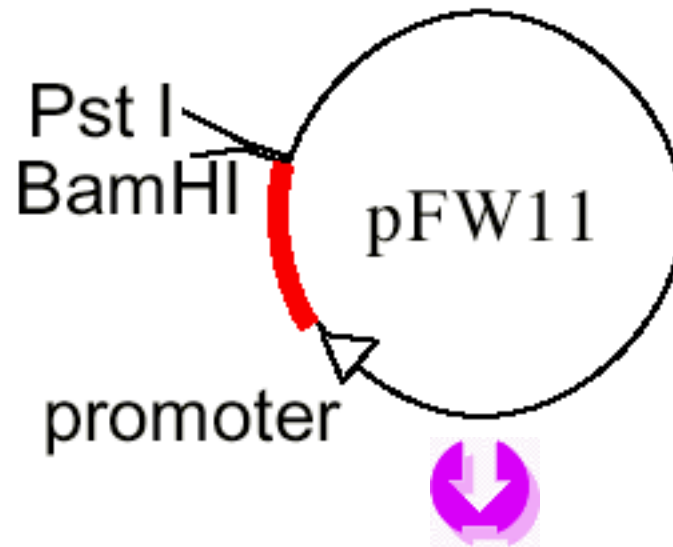
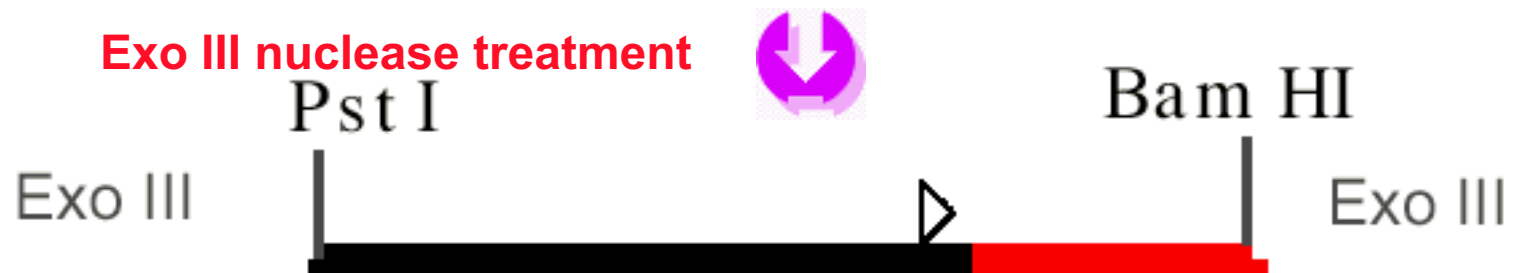


Incremental Truncation



Linearize vector by double digestion



Salt concentration (NaCl)

Exo III – helps stepwise removal of mononucleotides from 3'-hydroxyl termini of duplex DNA

Time-dependent sampling !!

PstI

Recognition Sequence

5'CTGCAG 3'
3'GACGTC 5'

Cut Site

5'--CTGCA G--3'
3'--G ACGTC--5'

BamHI

Recognition

5' -GGATTC-3'
3' -CCTAAG-5'

Cut site

5' -G **GATTC**-3'
3' -CCTAA **G**-5'

The 3' overhangs are resistant to Exo III digestion
whereas the 5' overhangs are SUSCEPTIBLE.

resistant

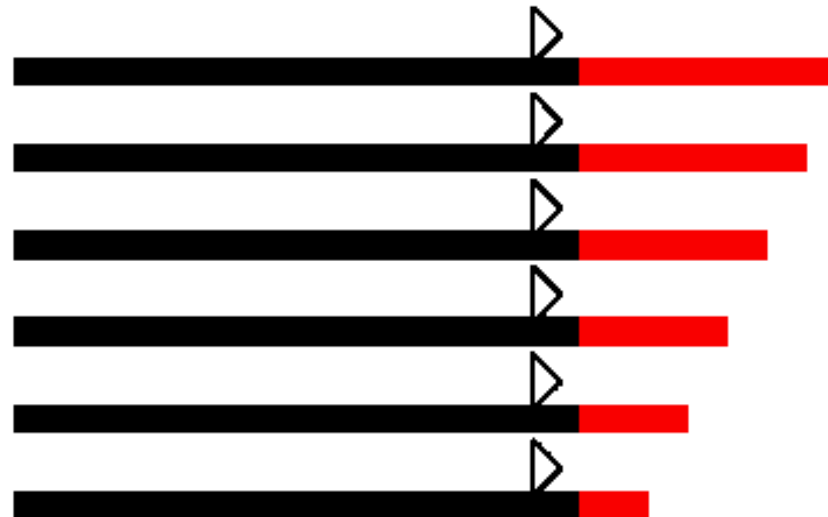


susceptible

Removal of single-stranded
extensions (3' and 5') to leave
ligatable blunt ends



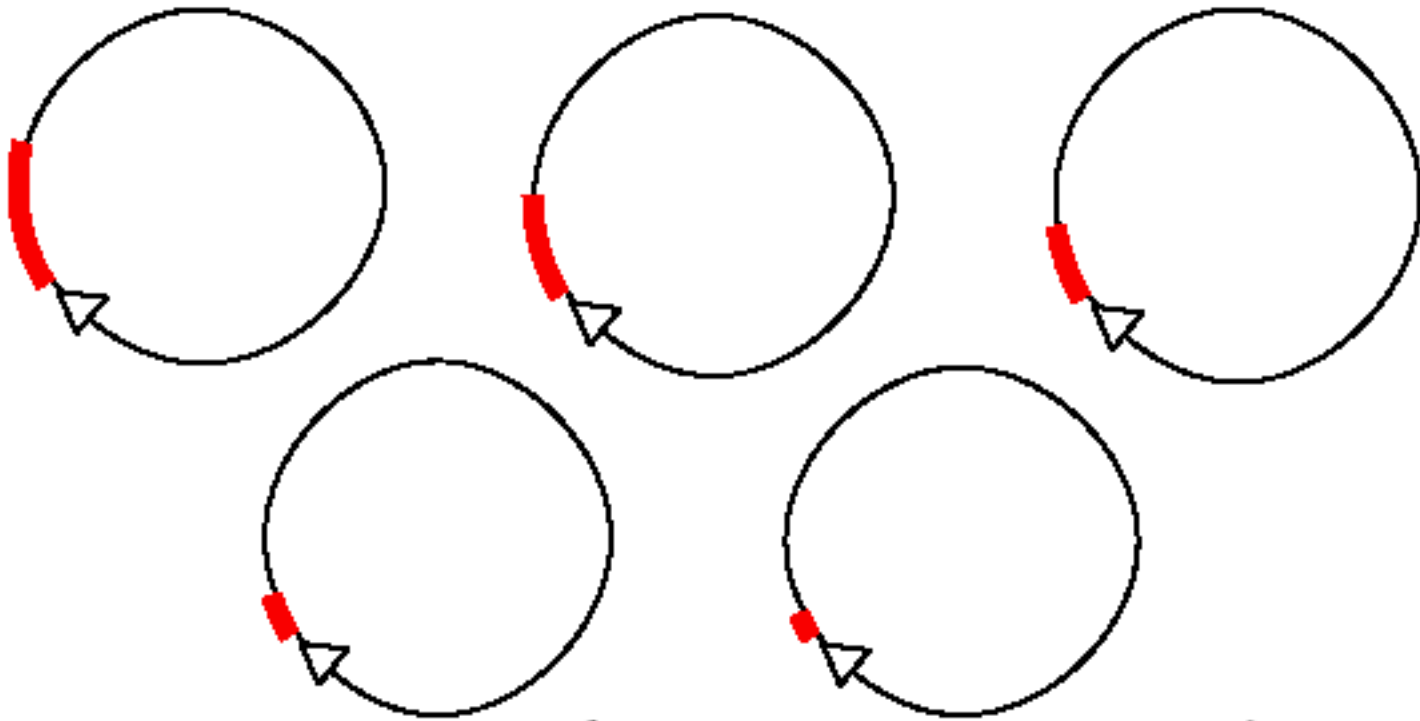
Mung bean nuclease treatment



Klenow treatment

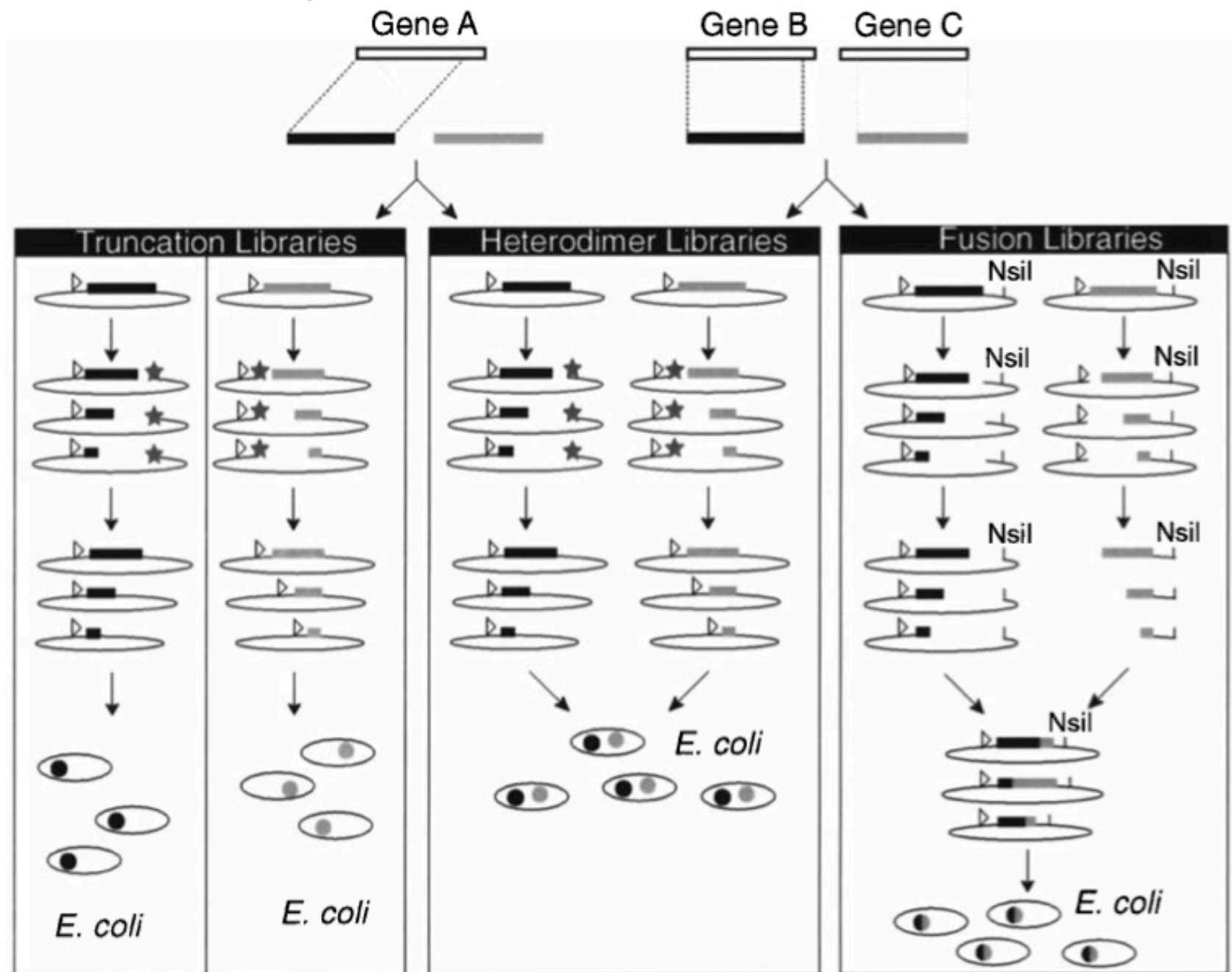
5'-3' polymerase
3'-5'-exonuclease

NO 5'-3' exonuclease



Diverse Library with every one base deletion

Combining Incremental Truncation Libraries



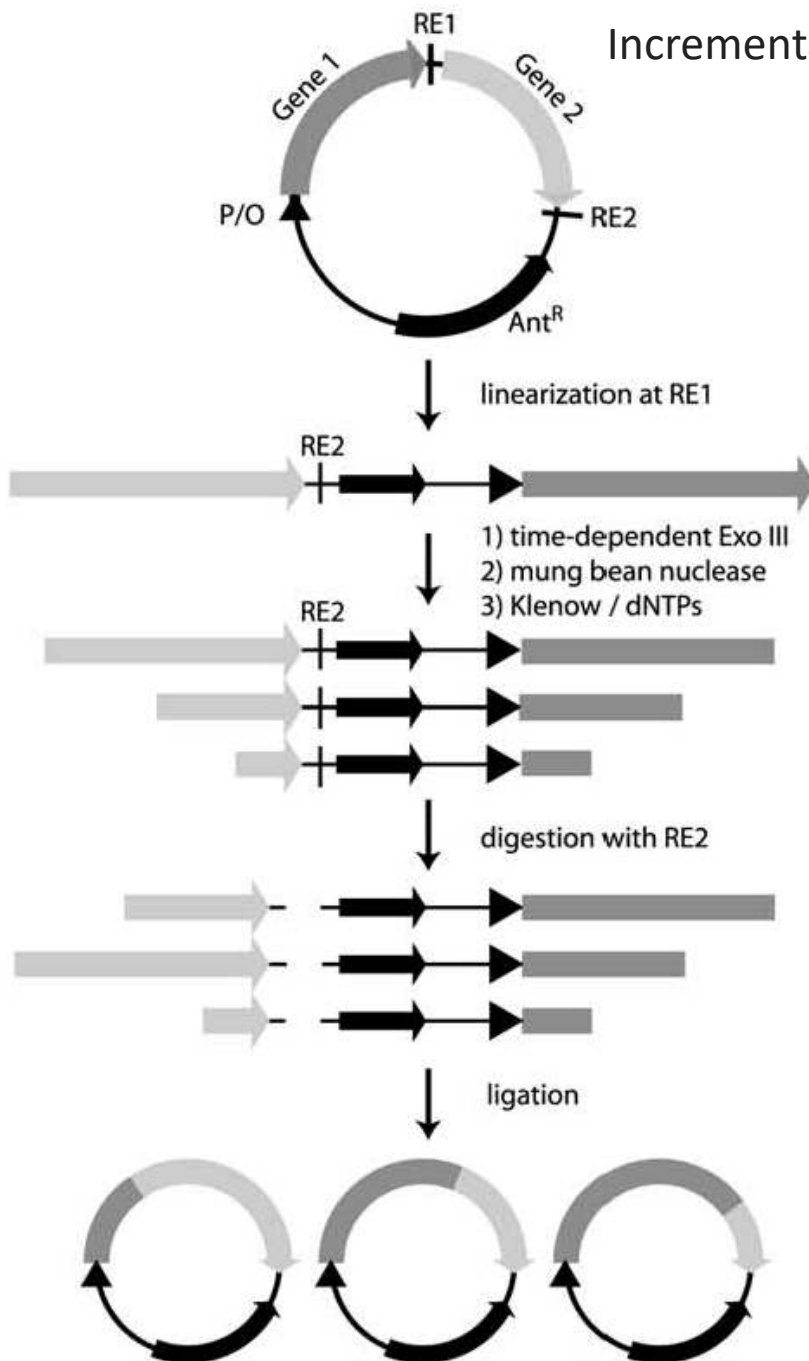
Incremental Truncation for the Creation of Hybrid enzYmes

ITCHY Hybrid Protein Library

The fusion of two incremental truncation libraries is called an ITCHY library.

An ITCHY library created from a **single gene** consists of genes with internal deletions and duplications.

An ITCHY library created between **two different genes** consists of gene fusions created in a DNA-homology independent fashion.



Case study

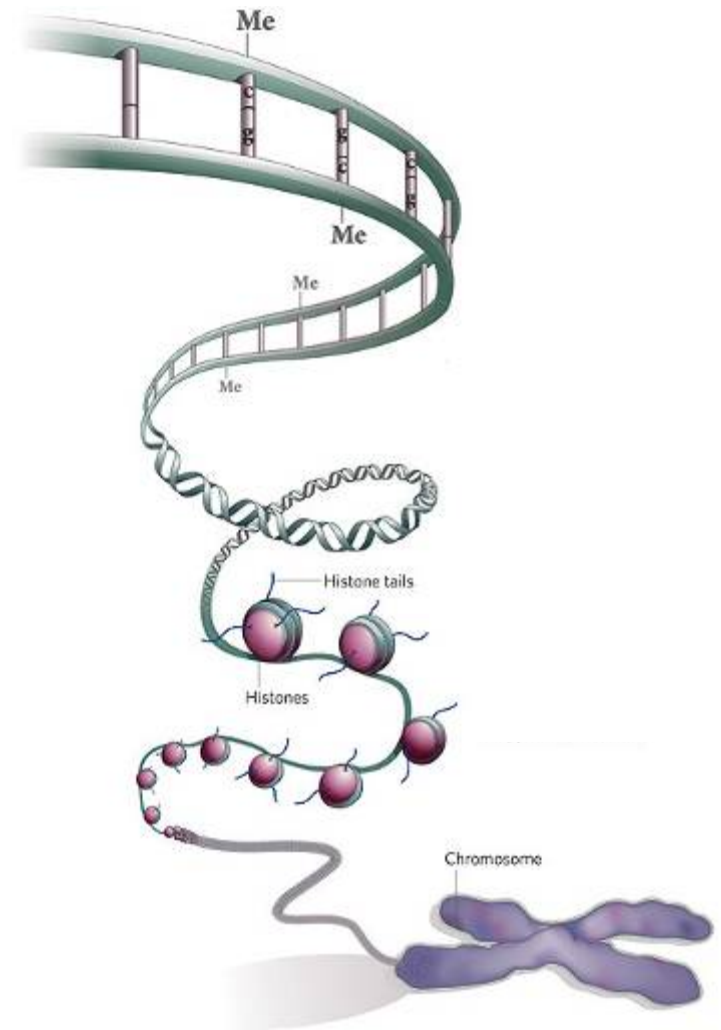
**Non-Associating Heterodimeric
DNA Methyltransferases as a
Platform for Developing
Site-Specific Methyltransferases**

Outline

- DNA methylation background
- Site-biased and site-specific DNA methyltransferases
- Splitting monomeric methyltransferases
- Naturally split methyltransferases

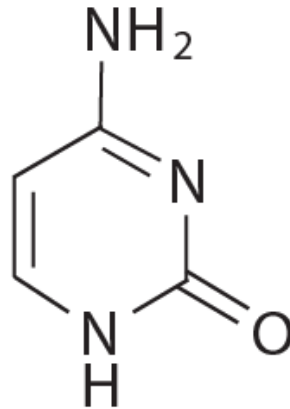
Genome Versus the Epigenome

- Higher eukaryotic cells contain two layers of heritable information
 - **Genome**: Instructions encoded in DNA for making RNA and proteins
 - **Epigenome**: Instructions for how to utilize genetic information
 - DNA methylation
 - Histone modification



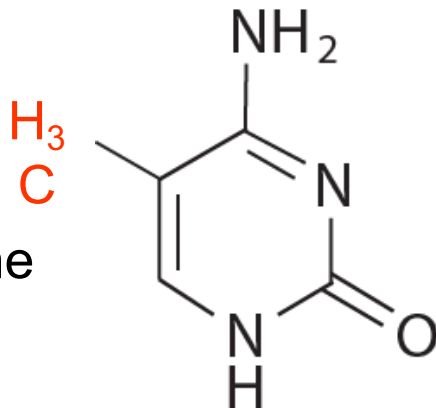
5-Methylcytosine in Mammalian and Plant Cells

cytosine



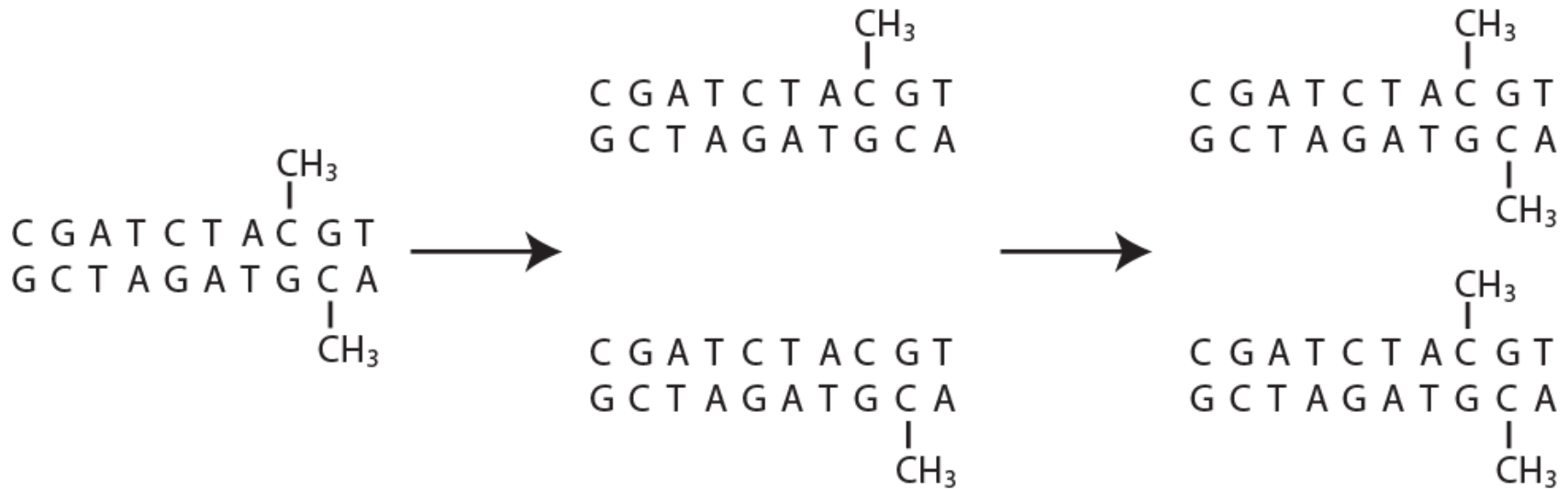
- In **plants**, 5-methylcytosine occurs at **CpG**, **CpHpG** and **CpHpH** sequences (where H = A, C or T).

5-methylcytosine
(5mC)



- In **fungi and animals**, 5-methylcytosine predominantly occurs at **CpG** dinucleotides.

Heritable DNA Methylation



- Enzymes recognize and methylate DNA at hemi-methylated sites

Why is DNA Methylation Important?

- Methylation patterns are important in establishing correct gene expression
 - Embryonic development and cell differentiation
 - DNA imprinting
- Epigenetic diseases characterized by abnormal methylation patterns
 - Cancer
 - Diseases caused by deregulation of imprinted genes