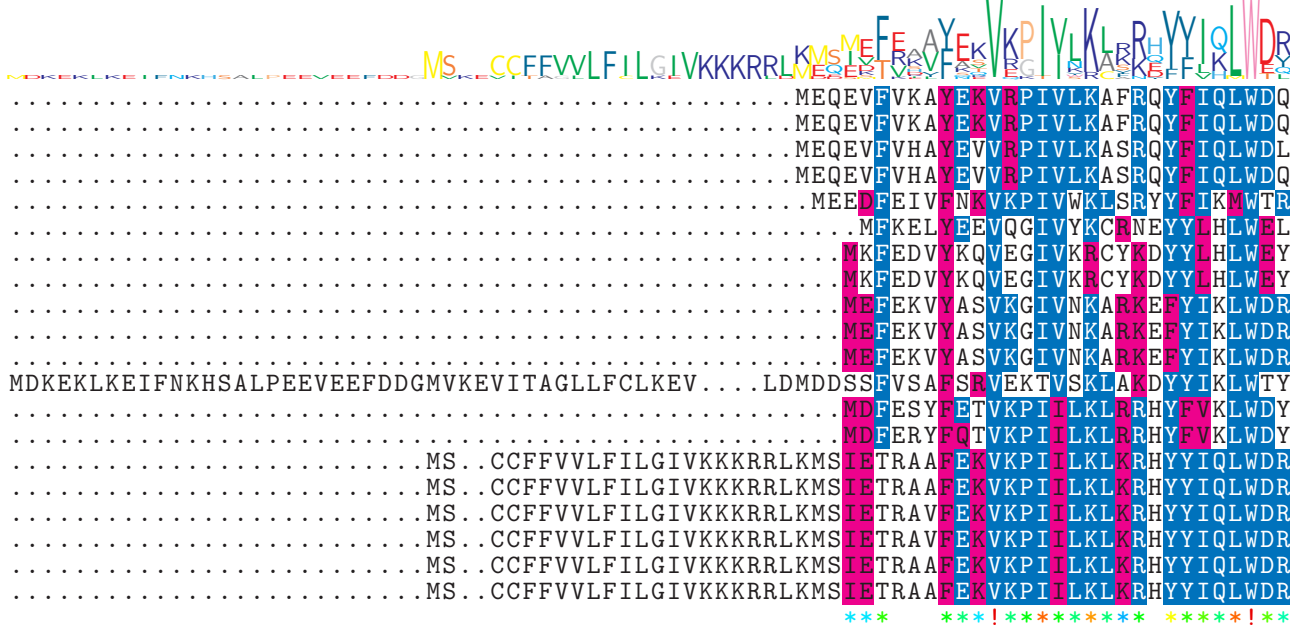


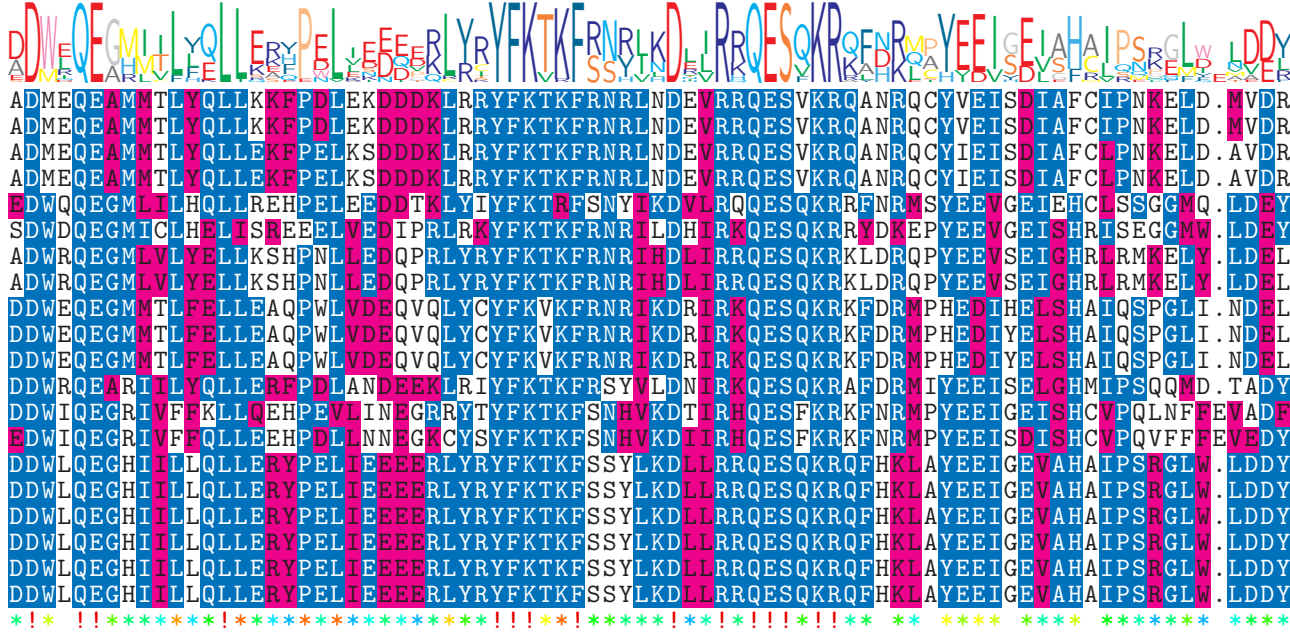
logo

NTDB id 320 STU RS10340 WP 002949220.1
NTDB id 47 STER RS00925 WP 002949220.1
NTDB id 50 STRSA0001 RS07545 WP 002885610.1
NTDB id 468 HSISS4 RS00725 WP 021143466.1
NTDB id 44 SMU RS09085 WP 002263585.1
NTDB id 489 SM12261 RS01070 WP 000471940.1
NTDB id 424 SGO RS10430 WP 012130755.1
NTDB id 423 SGO RS08370 WP 012130755.1
NTDB id 457 SSUD9 RS00080 WP 002936602.1
NTDB id 461 GPW51 RS00080 WP 011921639.1
NTDB id 454 SSU RS00080 WP 011921639.1
NTDB id 582 DLJ51 RS00505 WP 002962720.1
NTDB id 1127554 FGL06 RS01470 WP 024344741.1
NTDB id 554 STRINF RS00370 WP 115283725.1
NTDB id 450 SPYM18 RS08475 WP 011018214.1
NTDB id 449 SPYM18 RS01395 WP 011018214.1
NTDB id 442 SPYM3 RS08515 WP 011106595.1
NTDB id 441 SPYM3 RS01385 WP 011106595.1
NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1
NTDB id 447 SpyM6JRS4 RS01370 WP 010921894.1
consensus



logo

NTDB id 320 STU RS10340 WP 002949220.1
NTDB id 47 STER RS00925 WP 002949220.1
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NTDB id 468 HSISS4 RS00725 WP 021143466.1
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NTDB id 424 SGO RS10430 WP 012130755.1
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NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1
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consensus



logo

NTDB id 320 STU RS10340 WP 002949220.1	LAYDEQLNAFREQLSSEDSLKLDRLIGGECFRGRKKMIRELRFWMVDFDPCNEED	165
NTDB id 47 STER RS00925 WP 002949220.1	LAYDEQLNAFREQLSSEDSLKLDRLIGGECFRGRKKMIRELRFWMVDFDPCNEED	165
NTDB id 50 STRSA0001 RS07545 WP 002885610.1	LVIYDEQLNAFRNQLSSEDA TKLDRLLAGECFRGRKKMIRELKFWMVDFDPYNEDD	165
NTDB id 468 HSISS4 RS00725 WP 021143466.1	LVIYDEQLNAFRNQLSSEDA TKLDRLLAGECFRGRKKMIRELKFWMVDFDPYNEDD	165
NTDB id 44 SMU RS09085 WP 002263585.1	ILFRDSSLAYKQGLSTEKQELFERLVAGEHFLGRQSM LKDLRKKLSDFKEK....	160
NTDB id 489 SM12261 RS01070 WP 000471940.1	YLFHETLRD YRNKQSKDKQEELERVLRHERFRGRQVR LRD LRIVFKEDIRTH..	159
NTDB id 424 SGO RS10430 WP 012130755.1	VAFRSAMSEYRSRLSP EEA KQYERLMADERFKGRKAM LKDL SYHLRDFNPRLD..	160
NTDB id 423 SGO RS08370 WP 012130755.1	VAFRSAMSEYRSRLSP EEA KQYERLMADERFKGRKAM LKDL SYHLRDFNPRLD..	160
NTDB id 457 SSUD9 RS00080 WP 002936602.1	IMLRGALRDYRK NLSNDQLDKYEKLISGQCFNGRREMIRDLQIHLKDFR.....	156
NTDB id 461 GPW51 RS00080 WP 011921639.1	IMLRGALRDYRK NLSNDQLDKYEKLISGQCFNGRREMIRDLQIHLKDFR.....	156
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NTDB id 582 DLJ51 RS00505 WP 002962720.1	CALKEK LKELKQKLAPEDYAKILLLMRCGAFKGRSAFIKD IKVEFEDFDEDN...	207
NTDB id 1127554 FGL06 RS01470 WP 024344741.1	IAYRDSLSQLKATLSLEE QEKLA KVVVRGERFECKKAFLRQIEPYFSDFKH.....	158
NTDB id 554 STRINF RS00370 WP 115283725.1	VAYQDSLACLRR TLGREDRLKLEKVIRGERFECKKAFLKSLIELYFSDFRP.....	158
NTDB id 450 SPYM18 RS08475 WP 011018214.1	VAYQEVIASLENQLNSQERMQFKALIRGERFKGRRALLRKISPYFK EFAQQL...	183
NTDB id 449 SPYM18 RS01395 WP 011018214.1	VAYQEVIASLENQLNSQERMQFKALIRGERFKGRRALLRKISPYFK EFAQQL...	183
NTDB id 442 SPYM3 RS08515 WP 011106595.1	VAYQEVIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFK EFAQQL...	183
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NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1	VAYQEVIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFK EFAQQL...	183
NTDB id 447 SpyM6JRS4 RS01370 WP 010921894.1	VAYQEVIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFK EFAQQL...	183
consensus	*** * * ***** ***** ***** ! * ! ***** * **** !	

- non conserved
- similar
- ≥ 50% conserved