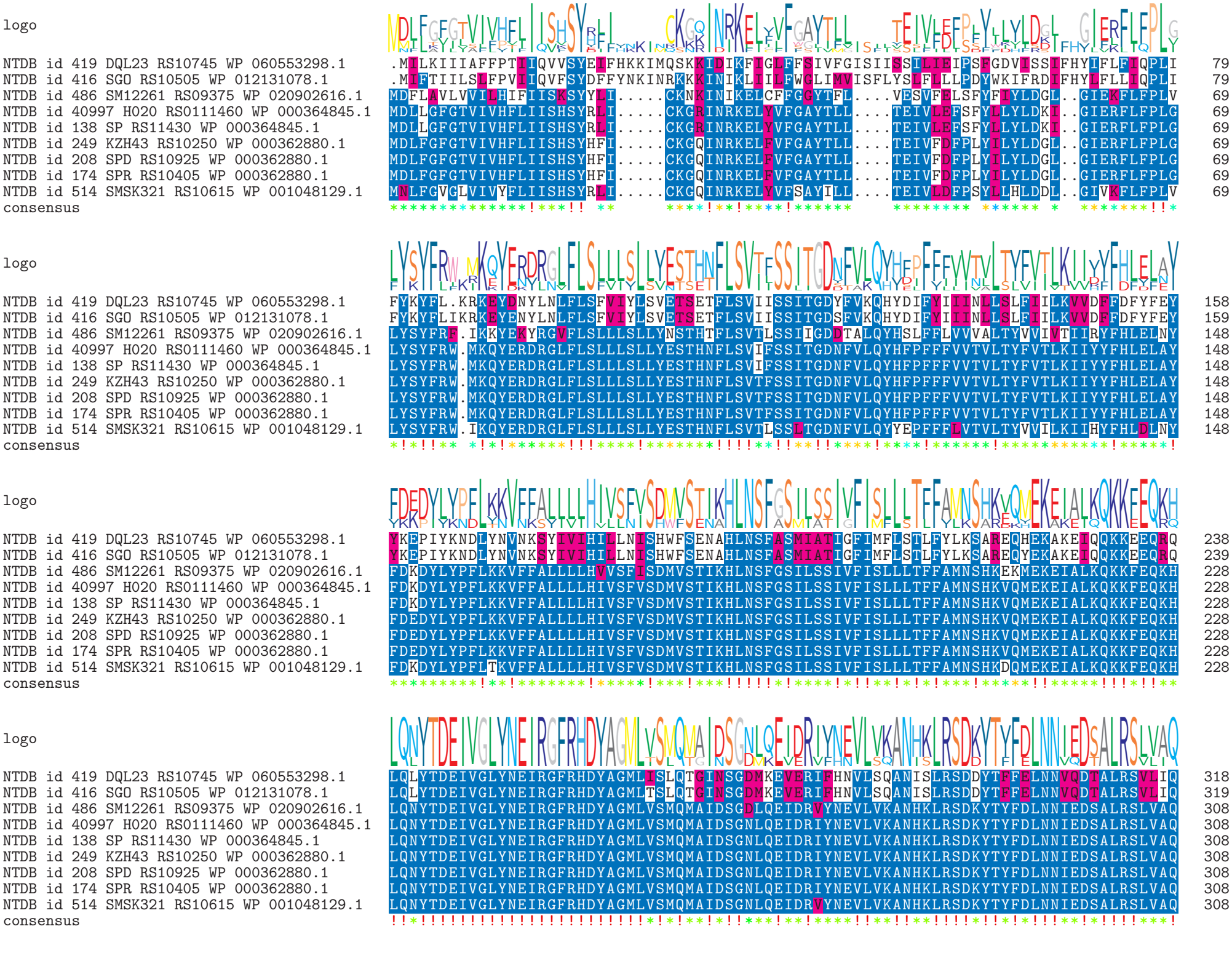
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NTDB id 416 SGO RS10505 WP 012131078.1 .MIFTIILSLFPVIIQVFSYDFFYNKINRKKKININKLIILFWGLIMVISFLYSLFLLLPDYWKIFRDIFHYLFLLIQPLI 79
NTDB id 486 SM12261 RS09375 WP 020902616.1 MDFLAVLVVILHIFIISKSYYLI.....CKNKINIKELCFFGGYTFL....VESVFELSFYFIYLDGL..GIEKFLFPLV 69
NTDB id 40997 HO20 RS0111460 WP 000364845.1 MDLLGFGTVIVHFLIISHSYRLI.....CKGRINRKELYVFGAYTLL....TEIVLEFSFYLLYLDKI..GIERFLFPLG 69
NTDB id 138 SP RS11430 WP 000364845.1 MDLLGFGTVIVHFLIISHSYRLI.....CKGRINRKELYVFGAYTLL....TEIVLEFSFYLLYLDKI..GIERFLFPLG 69
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NTDB id 208 SPD RS10925 WP 000362880.1 MDLFGFGTVIVHFLIISHSYHFI.....CKGQINRKELFVFGAYTLL....TEIVFDFPLYILYLDGL..GIERFLFPLG 69
NTDB id 174 SPR RS10405 WP 000362880.1 MDLFGFGTVIVHFLIISHSYHFI.....CKGQINRKELFVFGAYTLL....TEIVFDFPLYILYLDGL..GIERFLFPLG 69
NTDB id 514 SMSK321 RS10615 WP 001048129.1 MNLFGVGLVIVYFLIISHSYRLI.....CKGQINRKELYVFSAYILL....TEIVLDFPSYLLHLDDL..GIVKFLFPLV 69
consensus
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NTDB id 208 SPD RS10925 WP 000362880.1 LYSYFRW.MKQYERDRGLFLSLLLSLLYESTHNFLSVTFSSITGDNFVLQYHFPFFFVVTVLTYFVTLKIIYYFHLELAY 148
NTDB id 174 SPR RS10405 WP 000362880.1 LYSYFRW.MKQYERDRGLFLSLLLSLLYESTHNFLSVTFSSITGDNFVLQYHFPFFFVVTVLTYFVTLKIIYYFHLELAY 148
NTDB id 514 SMSK321 RS10615 WP 001048129.1 LYSYFRW.IKQYERDRGLFLSLLLSLLYESTHNFLSVTLSSLTGDNFVLQYYEPFFFLVTVLTYVVILKIIHYFHLDLNY 148
consensus
logo
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consensus
logo
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consensus

logo

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consensus                                    *!**!!**!!!***!*!!*!**!!**!!!!!!**!!!!!!!!!!!*!*!**!*!*!***!***!*!*!!!!!*********
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logo

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NTDB id 419 DQL23 RS10745 WP 060553298.1     EIYEVGFSTKGENRGLGLNNVKEIIDKYDEVILETDIETNYFIQVVRFKRKEKI   452
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NTDB id 514 SMSK321 RS10615 WP 001048129.1   DLFALGFSTKGRNRGVGLNNVKELLDKYNNIILETEMEGSTFRQIIRFKREFE.   441
consensus                                    *****!!!!!!*!!!*!!!!!!!**!!!***!!!!**!***!*!**!!!!***
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X non conserved
X similar
X ≥ 50% conserved