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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
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 113881 Q426 RS00365 WP 014623459.1
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
NTDB id 554 STRINF RS00370 WP 115283725.1

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MS...CCFFVVLFILGIVKKRRRLK...MS...CFEYFEVGVYKCRNEWYLHLEF

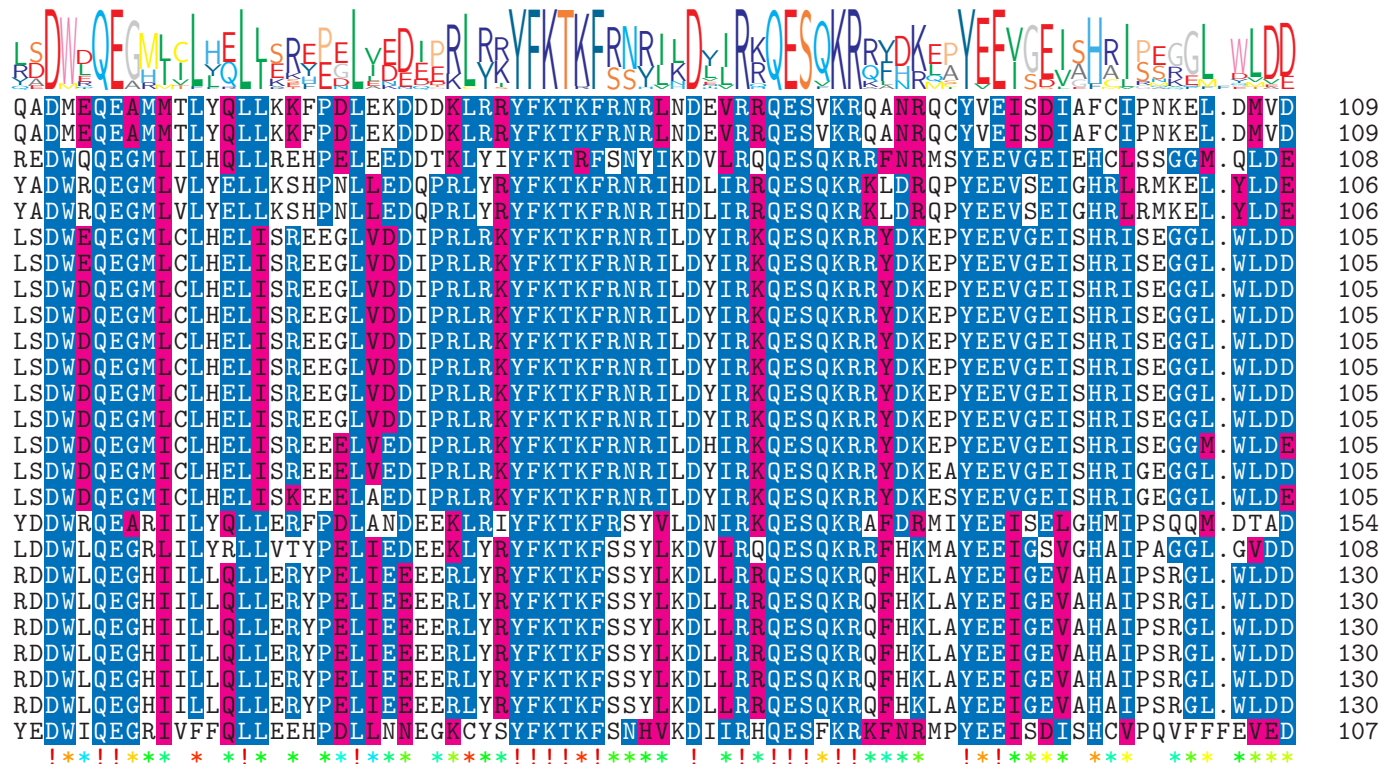
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MEQEV FVKAYEKVRPIVLKAFRQYFIQLWD 30
MEEDFEIVFNKVKPIVWKLSRYYFIKMWT 29
MKFEDVYKQVEGIVKRCYKDYYLHLWE 27
MKFEDVYKQVEGIVKRCYKDYYLHLWE 27
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
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MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYEEVQGTVYKCRNEYYLHLWE 26
MIKELYKEVGIVYKSRNEYYLHLWE 26
MIKELYKEVGIVYKSRNEYYLHLWE 26
MDKEKLKEIFNKHSAPEEVVEEFDDGMVKEVITAGLLFCLK... EVLDMDDSSFVSASF SRVEKT VSKLAKDYYIKLWT 75
MTDDLVR LFDKI KPTIMKL RHYYIQLWE 29
MS...CCFFVVLFILGIVKKRRRLK MSIETRAAF EKVKP IILKLKRHHYIQLWD 51
MS...CCFFVVLFILGIVKKRRRLK MSIETRAAF EKVKP IILKLKRHHYIQLWD 51
MS...CCFFVVLFILGIVKKRRRLK MSIETRAV FEKVKP IILKLKRHHYIQLWD 51
MS...CCFFVVLFILGIVKKRRRLK MSIETRAV FEKVKP IILKLKRHHYIQLWD 51
MS...CCFFVVLFILGIVKKRRRLK MSIETRAAF EKVKP IILKLKRHHYIQLWD 51
MS...CCFFVVLFILGIVKKRRRLK MSIETRAAF EKVKP IILKLKRHHYIQLWD 51
MDFER YFGTVKP IILKL RHYFVKLWD 27

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NTDB id 441 SPYM3 RS01385 WP 011106595.1
NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1
NTDB id 447 SpyM6JRS4 RS01370 WP 010921894.1
NTDB id 554 STRINF RS00370 WP 115283725.1
consensus

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logo

NTDB id 320 STU RS10340 WP 002949220.1	RLAYDEQLNAFREQLSSESLKLDRLLGGECEFRGRKKMIRELRFWMVDFDPCNEED	165
NTDB id 47 STER RS00925 WP 002949220.1	RLAYDEQLNAFREQLSSESLKLDRLLGGECEFRGRKKMIRELRFWMVDFDPCNEED	165
NTDB id 44 SMU RS09085 WP 002263585.1	YILFRDSL LAYKQGLSTEQELFERLVAGEHFLGRQSM LKDLRKKLSDFKEK....	160
NTDB id 424 SGO RS10430 WP 012130755.1	LVAFRSAMSEYRSRLSPEEAKQYERLMADERFKGRKAM LKDL SYHLRDFNPRLD..	160
NTDB id 423 SGO RS08370 WP 012130755.1	LVAFRSAMSEYRSRLSPEEAKQYERLMADERFKGRKAM LKDL SYHLRDFNPRLD..	160
NTDB id 141 SP RS10155 WP 000588925.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 140 SP RS00070 WP 000588925.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 252 KZH43 RS09050 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 251 KZH43 RS09050 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 211 SPD RS09675 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 210 SPD RS00065 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 177 SPR RS09210 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 176 SPR RS00065 WP 000588897.1	YYLFHETLRDYRNKQSKEKQEELERVLSNERFRGRQRVLRDLRIVFKEFTTIRTH..	159
NTDB id 489 SM12261 RS01070 WP 000471940.1	YYLFHETLRDYRNKQSKDKQEELERVLRHERFRGRQRVLRDLRIVFKEFDIRTH..	159
NTDB id 517 SMSK321 RS10700 WP 000471945.1	YYLFHETLRDYRSKQSKDKQEELERVLRHERFRGRQRVLRDLRIVFKEFDIRTQ..	159
NTDB id 516 SMSK321 RS09125 WP 000471944.1	YYLFHETLRDYRSKQSKDKQEELERVLRHERFRGRQRVLRDLRIVFKEFDIRT..	159
NTDB id 582 DLJ51 RS00505 WP 002962720.1	YCALKEK LKELKQKLAPEDYAK LLLMRGGAFKGRSAFLKDKVFEDEFDEDN...	207
NTDB id 113881 Q426 RS00365 WP 014623459.1	YVAYQLIVKEVEERL TESELSQFRALIRGERFEGRRALLRK L GPFYQDFTGEK...	161
NTDB id 450 SPYM18 RS08475 WP 011018214.1	YVAYQEV IASLENQLNSQERMQFKALIRGERFKGRRALLRK L SPYFKEFAAQL...	183
NTDB id 449 SPYM18 RS01395 WP 011018214.1	YVAYQEV IASLENQLNSQERMQFKALIRGERFKGRRALLRK L SPYFKEFAAQL...	183
NTDB id 442 SPYM3 RS08515 WP 011106595.1	YVAYQEV IASLENQLNSQERMQFQALIRGERFKGRRALLRK L SPYFKEFAAQL...	183
NTDB id 441 SPYM3 RS01385 WP 011106595.1	YVAYQEV IASLENQLNSQERMQFQALIRGERFKGRRALLRK L SPYFKEFAAQL...	183
NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1	YVAYQEV IASLENQLNSQERMQFQALIRGERFKGRRALLRK L SPYFKEFAAQL...	183
NTDB id 447 SpyM6JRS4 RS01370 WP 010921894.1	YVAYQEV IASLENQLNSQERMQFQALIRGERFKGRRALLRK L SPYFKEFAAQL...	183
NTDB id 554 STRINF RS00370 WP 115283725.1	YVAYQDSLACL RRT L GREDRLK L EKVI RGERFECKKAF L KSL E I VESDERP.....	158
consensus	***** ***** ** ***** *!*!* ***** **!* *	

- non conserved
- similar
- ≥ 50% conserved