

## Supplementary tables

**Supplementary table 1.** Distribution of restriction-modification proteins in Mollicutes. If at least one protein of particular type is encoded within the genome it is marked with 1, F indicates protein fusion. Columns: C – controller subunit (transcription factor), M I – methylase of type 1 RM-system, S I – specificity subunit of type 1 RM-system, R I – restrictase subunit of type 1 RM-system, M II - methylase of type 2 RM-system; Sequence – putative controller subunit binding sites upstream of respective RM-operon.

Species	C	M I	S I	R I	M II	Sequence
<i>A. axantum</i>	0	0	0	0	1	
<i>A. brassicae</i>	0	1	1	1	1	
<i>A. equifetale</i>	1	1	1	1	0	CAATTT <u>AACGATT</u> TGCTACTTTG <u>AACGATT</u> TATTCAACAAATG <u>AACGATT</u> AGGGG
<i>A. hippikon</i>	1	1	1	1	0	TATTT <u>AACGATT</u> ACCATACTTTT <u>AACGATT</u> TGACAACAATT <u>AACGATT</u> TGAGA
<i>A. granularum</i>	1	1	1	1	0	TTTTTATTTGTTAGTCACACTTT <u>AACGATT</u> ACCTTACTTTT <u>AACGATT</u> TGAGA
<i>A. laidlawii</i>	0	1	1	1	1	
<i>A. modicum</i>	1	1	1	1	0	TTTG <u>AACGTT</u> AGGGCTAGTTTG <u>AACGATT</u> TACTCAACTTTG <u>AACGATT</u> TGAGA
<i>A. oculi</i>	0	1	1	0	1	
<i>A. palmae</i>	0	1	1	0	1	
<i>M. agalictiae</i>	0	1	1	1	1	
<i>M. arginini</i>	1	1	1	1	0	ATAGCAGAAA <u>GTGCTA</u> TAATA <u>GTGTAT</u> ATATAATCA
<i>M. arthritidis</i>	0	1	0	0	0	
<i>M. bovis</i>	0	1	1	1	0	
<i>M. callifornicum</i>	F*	0	0	0	F*	TGTCAAGTAGAG <u>GGACATAAA</u> <u>GTCC</u> TTATATCAGCT
<i>M. capricolum</i>	0	0	0	0	1	
<i>M. conjunctivae</i>	1	1	1	0	1	CTCAAGTTTT <u>TTGTTA</u> TAATT <u>AAACAA</u> GATAAAAAGC
<i>M. crocodyli</i>	0	1	1	1	1	
<i>M. fermentans</i>	1	1	1	1	0	AAAAAAATAAA <u>ATGTTA</u> TAATT <u>TTGTTA</u> TAAGTTGTTAG
<i>M. gallinarum</i>	1	0	0	0	1	CGACTAAAAA <u>GTGCTA</u> AAAAT <u>GTGATA</u> TAATTGTGGC
<i>M. gallisepticum</i>	1	1	1	1	0	TATCGGCTTT <u>GTGTTA</u> AAATA <u>GTGTTA</u> ACGATTTGA
<i>M. genitalium</i>	0	0	1	0	1	
<i>M. hominis</i>	1	0	0	0	1	AATCGACAAA <u>GTGATA</u> GTTTT <u>GTGATA</u> TAGTTAAGAT
<i>M. hyopneumonia</i>	0	0	0	0	1	
<i>M. hyosynoviae</i>	1	0	0	0	1	TATAGAGTAA <u>ATGTTA</u> AAAAA <u>ATGATA</u> TAATTTTGTC
<i>M. imitans</i>	0	1	1	0	1	
<i>M. mobile</i>	F*	1	1	1	F*	TTGTATATTT <u>TAACAC</u> ACAA <u>ATGTTA</u> TAATGTAATT
<i>M. mycoides</i>	0	0	0	0	1	
<i>M. penetrans</i>	0	1	1	1	1	
<i>M. pneumonia</i>	0	1	1	1	1	
<i>M. pulmonis</i>	0	1	1	1	1	
<i>M. simbae</i>	F*	0	0	0	F*	TGTTAATAAGC <u>GGACAAATA</u> <u>GTCC</u> TAATTAATAAA
<i>M. synoviae</i>	0	0	0	0	1	
<i>U. canigenitalium</i>	F*	1	0	0	F*	TAGTTATAAT <u>ATGTC</u> ATTTA <u>GTGTC</u> ATATGAATT
<i>U. urealiticum</i>	0	1	1	1	0	

\* - Fusion of controller subunit with methylase subunit.