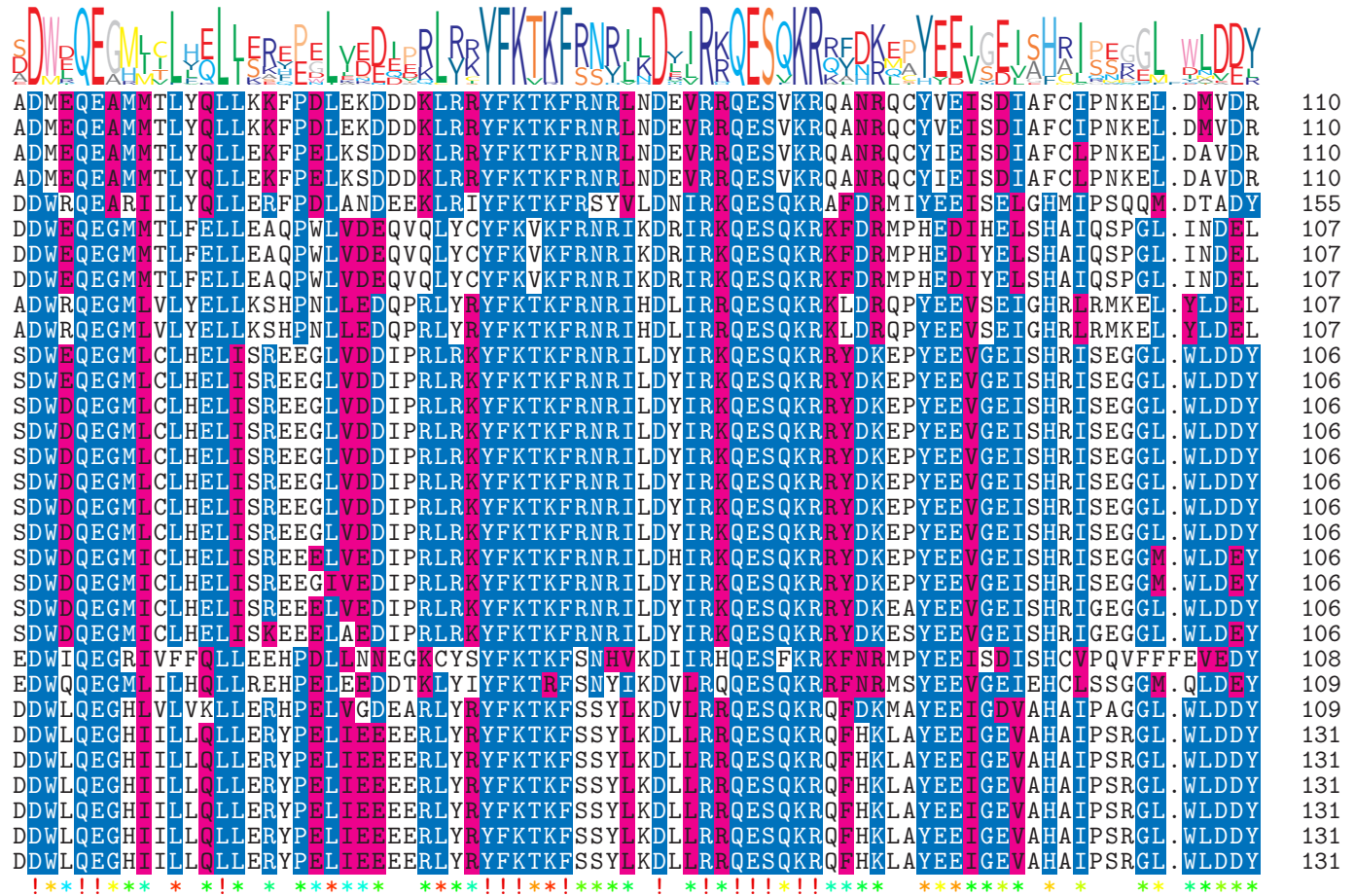


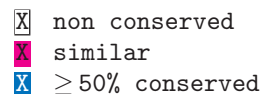
logo

NTDB id 320 STU RS10340 WP 002949220.1	MEQEVFVKAYEKVRPIVLKAFRQYFIQLWDQ	31
NTDB id 47 STER RS00925 WP 002949220.1	MEQEVFVKAYEKVRPIVLKAFRQYFIQLWDQ	31
NTDB id 468 HSISS4 RS00725 WP 021143466.1	MEQEVFVHAYEVVRPIVLKASRQYFIQLWDQ	31
NTDB id 50 STRSA0001 RS07545 WP 002885610.1	MEQEVFVHAYEVVRPIVLKASRQYFIQLWDL	31
NTDB id 582 DLJ51 RS00505 WP 002962720.1	MDKEKLKEIFNKHSAALPEEVEEFDDGMVKEVITAGLLFCLKEV....LDMDDSSFVSAF.....SRVEKTVSKLAKDYYIKLWTY	76
NTDB id 457 SSUD9 RS00080 WP 002936602.1	MEFEKYYASVKGIVNKAARKEFYIKLWDR	28
NTDB id 461 GPW51 RS00080 WP 011921639.1	MEFEKYYASVKGIVNKAARKEFYIKLWDR	28
NTDB id 454 SSU RS00080 WP 011921639.1	MEFEKYYASVKGIVNKAARKEFYIKLWDR	28
NTDB id 424 SGO RS10430 WP 012130755.1	MKFEDVYKQVEGIVKRCYKDYYLHLWEY	28
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NTDB id 252 KZH43 RS09050 WP 000588897.1	MIKELYEEVQCTVYKCRNEYYLHLWEL	27
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NTDB id 44 SMU RS09085 WP 002263585.1	MEEDFEIVFNKVKPIVWKLSRYFIKMWTR	30
NTDB id 34374 SDEG RS01795 WP 003057148.1	MSLETGEVFEKVKPIILKLRHHYYQLWET	30
NTDB id 450 SPYM18 RS08475 WP 011018214.1	MS..CCFFVVLFILGIVKKKRRRLKMSIETRAAF.....FEKVKPIILKLRHHYYIQLWDR	52
NTDB id 449 SPYM18 RS01395 WP 011018214.1	MS..CCFFVVLFILGIVKKKRRRLKMSIETRAAF.....FEKVKPIILKLRHHYYIQLWDR	52
NTDB id 442 SPYM3 RS08515 WP 011106595.1	MS..CCFFVVLFILGIVKKKRRRLKMSIETRAAF.....FEKVKPIILKLRHHYYIQLWDR	52
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consensus	* ** ! ** *	

NTDB id 320 STU RS10340 WP 002949220.1
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consensus



NTDB id 320 STU RS10340 WP 002949220.1
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consensus



X	non conserved
X	similar
X	> 50% conserved