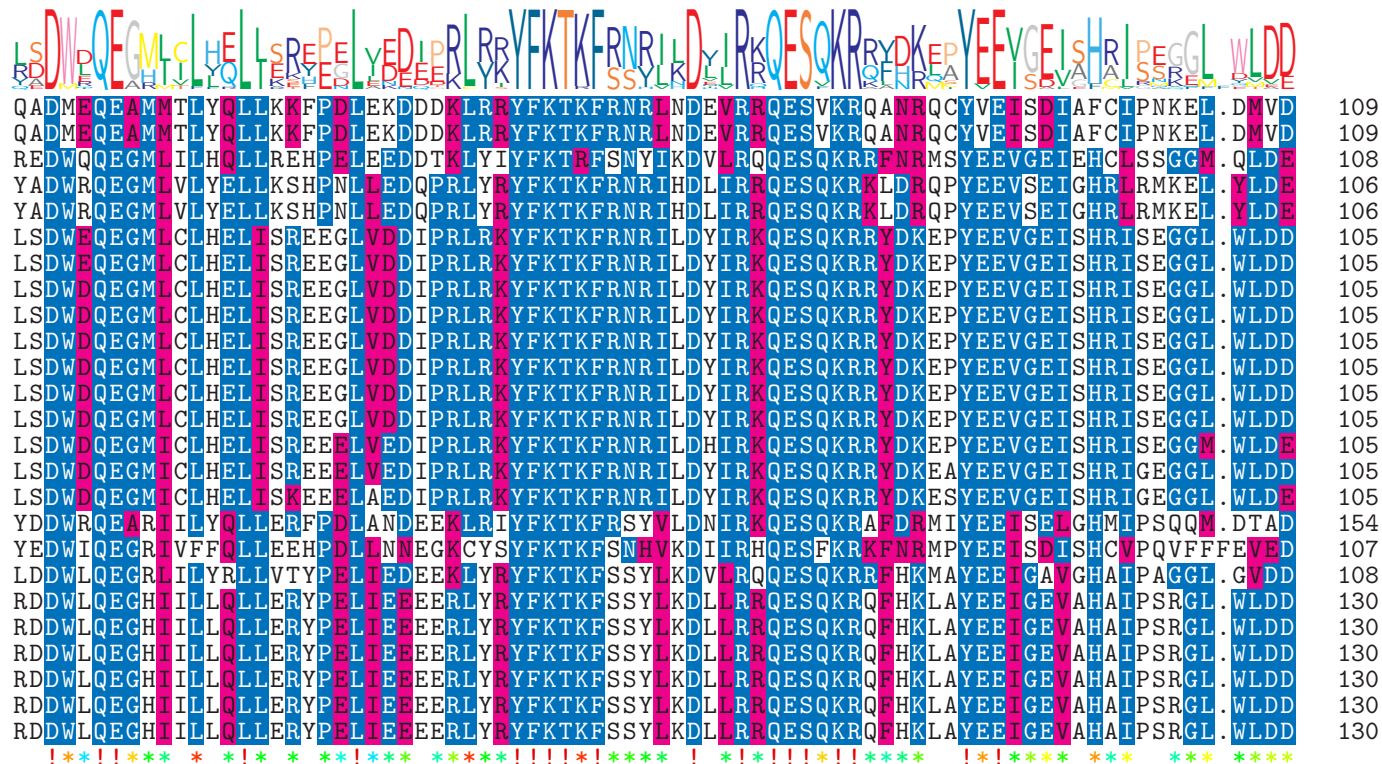


logo

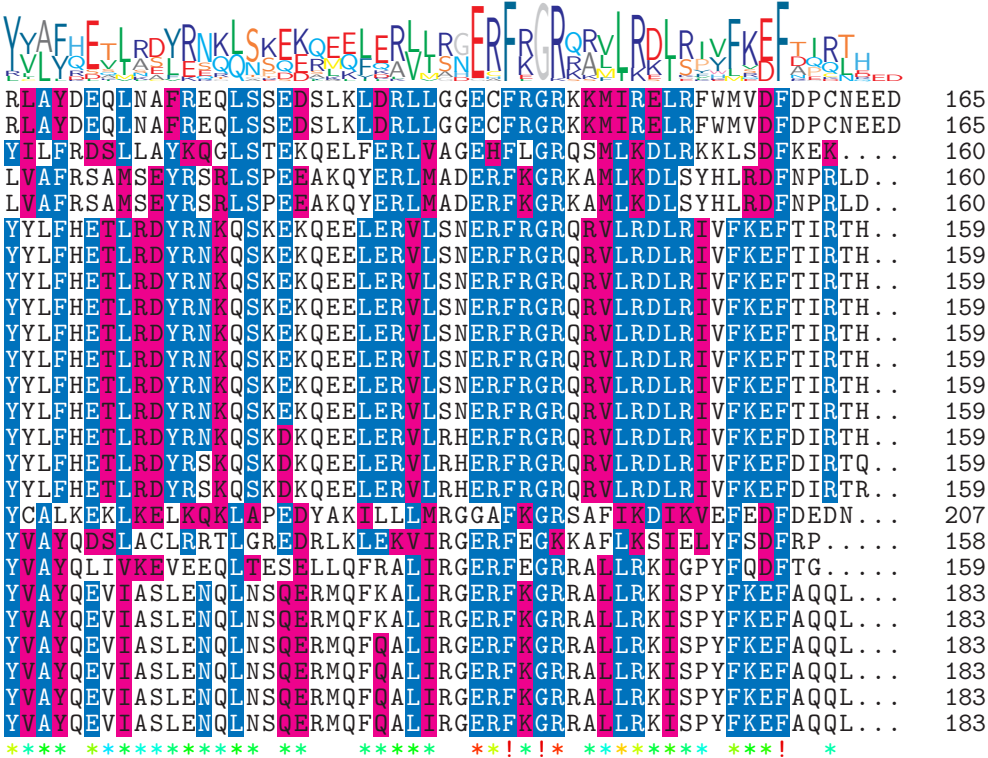
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NTDB id 44 SMU RS09085 WP 002263585.1	MEEDFEIVFNKVKPIVWKLSTYYFIKMT	29
NTDB id 424 SGO RS10430 WP 012130755.1	MKFEDVYKQVEGIVKRCYKDYLLHLWE	27
NTDB id 423 SGO RS08370 WP 012130755.1	MKFEDVYKQVEGIVKRCYKDYLLHLWE	27
NTDB id 141 SP RS10155 WP 000588925.1	MIKELYEEVQCTVYKCRNEYLLHLWE	26
NTDB id 140 SP RS00070 WP 000588925.1	MIKELYEEVQCTVYKCRNEYLLHLWE	26
NTDB id 252 KZH43 RS09050 WP 000588897.1	MIKELYEEVQCTVYKCRNEYLLHLWE	26
NTDB id 251 KZH43 RS09050 WP 000588897.1	MIKELYEEVQCTVYKCRNEYLLHLWE	26
NTDB id 211 SPD RS09675 WP 000588897.1	MIKELYEEVQCTVYKCRNEYLLHLWE	26
NTDB id 210 SPD RS00065 WP 000588897.1	MIKELYEEVQCTVYKCRNEYLLHLWE	26
NTDB id 177 SPR RS09210 WP 000588897.1	MIKELYEEVQCTVYKCRNEYLLHLWE	26
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NTDB id 489 SM12261 RS01070 WP 000471940.1	MFKELYEEVQGIVYKCRNEYLLHLWE	26
NTDB id 517 SMSK321 RS10700 WP 000471945.1	MFKELYKEVQGIVYKSRNEYLLHLWE	26
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NTDB id 554 STRINF RS00370 WP 115283725.1	MDFERYFGTVKPIILKLRHHYFVKLWD	27
NTDB id 1136768 DQL47 RS09065 WP 111690409.1	MSDDLVRILFDKIKPTIMKLRHHYIQLWE	29
NTDB id 450 SPYM18 RS08475 WP 011018214.1	MS..CCFFVVLFILGIVKKKRRRLK.MSIETRAAF EKVKPIILKLRHHYIQLWD	51
NTDB id 449 SPYM18 RS01395 WP 011018214.1	MS..CCFFVVLFILGIVKKKRRRLK.MSIETRAAF EKVKPIILKLRHHYIQLWD	51
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NTDB id 441 SPYM3 RS01385 WP 011106595.1	MS..CCFFVVLFILGIVKKKRRRLK.MSIETRAVFEKVKPIILKLRHHYIQLWD	51
NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1	MS..CCFFVVLFILGIVKKKRRRLK.MSIETRAAF EKVKPIILKLRHHYIQLWD	51
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consensus	* * * * *	

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NTDB id 47 STER RS00925 WP 002949220.1
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consensus



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NTDB	id	320	STU	RS10340	WP	002949220.1
NTDB	id	47	STER	RS00925	WP	002949220.1
NTDB	id	44	SMU	RS09085	WP	002263585.1
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consensus						



X non conserved
X similar
X ≥ 50% conserved