

# DNA METHYLATION IN *E. COLI*: EFFECT ON RESTRICTION ENZYMES

**1. The problem.** Many restriction enzymes are blocked (fully or partially) by methylation. A few enzymes require methylation (such as *DpnI*)

**2. Three methylation systems.** Most laboratory strains of *E. coli* contain three site-specific DNA methylases that can block digestion with restriction enzymes. Avoid designing restriction sites in primers or genes that may become methylated. Problems usually occur due to bases flanking the restriction site.

Dam methylase: methylates A in GATC

Dcm methylase: methylates C in CCAGG and CCTGG

EcoKI methylase: methylates A in AAC(N<sub>6</sub>)GTGC and AAC(N<sub>6</sub>)GTGC

**3. Sources of information.** Information about the sensitivity of restriction enzymes to methylation can be found on the internet, including at <http://rebase.neb.com>. The following table from NEB shows information about Dam and Dcm sensitivity.

Dam Methylation: G <sup>m</sup> ATC				Dcm Methylation: C <sup>m</sup> CWGG			
<b>Blocked by Overlapping Dam</b>				<b>Blocked by Overlapping Dcm</b>			
AlwI		GGATC		Acc65I <sup>2</sup>		GGTACCWGG	
BcgI <sup>1</sup>		CGATCNNNTGC		AlwNI		CAGNNCCTGG	
BclI		TGATCA		ApaI		GGGCCCWGG	
BsaBI		GATCNNNATC		AvaII		GGWCCWGG	
BspDI		ATCGATC		BanI		GGYRCWGG	
BspEI		TCCGGATC		BsaI		GAGACCWGG	
BspHI		TCATGATC		BsaHI <sup>2</sup>		GRCGCWGG	
ClaI		ATCGATC		BslI <sup>2</sup>		CCWGGNNNNGG	
DpnII		GATC		BsmFI		GGGACT	
HphI		GGTGATC		BssKI		CCWGG	
Hpy188I		TCNGATC		BstXI		CCAGGNNNTGG	
Hpy188III		TCNNGATC		EaeI		YGGCCAGG	
MboI		GATC		EcoO109I		RGGNCCTGG	
MboII		GAAGATC		MscI		TGGCCAAGG	
NruI		TCGCGATC		NlaIV		GGNNCCWGG	
TaqI		TCGATC		PfiMI		CCAAGNNNTGG	
XbaI		TCTAGATC		PpuMI		RGGWCCTGG	
<b>Not Blocked by Overlapping Dam</b>				<b>Not Blocked by Overlapping Dcm</b>			
AsiSI	BglII	BstYI	Sau3AI	BamHI	BsaJI	DraIII	HphI
BamHI	BsaWI	PvuI		BanII	Bsp1286I	FokI	KasI
				BciVI	BstEII	FseI	KpnI
				BglI	BstNI	HaeIII	NarI
				BpmI			

At rebase.neb.com, you can get specific information about enzymes of interest. This includes information, including EcoKI methylase effects. For example:

<u>SpeI</u> ACTAGT MS	<u>Dam</u>	<u>Dcm</u>	<u>CpG</u>	<u>EcoBI</u>	<u>EcoKI</u>
	m6 G A T C C T A G m6	m5 C C W G G G G W C C m5	m5 C G G C m5	m6 T G A N N N N N N N N T G C T A C T N N N N N N N N A C G A m6	m6 A A C N N N N N N N G T G C T T G N N N N N N C A C G m6
overlaps?	n	n	n	y	y
sensitivity?	-	-	-	blocked	blocked

**4. Cloning strains of *E. coli*.** Most cloning strains are Dam<sup>+</sup>, Dcm<sup>+</sup>, EcoKI<sup>+</sup> (most are derived from *E. coli* K12, which lack EcoB1). Methylation contributes to post-replication DNA repair, that is why we usually work with *E. coli* that has methylases.

The lab has a Dam<sup>-</sup>, Dcm<sup>-</sup> strain (2198) that can be used as needed; strains are also available from NEB (# E4109S, ER2925) and Invitrogen (INV110, derivative of JM110). SURE and XL10 cells are Dam<sup>+</sup>, Dcm<sup>+</sup>, EcoKI<sup>-</sup>. Note that some methylation systems are not 100% efficient, so you might get some digestion from regular *E. coli*.

The genotype of DH5-alpha is: F<sup>-</sup> Φ80lacZΔM15 Δ(lacZYA-argF) U169 recA1 endA1 hsdR17 (rK<sup>-</sup>, mK<sup>+</sup>) phoA supE44 λ<sup>-</sup> thi-1 gyrA96 relA1; the hsdR17 is in the EcoK restriction enzyme (AACNNNNNNGTGC) but the corresponding methylase gene (hsdM) is present.

Those other genotypes are:

ΔlacZYA: deletion of lac operon (the U169 mutation)

F<sup>-</sup> Φ80lacZΔM15: low-copy plasmid, for making single-stranded DNA with M13 phage, carries lacI and lacZΔM15 gene.

recA: mutation in gene for protein that stimulates replication.

endA: mutation in non-specific endonuclease, resulting in improved plasmid preps

gyrA96: resistance to nalidixic acid.

relA: uncouples transcription and translation from amino acid levels.

**5. What about DNA from a PCR reaction?** DNA amplified by PCR is not subject to methylation, and should therefore be digestible, unlike DNA from *E. coli*.

**6. What about cloning methylated DNA into *E. coli*?** There are two systems used to digest cytosine-methylated DNA, the mcrA and mcrB systems. Mammals, higher plants & many prokaryotes contain methylcytosine in their genomic DNA, so their DNA must be cloned into a mcrA<sup>-</sup>, mcrB<sup>-</sup> strain. In most cases, you only need to worry about this when going between species, not between *E. coli* strains.