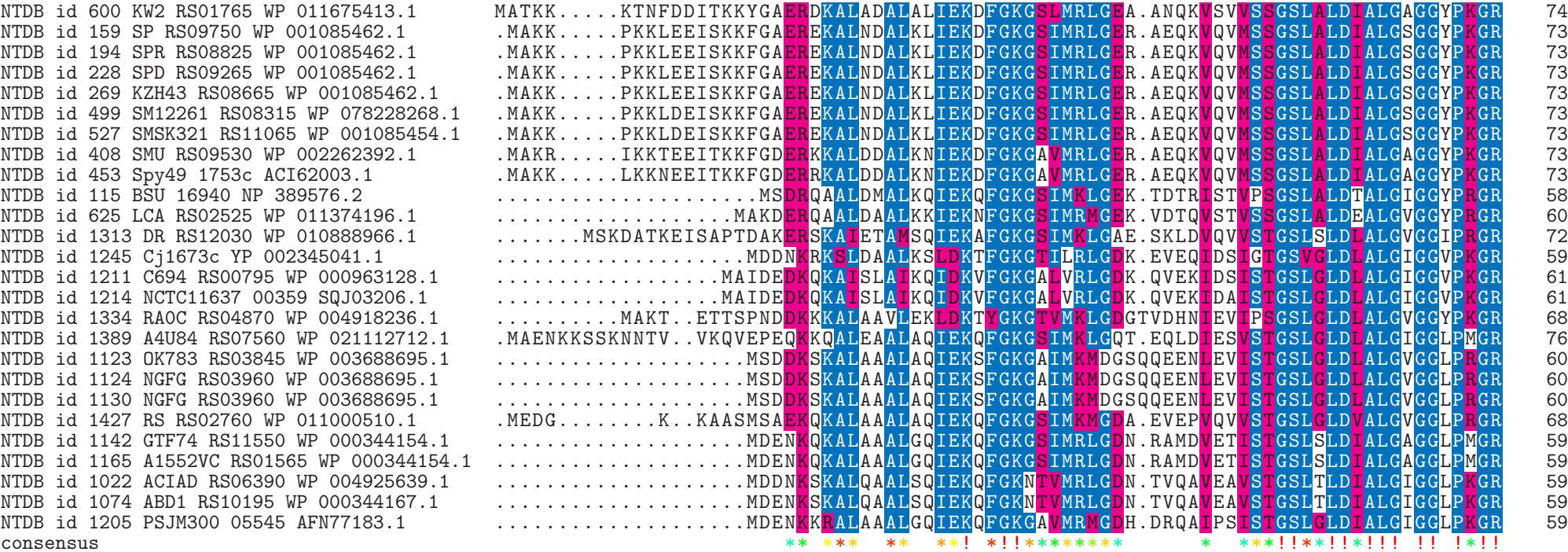


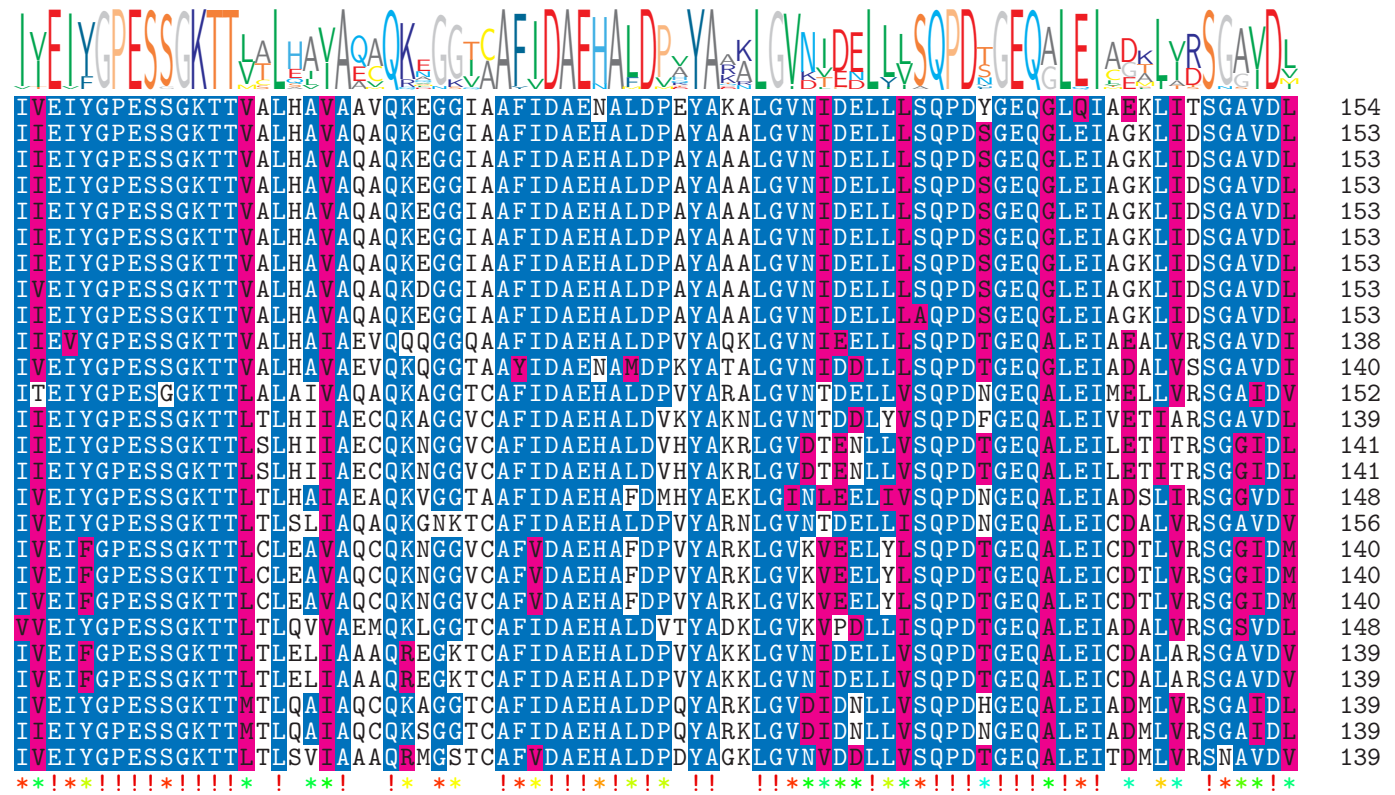
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consensus

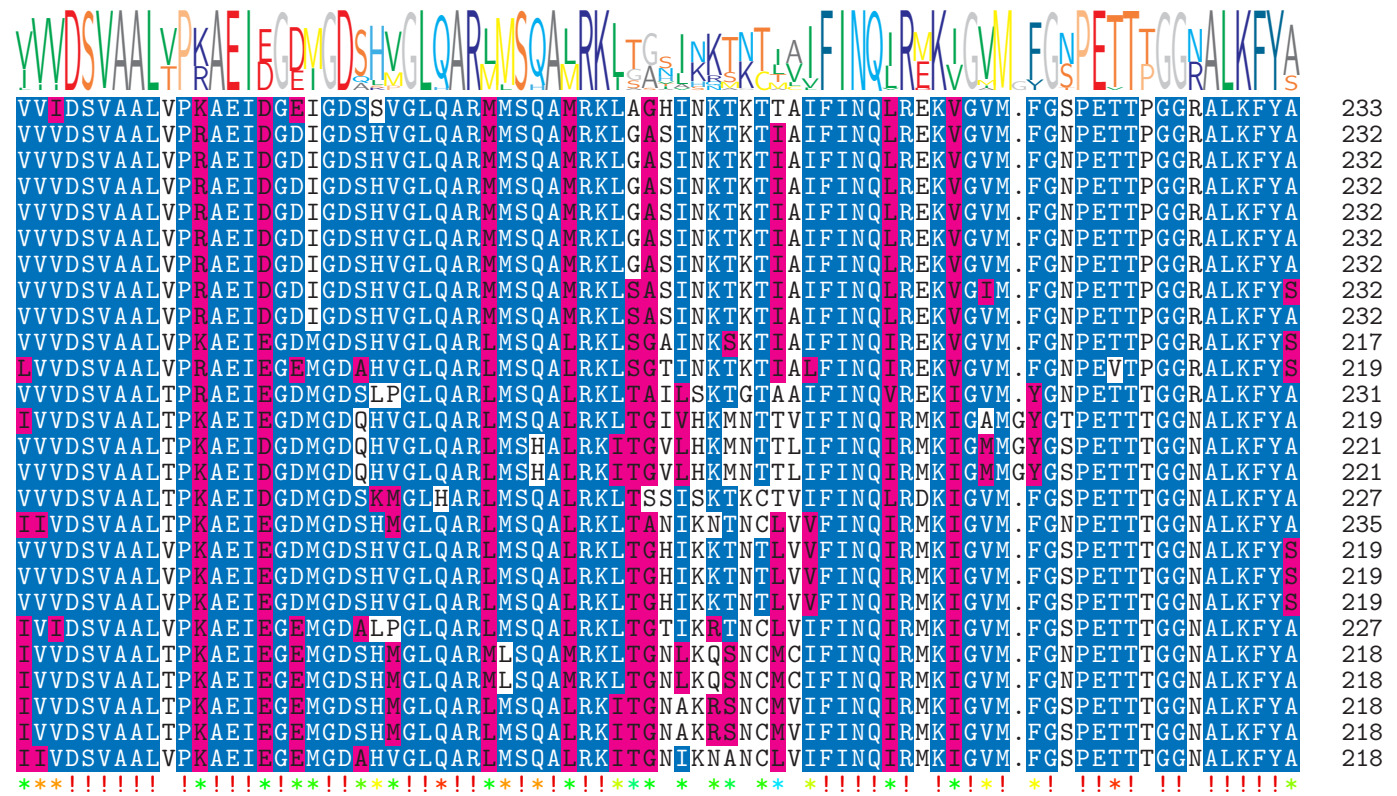
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consensus

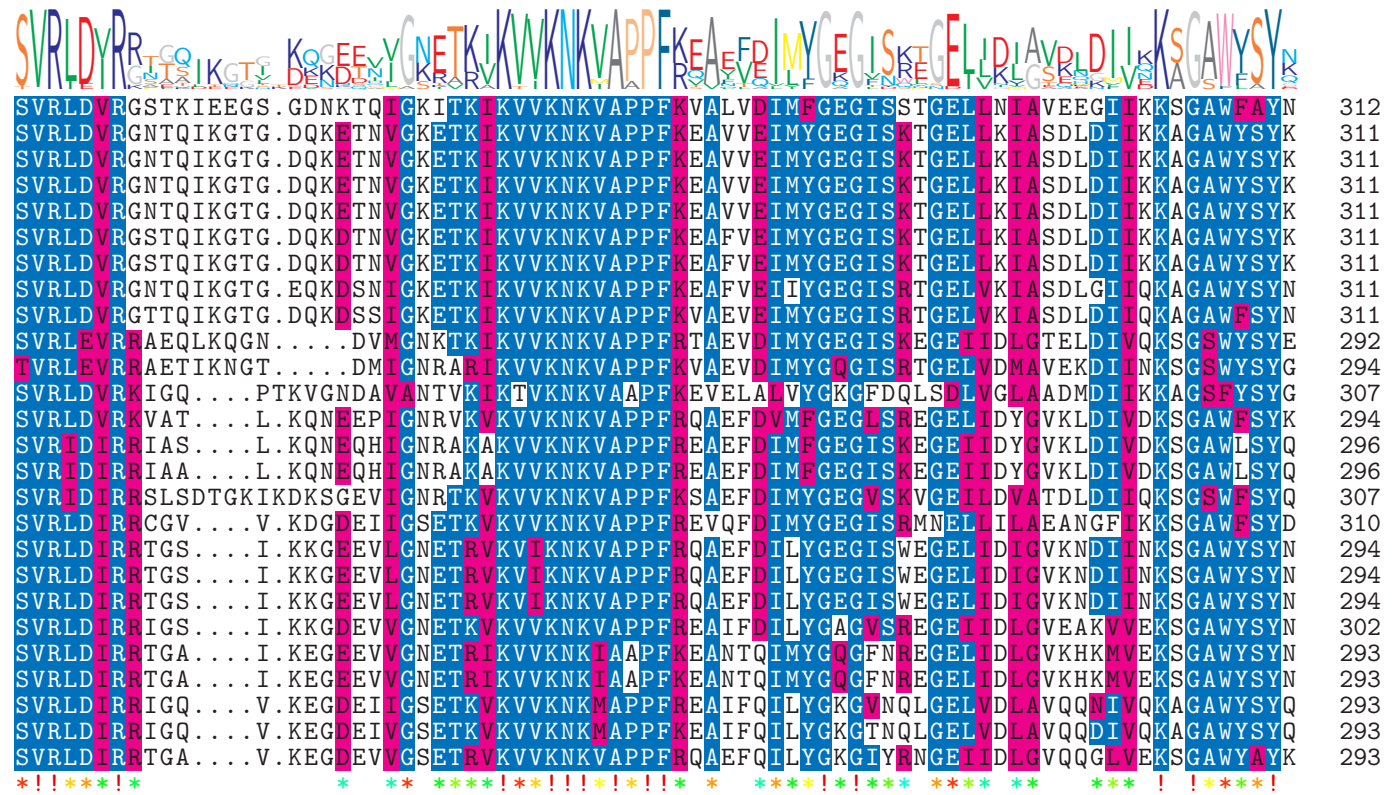
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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Sequence	Score
DEKIGQGA ⁺ E ⁺ KA ⁺ KN ⁺ YL ⁺ KE ⁺ HQ ⁺ DV ⁺ F ⁺ DE ⁺ ID ⁺ HK ⁺ VR ⁺ AAH ⁺ GI ⁺ LDDSEVAETEETTASKTKAKAKKEEKAVETEEIELELED....	387
DEKIGQGS ⁺ ENAK ⁺ KYLA ⁺ EH ⁺ PE ⁺ IF ⁺ DE ⁺ ID ⁺ KQ ⁺ VR ⁺ SKF ⁺ GL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ ENAK ⁺ KYLA ⁺ EH ⁺ PE ⁺ IF ⁺ DE ⁺ ID ⁺ KQ ⁺ VR ⁺ SKF ⁺ GL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ ENAK ⁺ KYLA ⁺ EH ⁺ PE ⁺ IF ⁺ DE ⁺ ID ⁺ KQ ⁺ VR ⁺ SKF ⁺ GL ⁺ IDGEEVSEQDTE ⁺ NKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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GEKIGQGS ⁺ ENAK ⁺ KRYLAD ⁺ HP ⁺ QL ⁺ FD ⁺ ED ⁺ HK ⁺ VR ⁺ VKF ⁺ GL ⁺ LEESEEEESAMAVASE.....E.....TDDLALDLNNGIEIE	377
EER ⁺ LGGGREN ⁺ AK ⁺ FLKEN ⁺ KDI ⁺ MLMI ⁺ QEL ⁺ IREHY ⁺ CL ⁺ DNNGVVQQQAETQEEL.EFE.E.....	348
SER ⁺ IGQGR ⁺ ENAK ⁺ NYLAD ⁺ HED ⁺ VE ⁺ DEV ⁺ LR ⁺ KV ⁺ RAAY ⁺ GI ⁺ SDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
DER ⁺ IGQGE ⁺ KETIAY ⁺ IA ⁺ ERP ⁺ EME ⁺ QEI ⁺ RD ⁺ RVMAA ⁺ IRAGNAGEAPALAPAPAAPEAAE..A.....	363
DKKLGGGREN ⁺ SKA ⁺ FLKEN ⁺ PEI ⁺ ADE ⁺ IT ⁺ KAI ⁺ QNSM ⁺ GI ⁺ EGMI.....SGSEDD..EGE..E.....	343
DKKLGGGREN ⁺ AK ⁺ ALLKED ⁺ KALADE ⁺ IT ⁺ LK ⁺ IKES ⁺ IG ⁺ SNEEI.....MPLPDE..PLE..EME.....	347
DKKLGGGREN ⁺ AK ⁺ ALLKED ⁺ KALADE ⁺ IT ⁺ LK ⁺ IKES ⁺ IG ⁺ SNEEI.....MPLPDE..PLE..EME.....	347
DTKLGGGRD ⁺ TV ⁺ KEL ⁺ IKD ⁺ NPE ⁺ LAE ⁺ LE ⁺ AK ⁺ IREKLQAK.....	343
GTKIGQGK ⁺ NNAVK ⁺ WLKEN ⁺ PEI ⁺ AD ⁺ KL ⁺ EQEI ⁺ RNTLA ⁺ SNPNS.....VLKATASDSND..ESDVEEFSEEEEL.....	372
GAKIGQGK ⁺ KDN ⁺ VR ⁺ VWLKEN ⁺ PEI ⁺ SDE ⁺ ID ⁺ AK ⁺ IRALN ⁺ GVEMHI.....TEGTQDET ⁺ DGE..R.....PEE.....	348
GAKIGQGK ⁺ KDN ⁺ VR ⁺ VWLKEN ⁺ PEI ⁺ SDE ⁺ ID ⁺ AK ⁺ IRALN ⁺ GVEMHI.....TEGTQDET ⁺ DGE..R.....PEE.....	348
GAKIGQGK ⁺ KDN ⁺ VR ⁺ VWLKEN ⁺ PEI ⁺ SDE ⁺ ID ⁺ AK ⁺ IRALN ⁺ GVEMHI.....TEGTQDET ⁺ DGE..R.....PEE.....	348
GER ⁺ IGQGRD ⁺ NCREF ⁺ LRENA ⁺ EL ⁺ LARE ⁺ EN ⁺ KV ⁺ REHL ⁺ GVTPMG.....AVTLA.EEEVE..D.....	352
GDKIGQGK ⁺ KANACK ⁺ YLKEN ⁺ PEI ⁺ AK ⁺ TL ⁺ DKK ⁺ LREMLLN ⁺ PNEM....QLIAETSSAAD..DVEFGAVPEEF.....	354
GDKIGQGK ⁺ KANACK ⁺ YLKEN ⁺ PEI ⁺ AK ⁺ TL ⁺ DKK ⁺ LREMLLN ⁺ PNEM....QLIAETSSAAD..DVEFGAVPEEF.....	354
GNKIGQGK ⁺ KNNVIR ⁺ YLEEN ⁺ PQI ⁺ STE ⁺ IE ⁺ AVI ⁺ REQLLT ⁺ TKASD....QTAA.HDETEE..EPDLL..ES.....	349
GNKIGQGK ⁺ KNNVIR ⁺ YFEEN ⁺ NQI ⁺ AE ⁺ EI ⁺ ERN ⁺ IREQLLT ⁺ TGTN....GAVQ.I.EDEE..EPDLL..LES.....	349
GSKIGQGK ⁺ KANAAK ⁺ YLEEN ⁺ PEI ⁺ GQEI ⁺ EQ ⁺ IRAK ⁺ LLV ⁺ TSAN....TKATPVAEDLA..DADI.....	347

logo

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consensus

 non conserved
 similar
 ≥ 50% conserved