

**Table S3.** Summary of putative transcription site mapping.

Strain	Gene	Putative TS*	-35	-10
<i>M. trichosporium</i> OB3b	<i>pmoC</i>	-273	<b>TTGTCA</b>	A <b>AT</b> GGTTG
	<i>pmoC</i>	-324	<b>TTGTGC</b>	TCCTACCC TA
	<i>mxoF</i>	-156	<b>CAGACA</b>	<b>TATATG</b>
	<i>faeI</i>	-214	<b>TTCGAT</b>	<b>TATAAT</b>
	<i>pqqA</i>	-90	<b>TTGCAC</b>	G <b>CTAAT</b>
<i>Methylocystis</i> sp. M <sup>1</sup>	<i>pmoC</i>		<b>TTGTCA</b>	GA <b>AT</b> GGTT G
<i>M. extorquens</i> AM1 <sup>2</sup>	<i>mxoF</i>		<b>AAGACA</b>	<b>TAGAAA</b>
	<i>mxoW</i>		<b>TTGGCA</b>	ACCCAT
<i>E. coli</i> <sup>3</sup>	s70		<b>TTGACA</b>	<b>TATAAT</b>
	s32		TCTCNCCCTTGAA	
	s54		CTGGNA	TTGCA
	s28		CTAAA	CCGATAT

1 Holmes *et al.*, 1995 (see Ref)

2. Zhang, M., and Lidstrom, M.E. (2003) Microbiol 149:1033-40.

3. Harley, C.B., and Reynolds, R.P. 1987, Nucleic Acids Res. 11:2343-61