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NTDB id 601 KW2 RS02615 WP 021036805.1
NTDB id 281 KZH43 RS03550 WP 000882517.1
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NTDB id 289 SPD RS10700 WP 001109677.1
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NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

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[illegible]

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NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

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Sequence	Position
...IHL ^Y TSV ^N NG ^I Q ^K Q ^I DL ^C QNCY ^Q ...IM ^K SG ^G Q ^E AL ^F GAGNASNGNSDEPFNP...FND ^I IFSAL ^L ...	69
...IHL ^Y TN ^L NG ^K Q ^K Q ^I DL ^C QNCY ^K ...II ^K TDPNNSL ^F KGM ^T D...LNNRD [.] FDP...FGD ^F FND ^L ...	66
...IHL ^Y TN ^L NG ^K Q ^K Q ^I DL ^C QNCY ^K ...II ^K TDPNNSL ^F KGM ^T D...LNNRD [.] FDP...FGD ^F FND ^L ...	66
...IHL ^Y TN ^L NG ^K Q ^K Q ^I DL ^C QNCY ^K ...II ^K TDPNNSL ^F KGM ^T D...LNNRD [.] FDP...FGD ^F FND ^L ...	66
...IHL ^Y TN ^L NG ^K Q ^K Q ^I DL ^C QNCY ^K ...II ^K TDPNNSL ^F KGM ^T D...LNNRD [.] FDP...FGD ^F FND ^L ...	66
...IHL ^Y TN ^L NG ^K Q ^Q Q ^V DL ^C HNCY ^Q ...IM ^K TDPNNA ^I LRG ^L GD...LTNPNN ^M DP...FS ^E FFNH ^L ...	67
...IHL ^Y AN ^V NG ^K Q ^K Q ^I DL ^C QNCY ^Q ...IM ^K TDP ^F ENG ^P LNHL ^S Q...QNSSS ^I NPF...FD ^D FFGD ^L ...	67
GQEMSQT ^I HYTP ^R AKKV ^I EL ^S MSDEARKLGHSYV ^G TEH ^I L ^L GL ^I RE ^G EGVAARV ^I NNLGVS ^L NKARQ ^V VLQL...141	141
VKVKRSTL ^R FS ^P RAE ^E VM ^T AASFLAIHNNSEAV ^G TEH ^L L ^L YALL ^L QVEDG ^F GLQL ^L KL...QKINIV ^S LRKE ^L LEKR	143
QDET ^F TELP ^F SRR ^L QL ^V FD ^E AEYVASV ^V HAKVL ^G TEH ^V L ^L YAIL ^H DGNALATRIL ^E RAGFSYEDKKDQVKIAALRRN ^L LEER	150
QDET ^F TELP ^F SRR ^L QL ^V FD ^E AEYVASV ^V HAKVL ^G TEH ^V L ^L YAIL ^H DGNALATRIL ^E RAGFSYEDKKDQVKIAALRRN ^L LEER	150
QDET ^F TELP ^F SRR ^L QL ^V FD ^E AEYVASV ^V HAKVL ^G TEH ^V L ^L YAIL ^H DGNALATRIL ^E RAGFSYEDKKDQVKIAALRRN ^L LEER	150
N...D ^I ML ^L EQ ^S HALKRT ^L AEAAAISQVTHAKEV ^G TEH ^V L ^L MAFL ^L NP ^N LLATRIL ^E LVGFHAKDDGESIRLL ^D LRKV ^I ERY	154
ESDI ^I DLRAQSPAL ^E AMLQEAQEIASVTGAV ^E GVSEH ^V LMAFL ^L LHKDLMVCR ^L LEVAGFQYKDDSDKPRII ^D LRRL ^S LERN	156
ESDI ^I DLRAQSPAL ^E AMLQEAQGIASVTGAV ^E GVSEH ^V LMAFL ^L LHKDLMVCR ^L LEVAGFQYKDDSDKPRII ^D LRRL ^S LERN	156

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consensus

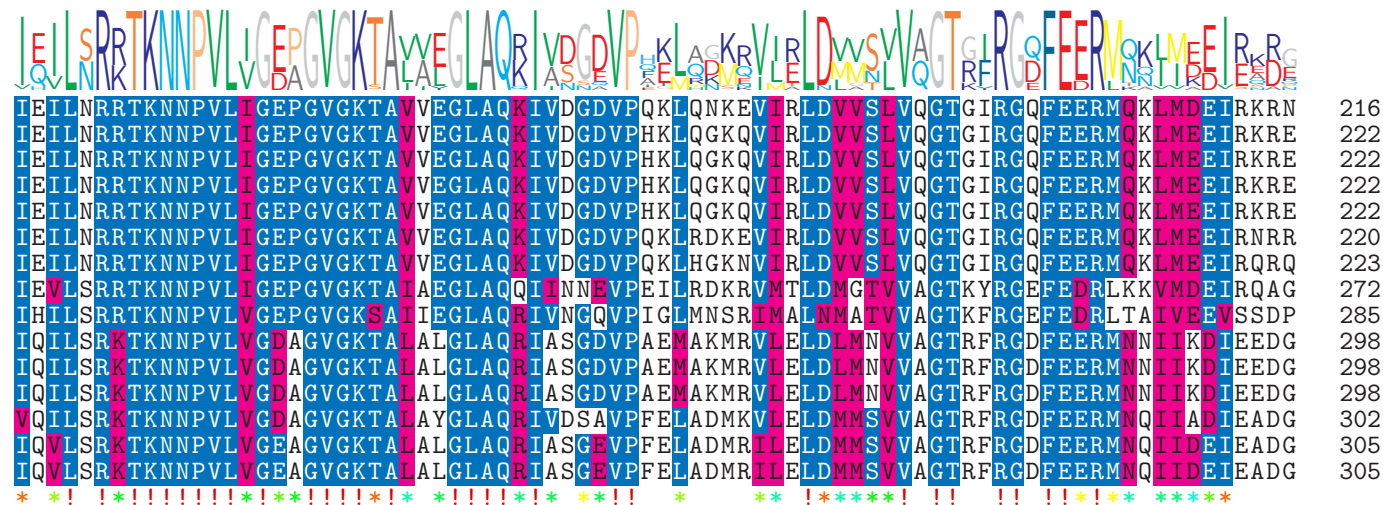
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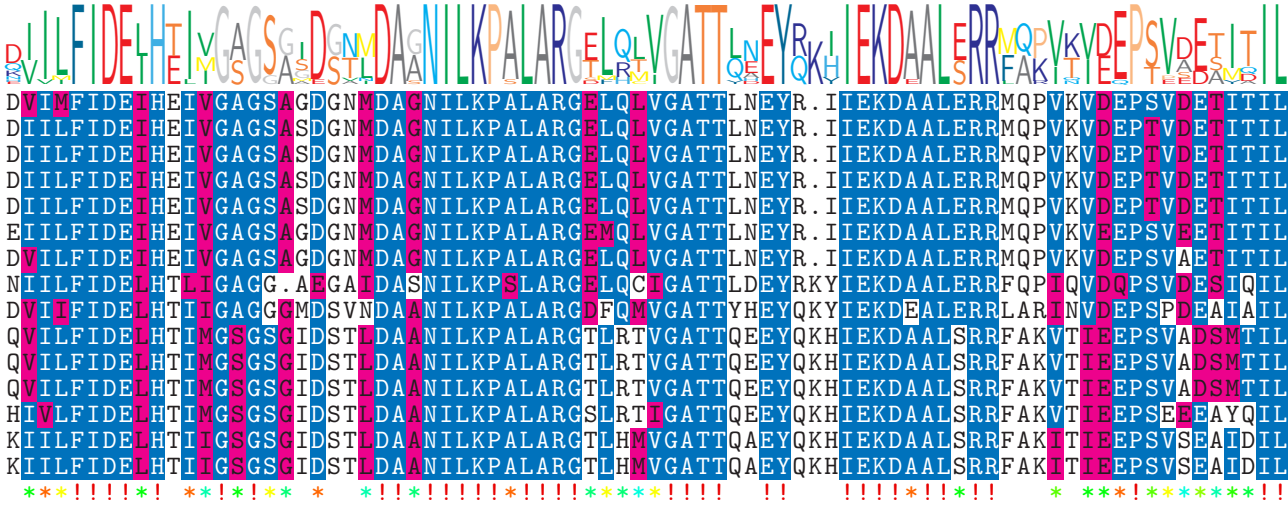
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consensus

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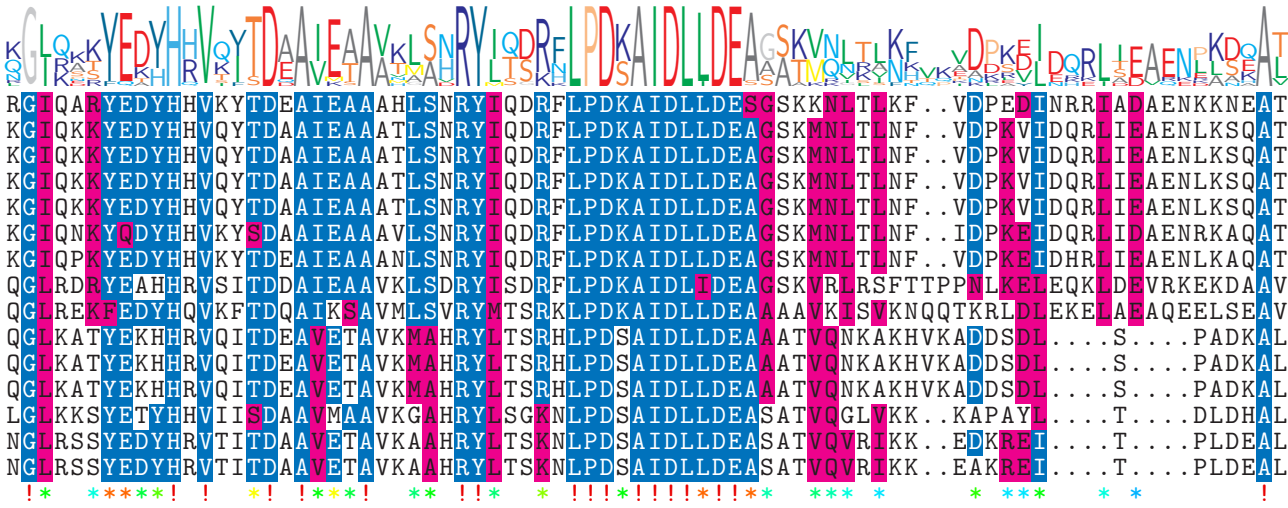
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consensus					



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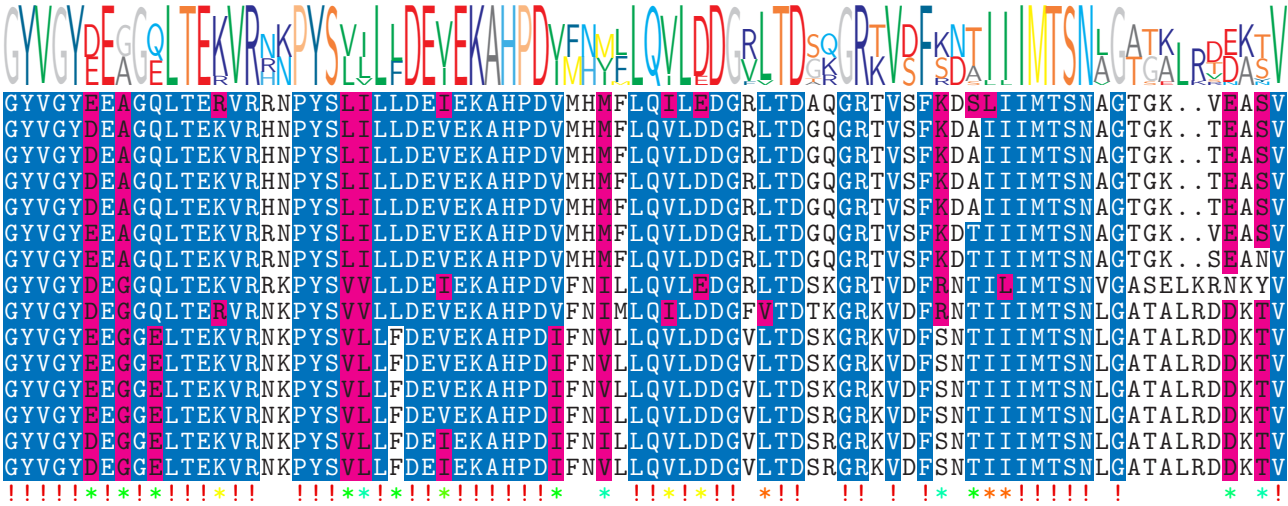
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consensus					



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logo

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consensus					



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751
760
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755
755
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761

logo



X non conserved
X similar
X ≥ 50% conserved