

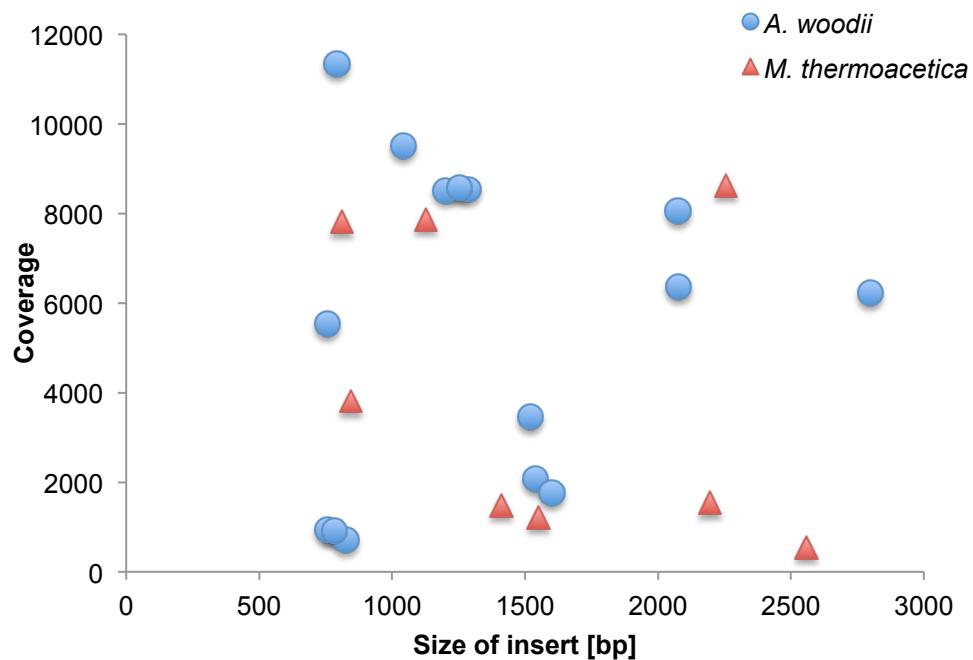
Supplementary Information

Genome-wide systematic identification of methyltransferase recognition and modification patterns

Jensen et al.

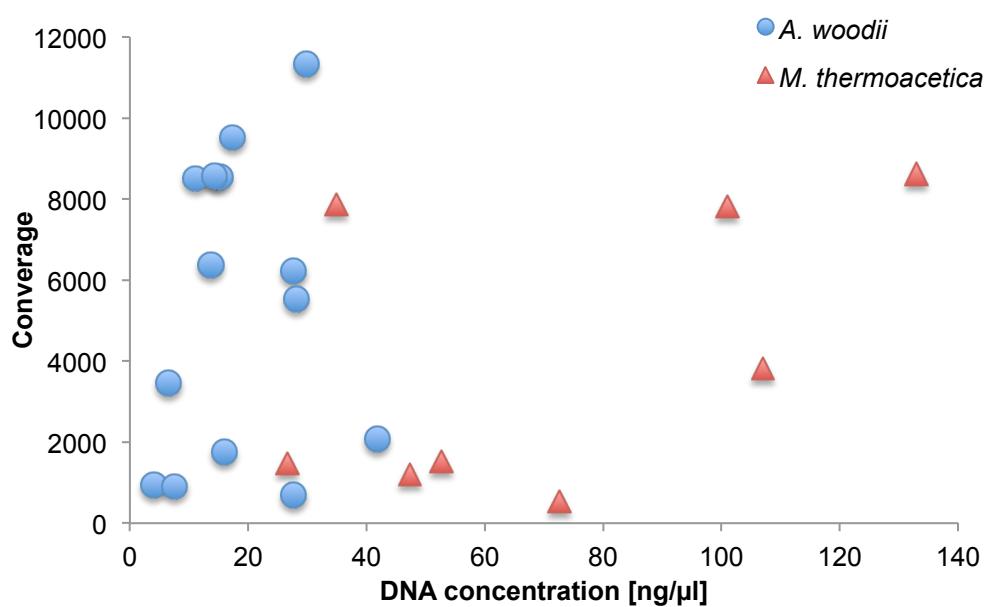
Supplementary Figure 1

Graph showing the coverage (generated at std. parameters) as a function of the size of the insert (methyltransferase) in basepairs (bp).



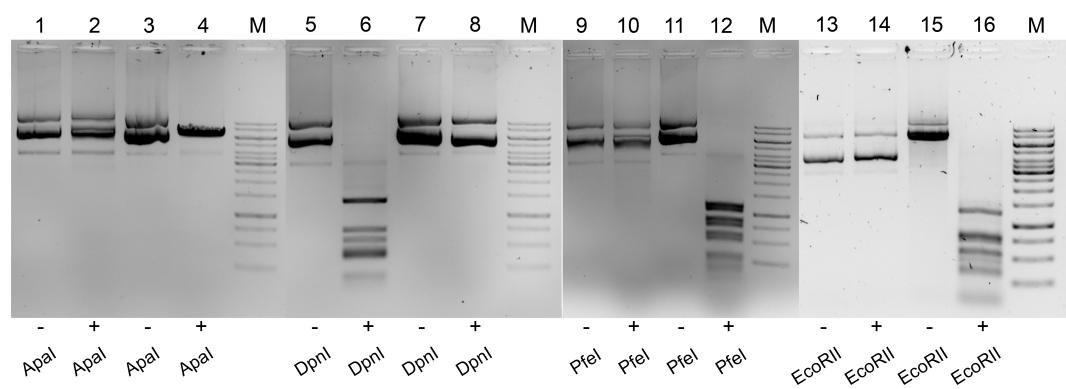
Supplementary Figure 2

Graph showing the coverage (generated at std. parameters) as a function of the DNA concentration (plasmid) in ng/ μ l coverage.



Supplementary Figure 3

Modification verification by restriction enzymes. Plasmid backbone is pRSFduetPA1lacO-1. Lanes marked with M are GeneRuler™ 1 kb DNA ladder (Thermo Scientific). Lanes 1 and 2 are MothHH_02467, lanes 5 and 6 are MothHH_01869, lanes 9 and 10 are MothHH_00029, and lanes 13 and 14 are Awo_c18590. Lanes 3,4,7,8,11,12,15 and 16 are the negative control *bgaB*.



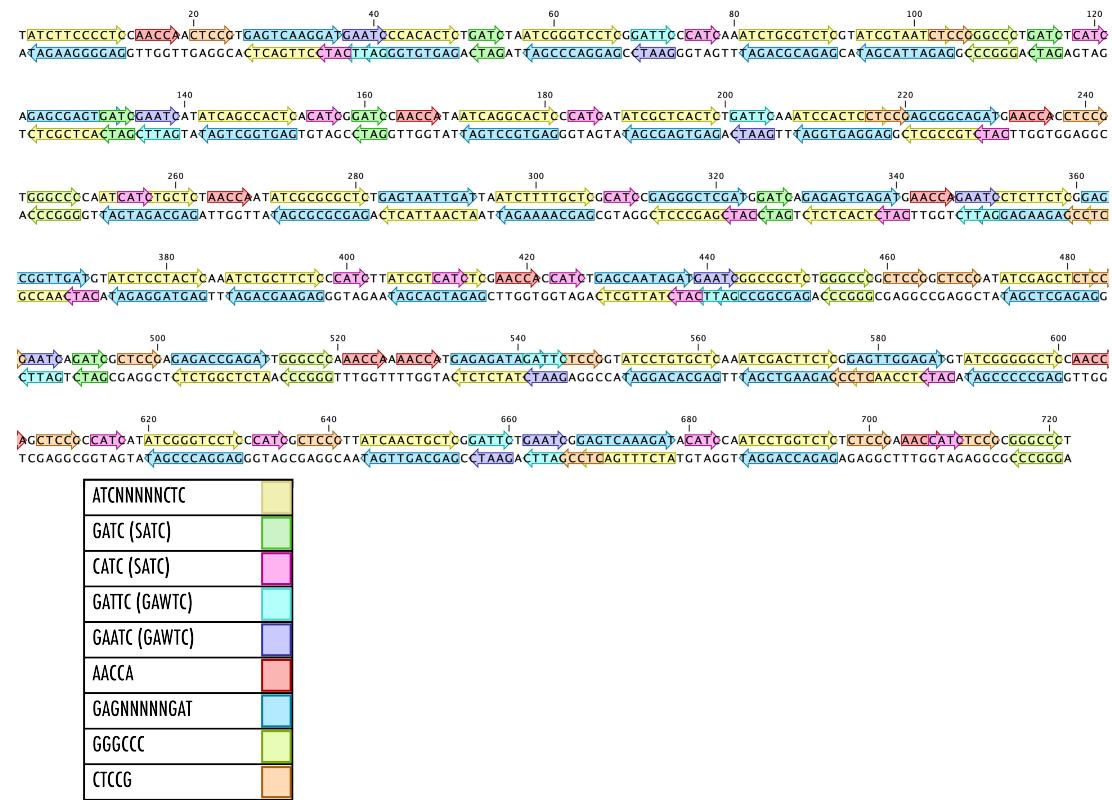
Supplementary Table 1

Numbers of restriction digestion sites on the plasmids tested by restriction digestion.

pRSFduetPA1lacO-1 with the following genes inserted	Size (bp)	Apal GGGCCC	DpnI G ^{m6} ATC	PfeI GAWTC	EcoRII CCWGG
MothHH_02467	6474	6			
MothHH_01869	5031		14		
MothHH_00029	5346			21	
Awo_c18590	5504				27
<i>bgaB</i> (negative cont.)	6302	1	14	12	14

Supplementary Figure 4

Sequence of the motif-cassette with motifs for assessing the methyltransferases of *M. thermoacetica*.



Raw sequence

TATCTTCCCCCTCCAACCAACTCCGTGAGTCAGGATGAATCCCACACTCTGATCTAAT
CGGGTCCTCGGATTCCCATCAAATCTGCCTCGTATCGTAATCTCCGGGCCCTGATC
TCATCAGAGCGAGTGATCGAACATCATATCAGCCACTCACATCGGATCCAACCATAATC
AGGCACCTCCCATCATATCGCTACTCTGATTCAAATCCACTCCTCCGAGCGGCAGATG
AACCACCTCCGTGGGCCCCAATCATCTGCTCTAACCAATATCGCGCCTCTGAGTAAT
TGATTAATCTTTGCTCGCATCCGAGGGCTCGATGGATCAGAGAGTGAGATGAACCA
GAATCCTCTTCTCGGAGCGGTTGATGTATCTCCTACTCAAATCTGCTTCTCCATCTT
ATCGTCATCTCGAACCCACCATCTGAGCAATAGATGAATCGGCCGCTCTGGGCCGCTC
CGGCTCCGATATCGAGCTCTCCGAATCAGATCGCTCCGAGAGACCGAGATTGGGCCA
AACCAAAACCATGAGAGATAGATTCTCCGGTATCCTGTGCTCAAATCGACTTCTCGG
AGTTGGAGATGTATCGGGGGCTCCAACCAAGCTCCGCCATCATATCGGGTCTCCCATC
GCTCCGTTATCAACTGCTCGGATTCTGAATCGGAGTCAAAGATACTCCAATCCTGGT
CTCTCTCCGAAACCATCTCCGCGGGCCCT -3'

Supplementary Figure 5

Sequence of the motif cassette for assessing the methyltransferases of *A. woodii*.



GCCAAG (GCCRAG)	
GCGAG (GCCRAG)	
CCAGG (CCWGG)	
CCTGG (CCWGG)	
TAAGNNNNNTCC	
GATGNNNNNNTGC	
CAAAAAA (CAAAAAR)	
CAAAAG (CAAAAAR)	

Raw sequence

TAAGTCCTGTCCAGCCAAGCGCCGAGGCAAAAAATAACCAGCACGAATCCTGCTGGT
 CCTCACCCCCAAAAAGGGATGAAACATTGCCTAACGATATCTCCGCCTGGACGCCAAGC
 GCCGAGCCCTGGTAAGTCTTCCAGATGGCACGTGACAAAAAAAGCCAAGGGA
 TGACAATATGCCCAAAAAGGGCCGAGGCCAGGTTAACGGCTATCCTCAAAAGGGCC
 GAGGCCAGGCTAACGGCTGTCCAGCCAAGTTAACGTTGGGTCCACCAGGTAAAAAAA
 GCCAAGCGCCGAGACAAAAAGGCCAGACCAGGCTAACGCTAGTCCAGCCGAGAT
 AAGGAAAGTCCGGCCGAGTCAAAAAATAAGAGATCTCCGCCTGGTGCCAGGCCAAA
 AAGTACAGTTTCAGGCCTTCCTTCAGGTCCAGTCCCTGCAAGATGCACGAATGCAGC
 CAAGTCCAGGCCAAAAAGGCCAGAGAGATGTCTGGTGCCTCCAAAAAGACCTGGGAT
 GCCCAGTTGCCAGGCCAAAAAGGCCAAAAACCTGGAGGCCAGAGATGTTGGT
 TGCCCTGGTGTAGAGGAATGCACCTGGTGCCTGAGCTAACGTCAGGTCCACCAGGC
 ATGAGACCCTGCGCAAAAAGGCCAGGCAAGCCAAAAAGGCCAGGTCAAAAAAA
 ATTAAGATGCCGTCCAGATGCAGATGTGCTCAAAAAGCGATGTCCGCTGCCTAACAG
 GTGTCCAGCCAAGGGATGCGCGGCTGCCAAAAAAACAAAAAGCTCACACTGCTTCC
 GGTAG -3'

Supplementary Table 2Media composition for cultivating *M. thermoacetica*, according Daniel et al.⁶².

Components	Concentration
AlK(SO ₄) ₂ ·12H ₂ O	50 µg/l
Biotin	0.1 mg/l
CaCl ₂ ·2H ₂ O	50 mg/l
Calcium D-(+)-panthothenate	0.25 mg/l
Co(NO ₃) ₂ · 6H ₂ O	0.5 mg/l
CuSO ₄ ·5H ₂ O	0.05 mg/l
Cysteine-HCl	158 mg/l
FeSO ₄ ·7H ₂ O	0.5 mg/l
Folic acid*	0.1 mg/l
Fructose	5 g/l
H ₃ BO ₃	0.05 mg/l
KH ₂ PO ₄	500 mg/l
Lipoic acid	0.15 mg/l
MgCl ₂	330 mg/l
MnSO ₄ ·H ₂ O	2.5mg/l
Na ₂ MoO ₄ ·2H ₂ O	0.05 mg/l
NaCl	400 mg
NaHCO ₃	3.5 g/l
NH ₄ Cl	400 mg/l
NiCl ₂ ·6H ₂ O	0.25 mg/l
Nicotinic acid	0.25 mg/l
Nitrilotriacetate	7.5 mg/l
p-Aminobenzoic acid	0.25 mg
Pyridoxine hydrochloride	0.05 mg/l
Resazurin	1 mg/l
Riboflavin	0.25 mg/l
Thiamine HCL	0.25 mg/l
Vitamin B12 (Cyanocobalamin)	0.25 mg/l
Yeast extract	1 g/l
ZnCl ₂	0.5 mg/l