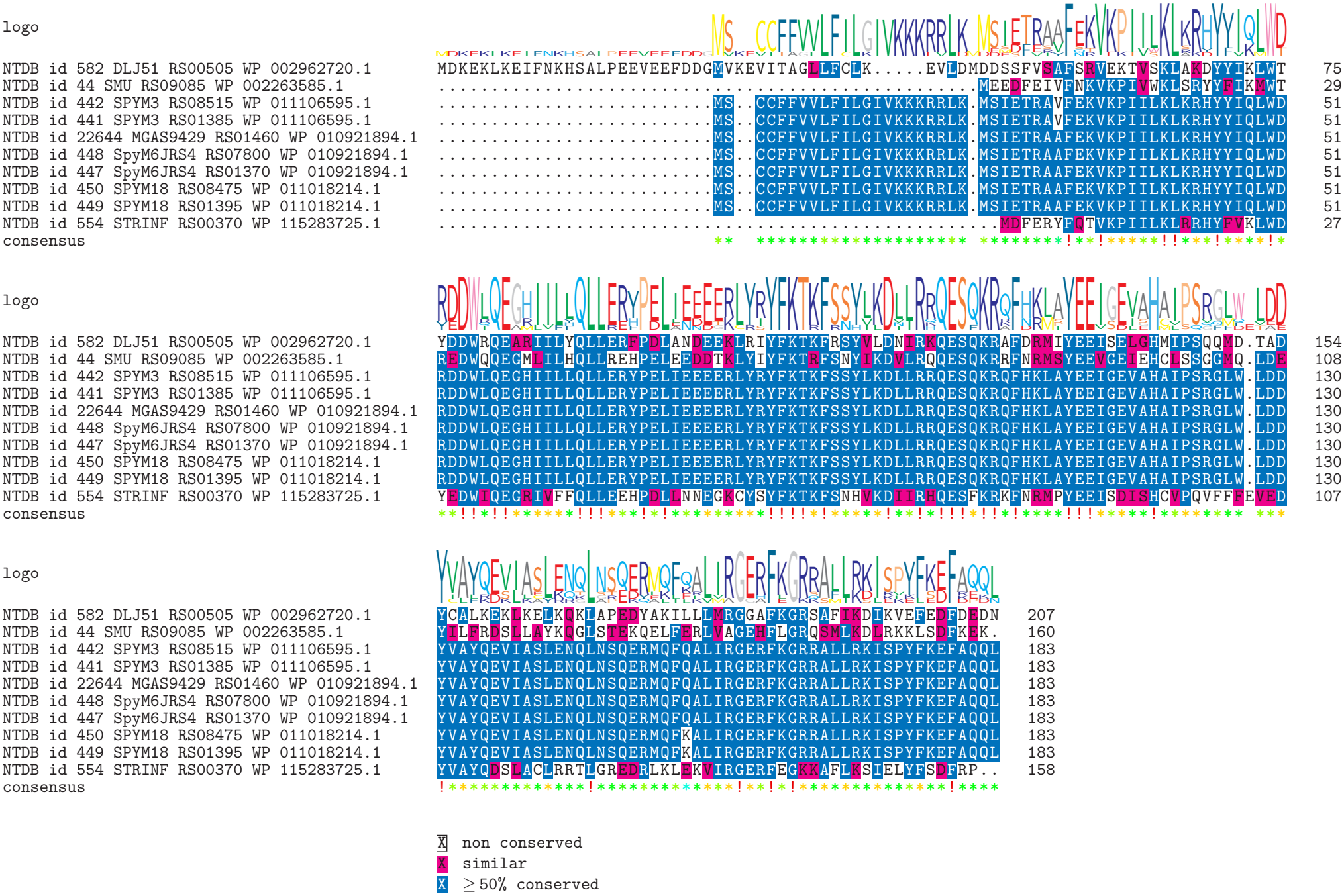
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NTDB id 44 SMU RS09085 WP 002263585.1 .....MEEDFEIVFNKVKPIVWKLSRYYFIKMWT 29
NTDB id 442 SPYM3 RS08515 WP 011106595.1 .....MS..CCFFVVLFILGIVKKKRRLK.MSIETRAVFEKVKPIILKLKRHYYIQLWD 51
NTDB id 441 SPYM3 RS01385 WP 011106595.1 .....MS..CCFFVVLFILGIVKKKRRLK.MSIETRAVFEKVKPIILKLKRHYYIQLWD 51
NTDB id 22644 MGAS9429 RS01460 WP 010921894.1 .....MS..CCFFVVLFILGIVKKKRRLK.MSIETRAAFEKVKPIILKLKRHYYIQLWD 51
NTDB id 448 SpyM6JRS4 RS07800 WP 010921894.1 .....MS..CCFFVVLFILGIVKKKRRLK.MSIETRAAFEKVKPIILKLKRHYYIQLWD 51
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NTDB id 554 STRINF RS00370 WP 115283725.1 .....MDFERYFQTVKPIILKLRRHYFVKLWD 27
consensus
logo
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NTDB id 442 SPYM3 RS08515 WP 011106595.1 RDDWLQEGHIILLQLLERYPELIEEEERLYRYFKTKFSSYLKDLLRRQESQKRQFHKLAYEEIGEVAHAIPSRGLW.LDD 130
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NTDB id 449 SPYM18 RS01395 WP 011018214.1 RDDWLQEGHIILLQLLERYPELIEEEERLYRYFKTKFSSYLKDLLRRQESQKRQFHKLAYEEIGEVAHAIPSRGLW.LDD 130
NTDB id 554 STRINF RS00370 WP 115283725.1 YEDWIQEGRIVFFQLLEEHPDLLNNEGKCYSYFKTKFSNHVKDIIRHQESFKRKFNRMPYEEISDISHCVPQVFFFEVED 107
consensus
logo
NTDB id 582 DLJ51 RS00505 WP 002962720.1 YCALKEKLKELKQKLAPEDYAKILLLMRGGAFKGRSAFIKDIKVEFEDFDFDN 207
NTDB id 44 SMU RS09085 WP 002263585.1 YILFRDSLLAYKQGLSTEKQELFERLVAGEHFLGRQSMLKDLRKKLSDFKEK. 160
NTDB id 442 SPYM3 RS08515 WP 011106595.1 YVAYQEVIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFKEFAQQL 183
NTDB id 441 SPYM3 RS01385 WP 011106595.1 YVAYQEVIASLENQLNSQERMQFQALIRGERFKGRRALLRKISPYFKEFAQQL 183
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NTDB id 554 STRINF RS00370 WP 115283725.1 YVAYQDSLACLRRTLGREDRLKLEKVIRGERFEGKKAFLKSIELYFSDFRP.. 158
consensus
X non conserved
X similar
X ≥ 50% conserved