⁰ IM ¹¹ LM ¹² IQE ¹³ QIREHY ¹⁴ CL ¹⁵ DNNGVVQQQA ¹⁶ EETQE ¹⁷ ..ELEF ¹⁸EE ¹⁹ ...348	348
EDR ¹ IG ² GG ³ RE ⁴ NA ⁵ KYL ⁶ VE ⁷ HP ⁸ PDV ⁹ MT ¹⁰ EIR ¹¹ QK ¹² VR ¹³ NAY ¹⁴ CM ¹⁵ DASADEETEAVD ¹⁶ GKRTEPKAANDKAAGKAPDKKAAIKL ¹⁷ DLDDTKK ¹⁸ ...373	373
SERIGGG ¹ RE ² NA ³ KEN ⁴ YL ⁵ AD ⁶ HD ⁷ VED ⁸ VR ⁹ LKV ¹⁰ RAAY ¹¹ GIS ¹² DPVEEDLPTTEDEQINI ¹³LPDDSTE ¹⁴354	354
DERIGGG ¹ E ² KTI ³ AI ⁴ YIA ⁵ ER ⁶ PEME ⁷ QIE ⁸ IRDRV ⁹ MAAIRAGNAGEAPALAPAPAAPEAAE ¹⁰ ..A ¹¹363	363
DK ¹ KL ² GG ³ GRE ⁴ NS ⁵ KAF ⁶ L ⁷ KE ⁸ NPEI ⁹ AD ¹⁰ EIT ¹¹ KAI ¹² QNSM ¹³ CT ¹⁴ EGMI ¹⁵SGSEDD ¹⁶ ..EGE ¹⁷ ..E ¹⁸343	343
DK ¹ KL ² GG ³ GRE ⁴ NA ⁵ KAL ⁶ L ⁷ KE ⁸ DKAL ⁹ AD ¹⁰ EIT ¹¹ LK ¹² IKES ¹³ GSNEEI ¹⁴MPLPDE ¹⁵ ..PLE ¹⁶ ..EME ¹⁷347	347
DK ¹ KL ² GG ³ GRE ⁴ NA ⁵ KAL ⁶ L ⁷ KE ⁸ DKAL ⁹ AD ¹⁰ EIT ¹¹ LK ¹² IKES ¹³ GSNEEI ¹⁴MPLPDE ¹⁵ ..PLE ¹⁶ ..EME ¹⁷347	347
DT ¹ KL ² GG ³ GRD ⁴ TV ⁵ KE ⁶ L ⁷ IKD ⁸ NPE ⁹ LA ¹⁰ EEL ¹¹ AKI ¹² REKLQAK ¹³343	343
GT ¹ KIG ² GQ ³ KNN ⁴ AV ⁵ WL ⁶ KE ⁷ NPEI ⁸ AD ⁹ KL ¹⁰ EQE ¹¹ IRNTLA ¹² SNPNS ¹³VLKATASDSND ¹⁴ ..ESDVEEFS ¹⁵EEEL ¹⁶ ...372	372
GAKIGGQ ¹ KD ² NVR ³ VL ⁴ KE ⁵ NPEI ⁶ SD ⁷ EID ⁸ AKI ⁹ RALNG ¹⁰ VEMHI ¹¹TEGTQDET ¹² DGE ¹³ ..R ¹⁴P ¹⁵EE ¹⁶ ...348	348
GAKIGGQ ¹ KD ² NVR ³ VL ⁴ KE ⁵ NPEI ⁶ SD ⁷ EID ⁸ AKI ⁹ RALNG ¹⁰ VEMHI ¹¹TEGTQDET ¹² DGE ¹³ ..R ¹⁴P ¹⁵EE ¹⁶ ...348	348
GAKIGGQ ¹ KD ² NVR ³ VL ⁴ KE ⁵ NPEI ⁶ SD ⁷ EID ⁸ AKI ⁹ RALNG ¹⁰ VEMHI ¹¹TEGTQDET ¹² DGE ¹³ ..R ¹⁴P ¹⁵EE ¹⁶ ...348	348
GE ¹ RIG ² GRD ³ NC ⁴ REF ⁵ LR ⁶ EN ⁷ AE ⁸ LARE ⁹ EN ¹⁰ KV ¹¹ REHL ¹² CV ¹³ TPMG ¹⁴AVTLA ¹⁵ ..EEVEE ¹⁶ ..D ¹⁷352	352
GD ¹ KIG ² GQ ³ KAN ⁴ ACK ⁵ YL ⁶ KE ⁷ NPEI ⁸ AKT ⁹ LD ¹⁰ KKL ¹¹ REMLLN ¹² PENM ¹³QLIAETSSAAD ¹⁴ ..DVEFGAVP ¹⁵EEF ¹⁶ ..354	354
GD ¹ KIG ² GQ ³ KAN ⁴ ACK	

logo

NTDB id 600 KW2 RS01765 WP 011675413.1		387
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 non conserved
 similar
 ≥ 50% conserved