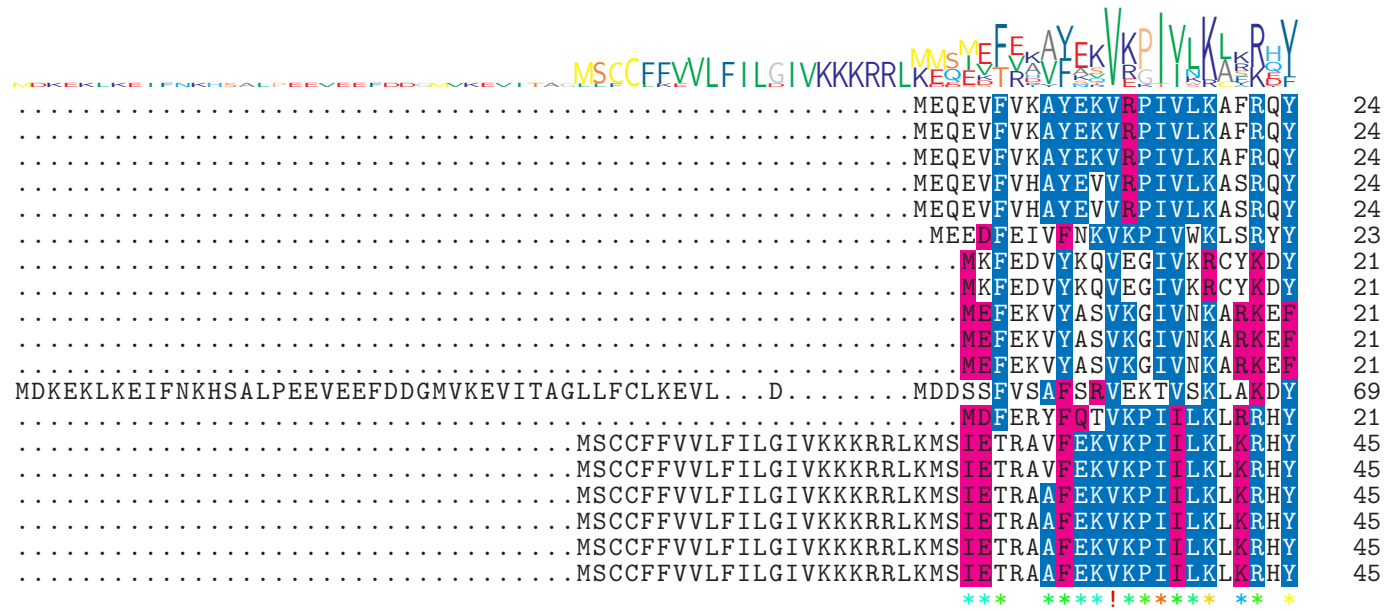


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NTDB id 442 SPYM3 RS08515 WP 011106595.1
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consensus

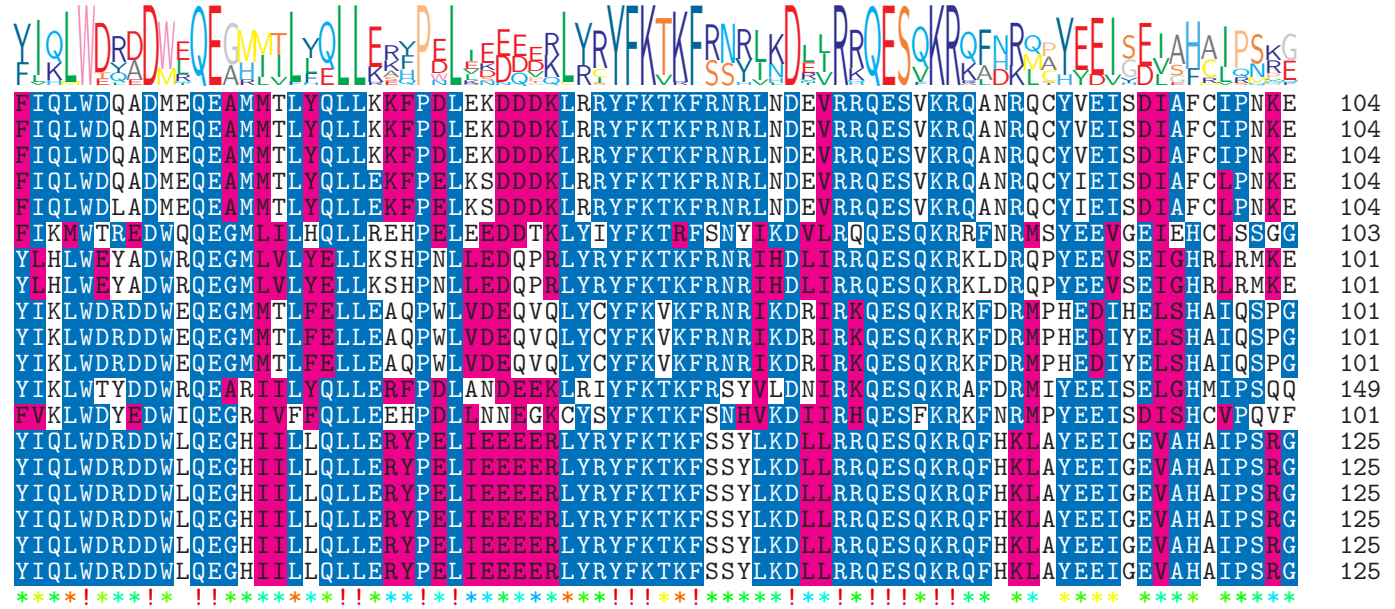
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consensus

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logo

NTDB id 1130591 H0507 RS00860 WP 002949220.1	LD.MVDRLAYDEQLNAFREQLSSEDSLKLDRLIGGECFRGRKKMIRELRFWMVDFDPCNEED	165
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NTDB id 468 HSISS4 RS00725 WP 021143466.1	LD.AVDRLIVYDEQLNAFRNQLSSEDATKLDRLIAGECFRGRKKMIRELKFWMVDFDPYNEDD	165
NTDB id 50 STRSA0001 RS07545 WP 002885610.1	LD.AVDRLIVYDEQLNAFRNQLSSEDATKLDRLIAGECFRGRKKMIRELKFWMVDFDPYNEDD	165
NTDB id 44 SMU RS09085 WP 002263585.1	MQ.LDEYILFRDLSLAYKQGLSTEKQELFERLVAGEHFLGRQSMKDLRKKLSDFKEK....	160
NTDB id 424 SGO RS10430 WP 012130755.1	LY.LDELVAFRSAMSEYRSRLSPEEAKQYERLMADERFKGRKAMLKDLSYHLRDFNPRLD..	160
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NTDB id 461 GPW51 RS00080 WP 011921639.1	LI.NDELLMLRGALRDYRKNSNDQLDKYEKLISGCFNGRREMIRDQLQIHLKDFR.....	156
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NTDB id 582 DLJ51 RS00505 WP 002962720.1	MD.TADYCALKEKELKELKQKLAPEDYAKILLLMRCGAFKGRSAFIKDIKVEFEDFDEDN...	207
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NTDB id 442 SPYM3 RS08515 WP 011106595.1	LW.LDDYVAYQEVIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFKEFAQQL...	183
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NTDB id 450 SPYM18 RS08475 WP 011018214.1	LW.LDDYVAYQEVIASLENQLNSQERMQFKALIRGERFKGRRALLRKISPYFKEFAQQL...	183
NTDB id 449 SPYM18 RS01395 WP 011018214.1	LW.LDDYVAYQEVIASLENQLNSQERMQFKALIRGERFKGRRALLRKISPYFKEFAQQL...	183
consensus	* ** * * * * ! * * * * * * * * * * * * ! * ! * * * * * * * * * * * * !	

- non conserved
- similar
- ≥ 50% conserved