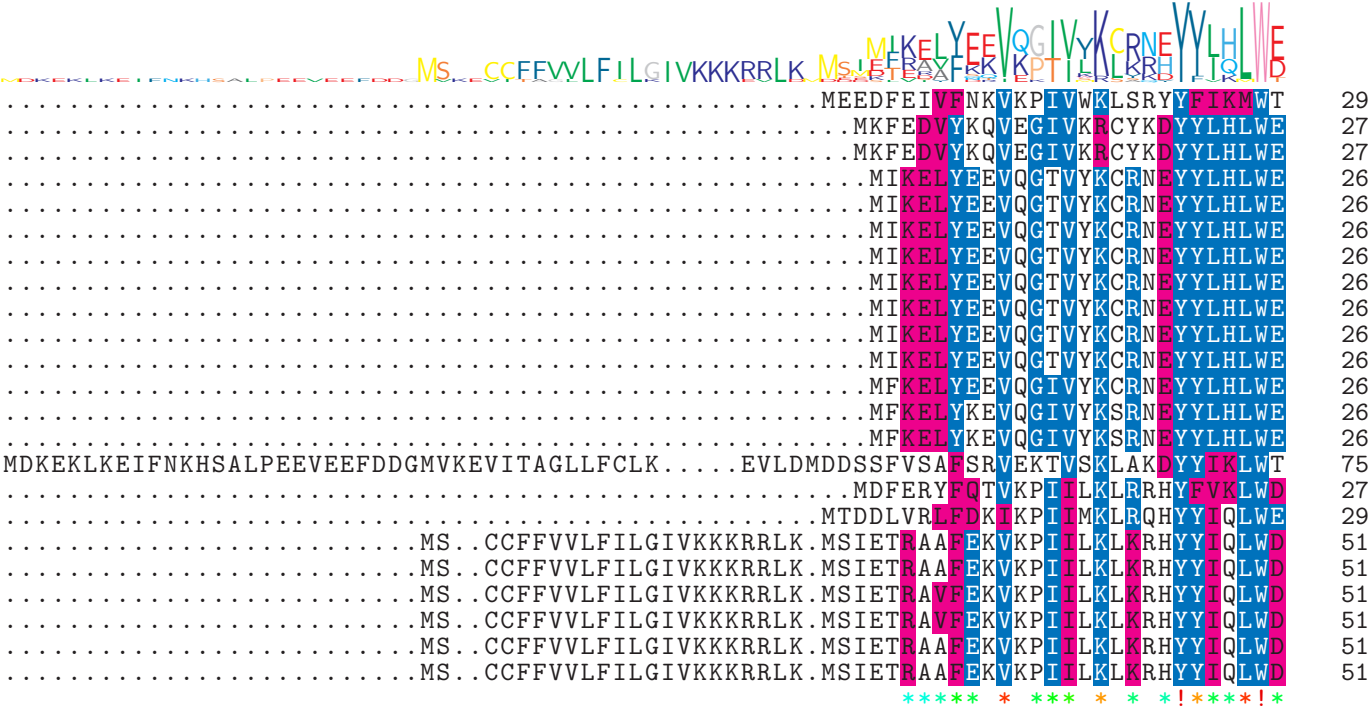


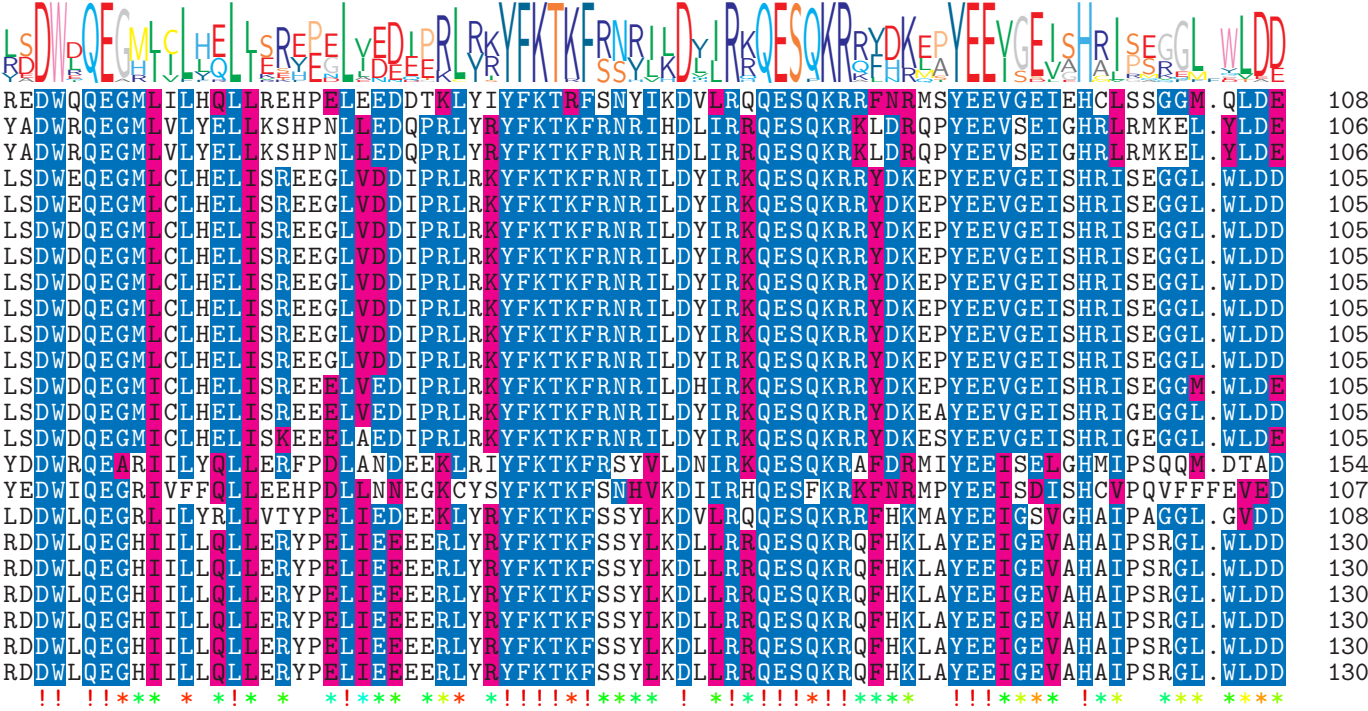
logo

NTDB id 44	SMU RS09085 WP 002263585.1
NTDB id 424	SGO RS10430 WP 012130755.1
NTDB id 423	SGO RS08370 WP 012130755.1
NTDB id 141	SP RS10155 WP 000588925.1
NTDB id 140	SP RS00070 WP 000588925.1
NTDB id 252	KZH43 RS09050 WP 000588897.1
NTDB id 251	KZH43 RS09050 WP 000588897.1
NTDB id 211	SPD RS09675 WP 000588897.1
NTDB id 210	SPD RS00065 WP 000588897.1
NTDB id 177	SPR RS09210 WP 000588897.1
NTDB id 176	SPR RS00065 WP 000588897.1
NTDB id 489	SM12261 RS01070 WP 000471940.1
NTDB id 517	SMSK321 RS10700 WP 000471945.1
NTDB id 516	SMSK321 RS09125 WP 000471944.1
NTDB id 582	DLJ51 RS00505 WP 002962720.1
NTDB id 554	STRINF RS00370 WP 115283725.1
NTDB id 33549	SEQ RS09565 WP 012680087.1
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	



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NTDB id 44	SMU RS09085 WP 002263585.1
NTDB id 424	SGO RS10430 WP 012130755.1
NTDB id 423	SGO RS08370 WP 012130755.1
NTDB id 141	SP RS10155 WP 000588925.1
NTDB id 140	SP RS00070 WP 000588925.1
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NTDB id 554	STRINF RS00370 WP 115283725.1
NTDB id 33549	SEQ RS09565 WP 012680087.1
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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 44 SMU RS09085 WP 002263585.1	YILFRDSLLAYKQGLSTEKQELFERLVAGEHFLGRQSMKKDLKKKLSDFKEK..	160
NTDB id 424 SGO RS10430 WP 012130755.1	LVAFRSAMSEYRSRLSPEEAKQYERLMADERFKGRKAMKKDLSYHLRDFNPRLD	160
NTDB id 423 SGO RS08370 WP 012130755.1	LVAFRSAMSEYRSRLSPEEAKQYERLMADERFKGRKAMKKDLSYHLRDFNPRLD	160
NTDB id 141 SP RS10155 WP 000588925.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 140 SP RS00070 WP 000588925.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 252 KZH43 RS09050 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 251 KZH43 RS09050 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 211 SPD RS09675 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 210 SPD RS00065 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 177 SPR RS09210 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 176 SPR RS00065 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFGRQRVLRDLRIVFKEFTIRTH	159
NTDB id 489 SM12261 RS01070 WP 000471940.1	YYLFHETLRDYRNKQSKDKQEELERVLRHERFGRQRVLRDLRIVFKEFDIRTH	159
NTDB id 517 SMSK321 RS10700 WP 000471945.1	YYLFHETLRDYRSKQSKDKQEELERVLRHERFGRQRVLRDLRIVFKEFDIRTQ	159
NTDB id 516 SMSK321 RS09125 WP 000471944.1	YYLFHETLRDYRSKQSKDKQEELERVLRHERFGRQRVLRDLRIVFKEFDIRTR	159
NTDB id 582 DLJ51 RS00505 WP 002962720.1	YCALKEKLEKELKQKLAPEDYAKILLLMRGGAFFGRSAFTKDIKVEFEDFDEDN.	207
NTDB id 554 STRINF RS00370 WP 115283725.1	YVAYQDSLACLRLTLGREDRCLKLEKVIIRGERFECCKAFLKSLIELYFSDFRP...	158
NTDB id 33549 SEQ RS09565 WP 012680087.1	YVAYQLIVKEVEEHLTEGELSQFRALIRGERFEGRRALLRKIGPYFQDFTGEK.	161
NTDB id 450 SPYM18 RS08475 WP 011018214.1	YVAYQEVIIASLENQLNSQERMQFKALIRGERFKGRRALLRKISPYFKEFAAQL.	183
NTDB id 449 SPYM18 RS01395 WP 011018214.1	YVAYQEVIIASLENQLNSQERMQFKALIRGERFKGRRALLRKISPYFKEFAAQL.	183
NTDB id 442 SPYM3 RS08515 WP 011106595.1	YVAYQEVIIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFKEFAAQL.	183
NTDB id 441 SPYM3 RS01385 WP 011106595.1	YVAYQEVIIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFKEFAAQL.	183
NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1	YVAYQEVIIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFKEFAAQL.	183
NTDB id 447 SpyM6JRS4 RS01370 WP 010921894.1	YVAYQEVIIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFKEFAAQL.	183
consensus	* * ***** * * ***** **!*!* *****! *	

- non conserved
- similar
- ≥ 50% conserved