

Your Genes and Your Health

<http://bio84.stanford.edu/>

Epigenetics

<http://bio84.stanford.edu/08%20miRNA%20&%20Epigenetics.htm>

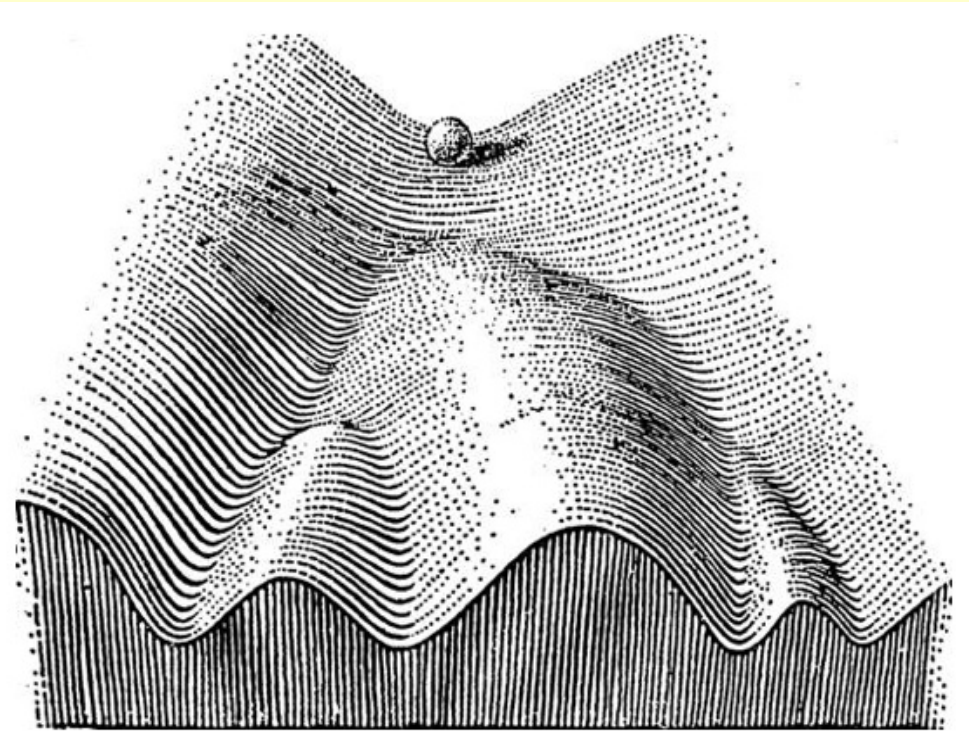


Doug Brutlag, Professor Emeritus
Biochemistry and Medicine (by courtesy)
Stanford University School of Medicine

What is Epigenetics?

- C.H. Waddington coined the term epigenetics to mean above or in addition to genetics to explain differentiation.
- How do different adult stem cells know their fate?
 - Myoblasts can only form muscle cells
 - Keratinocytes only form skin cells
 - Hematopoietic stem cells only become blood cells
 - But all have identical DNA sequences.

C.H. Waddington



Waddington's Epigenetic Landscape

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- Modern definition is non-sequence dependent inheritance.
- How can identical twins have different natural hair colors?

Identical Twins with Different Hair Color



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- How can a single individual have two different eye colors?

An Individual with Two Different Eye Colors



“Diego”

An Individual with Two Different Eye Colors



“Josie Too”

A Woman with Two Different Eye Colors



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- How can identical twin litter mates show different coat colors?

Coat Colors of Genetically Identical Agouti Mice Litter Mates



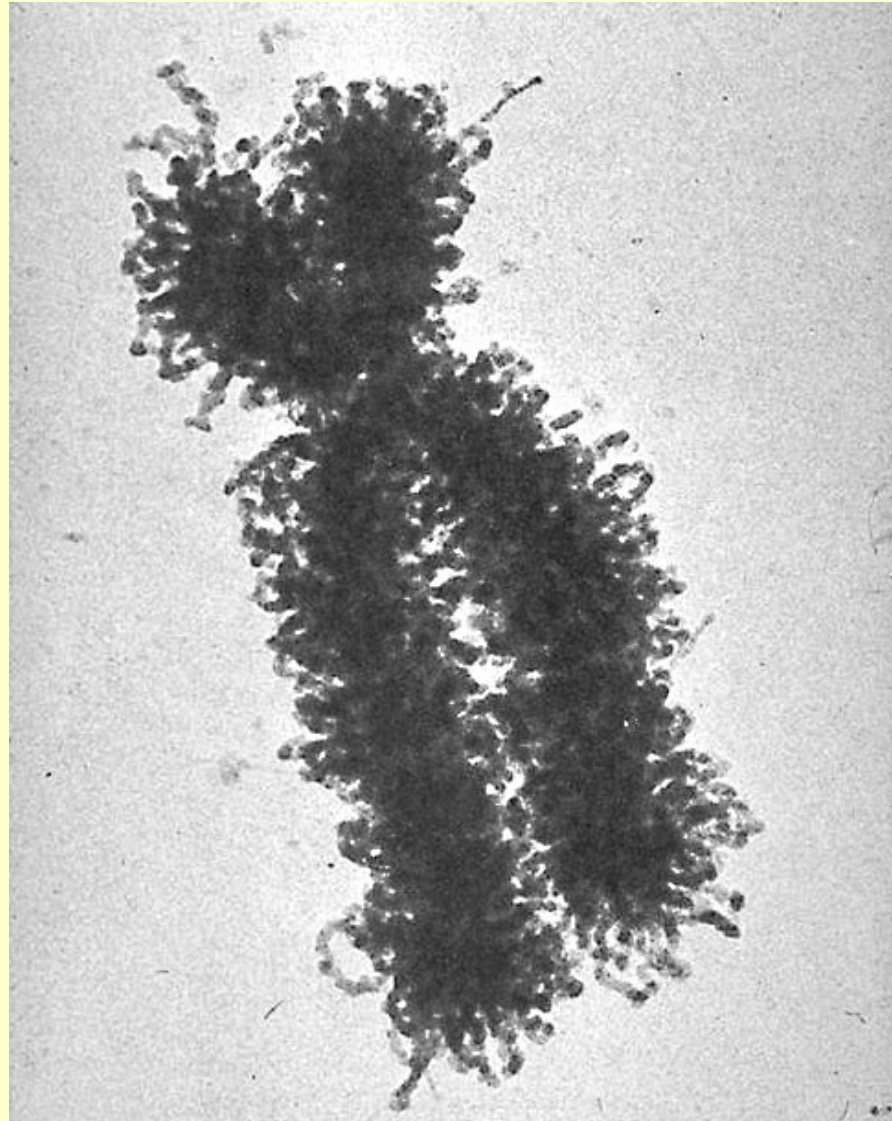
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- How can females express only one X chromosome per cell?
- How can acquired traits be passed on to offspring?
- Some changes in gene expression that are, in fact, heritable!

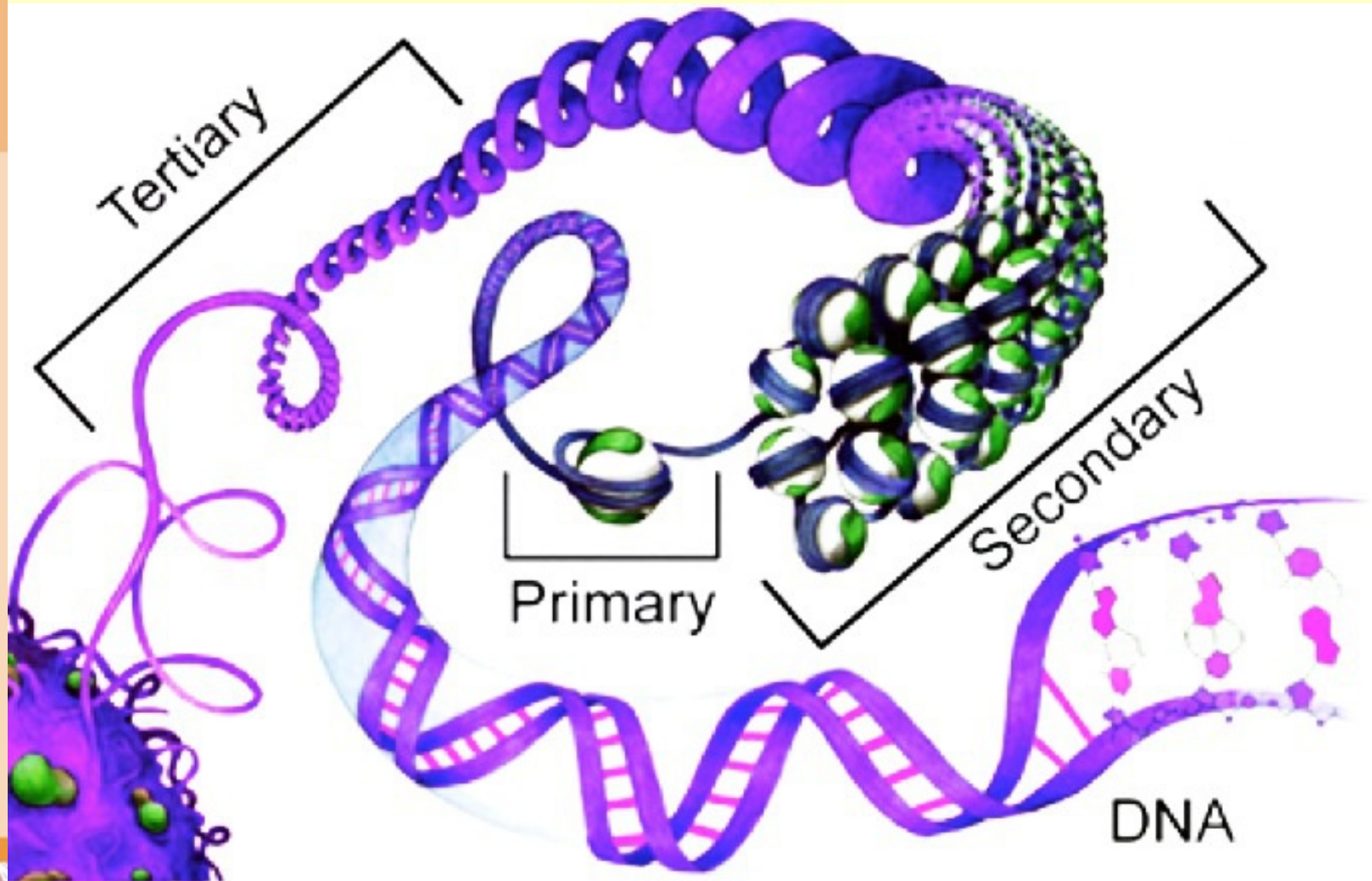
What is Epigenetics?

- Nova Show on Epigenetics
- http://www.teachersdomain.org/asset/biot09_vid_epigenetics/

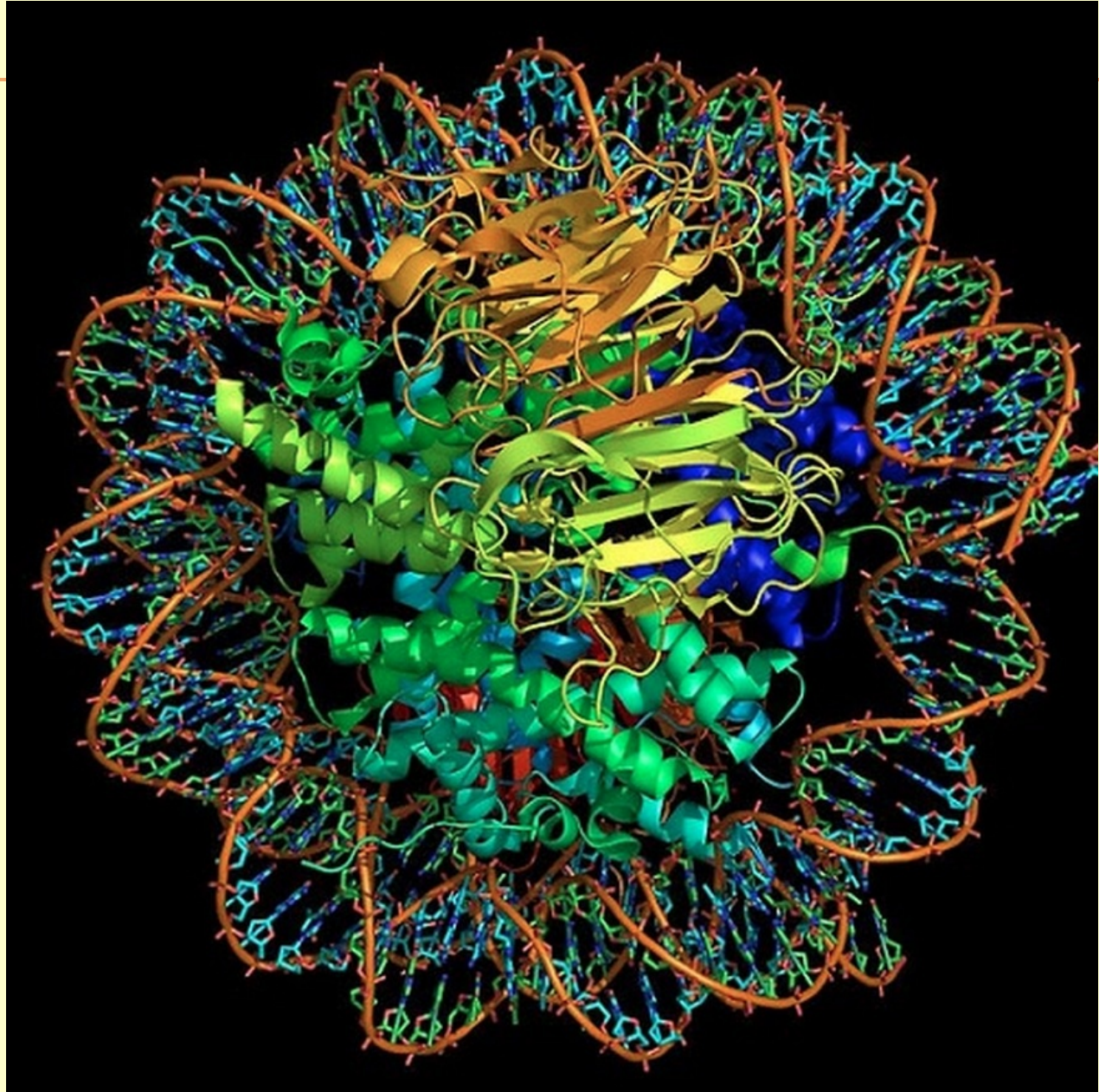
Human Mitotic Chromosome



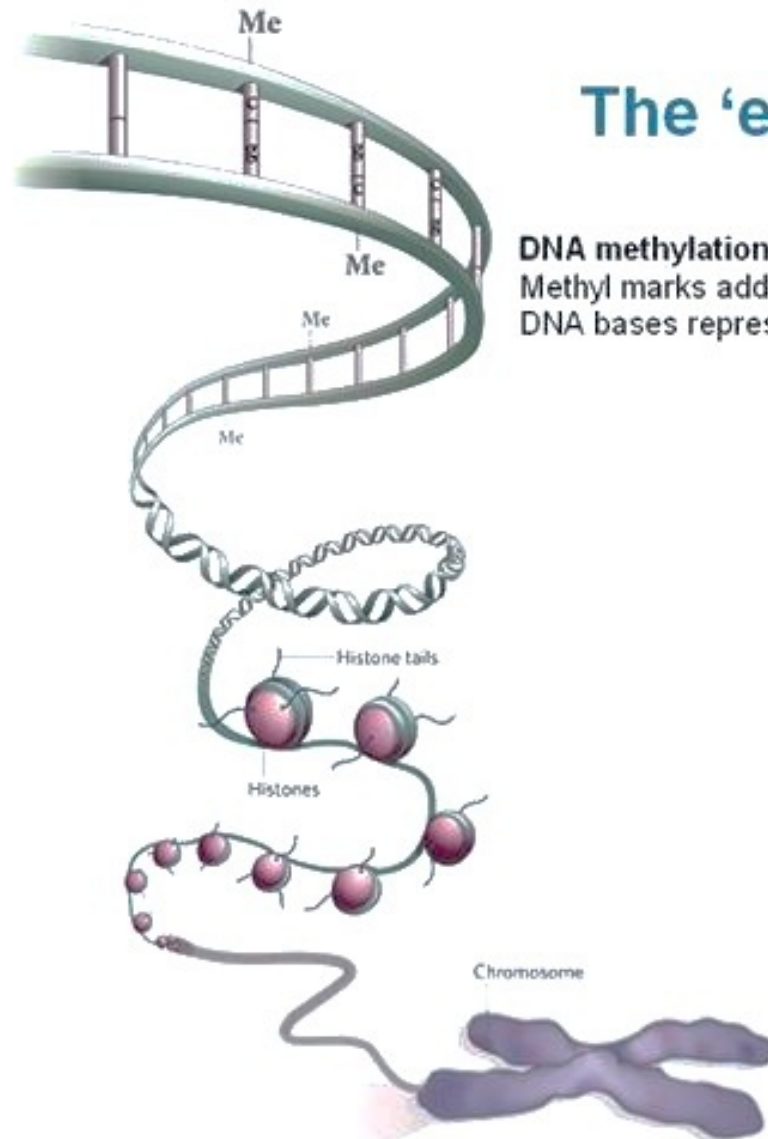
Three Levels of Folding of DNA in Chromatin



Nucleosome Core Structure



DNA Methylation & the Epigenetic Code

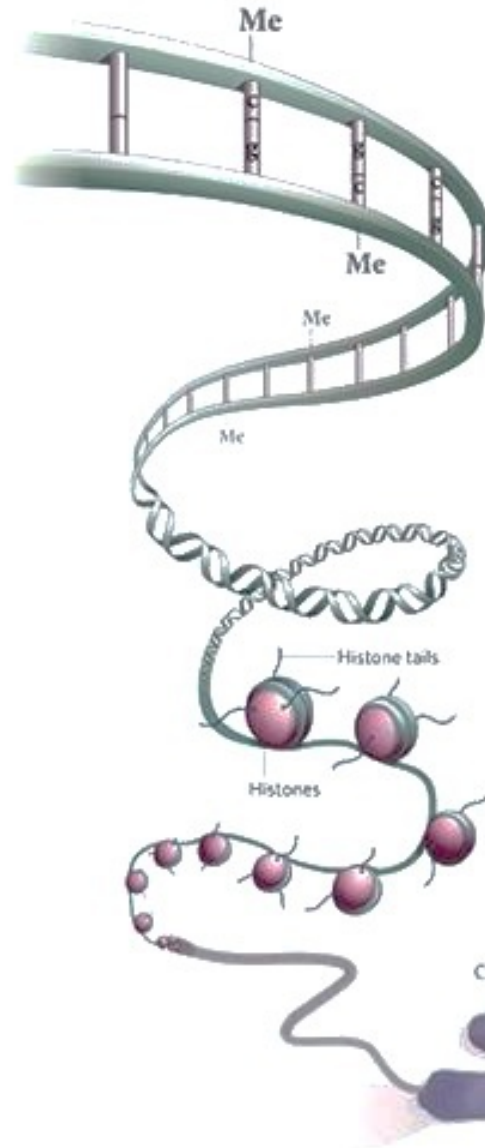


The 'epigenetic' code

DNA methylation
Methyl marks added to certain
DNA bases repress gene activity



DNA Methylation & Histone Modifications Form the Epigenetic Code



The 'epigenetic' code

DNA methylation

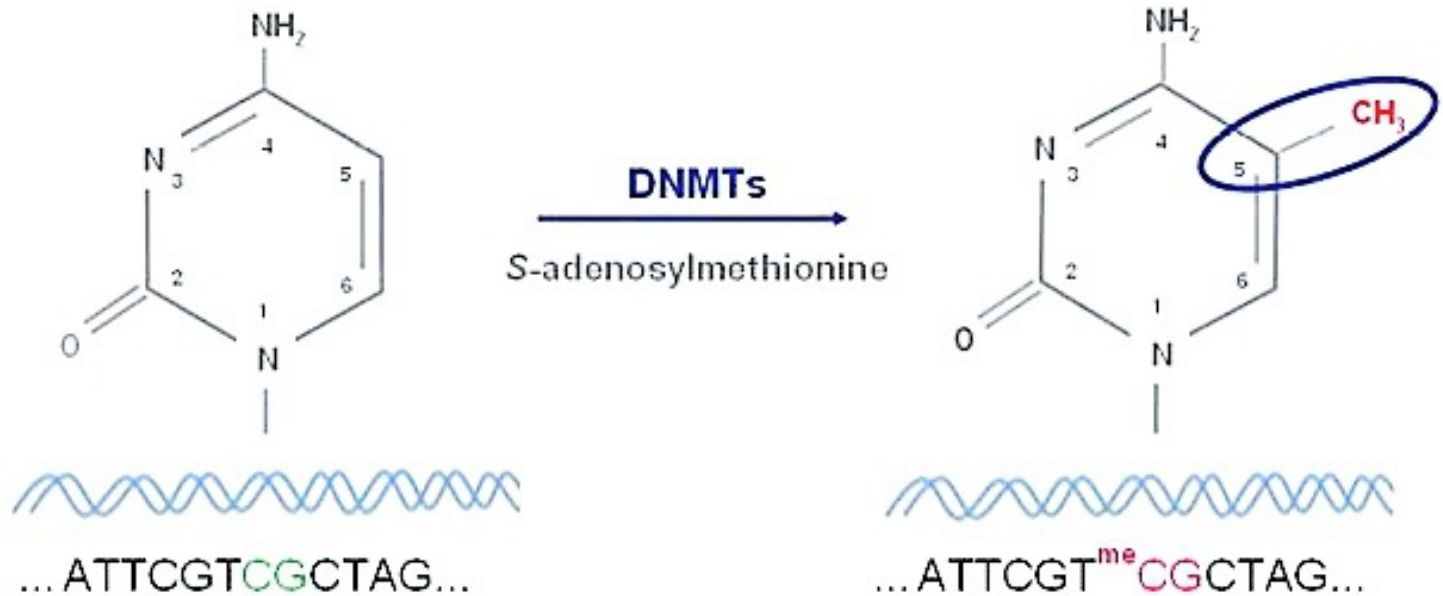
Methyl marks added to certain DNA bases repress gene activity

Histone modification

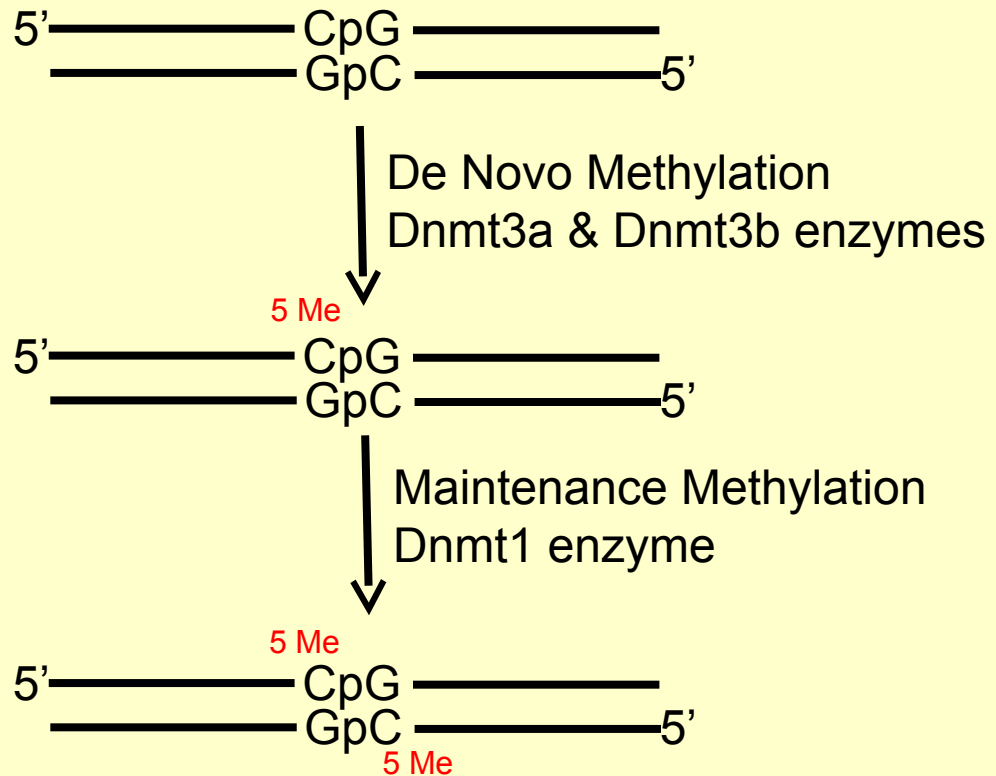
A combination of different molecules can attach to the "tails" of proteins called histones. These alter the activity of the DNA wrapped around them

Methylation of Cytosine in DNA

Cytosine methylation

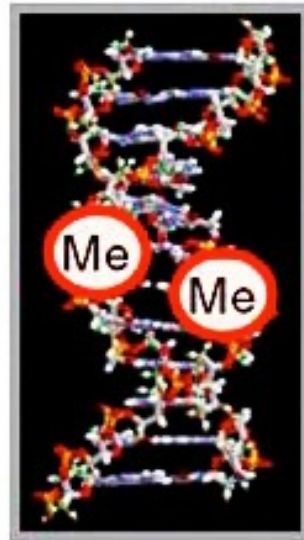


Only Cs in CG sequences are Methylated

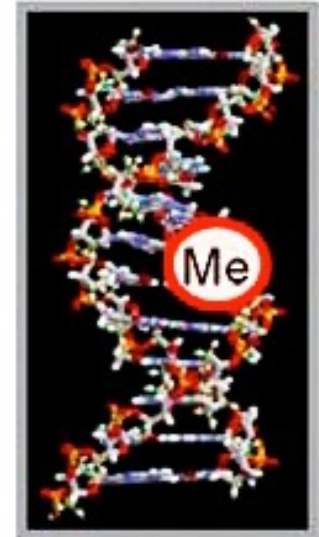


Maintenance of Cytosine Methylation

Establishment and maintenance

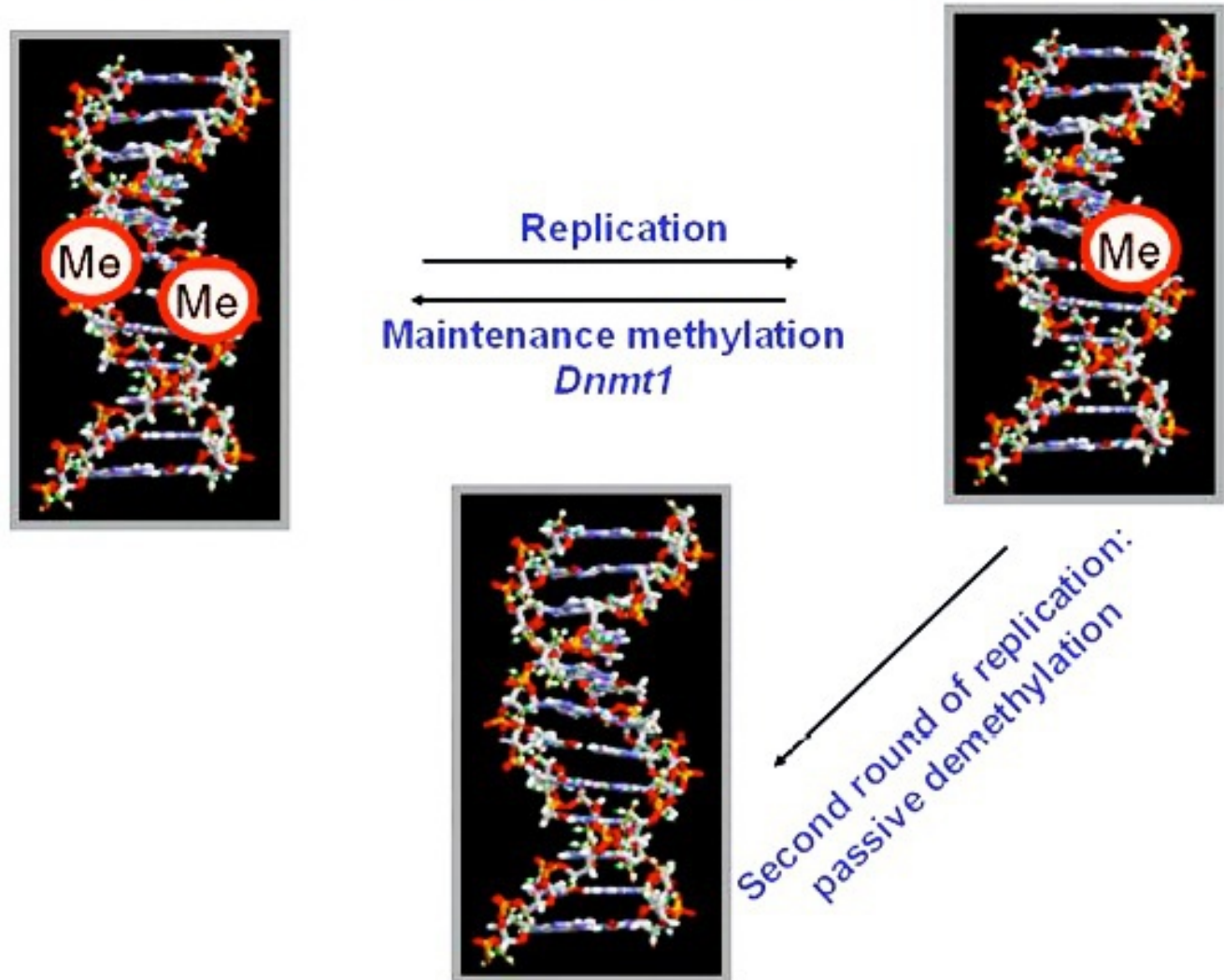


Replication
→
←
Maintenance methylation
Dnmt1

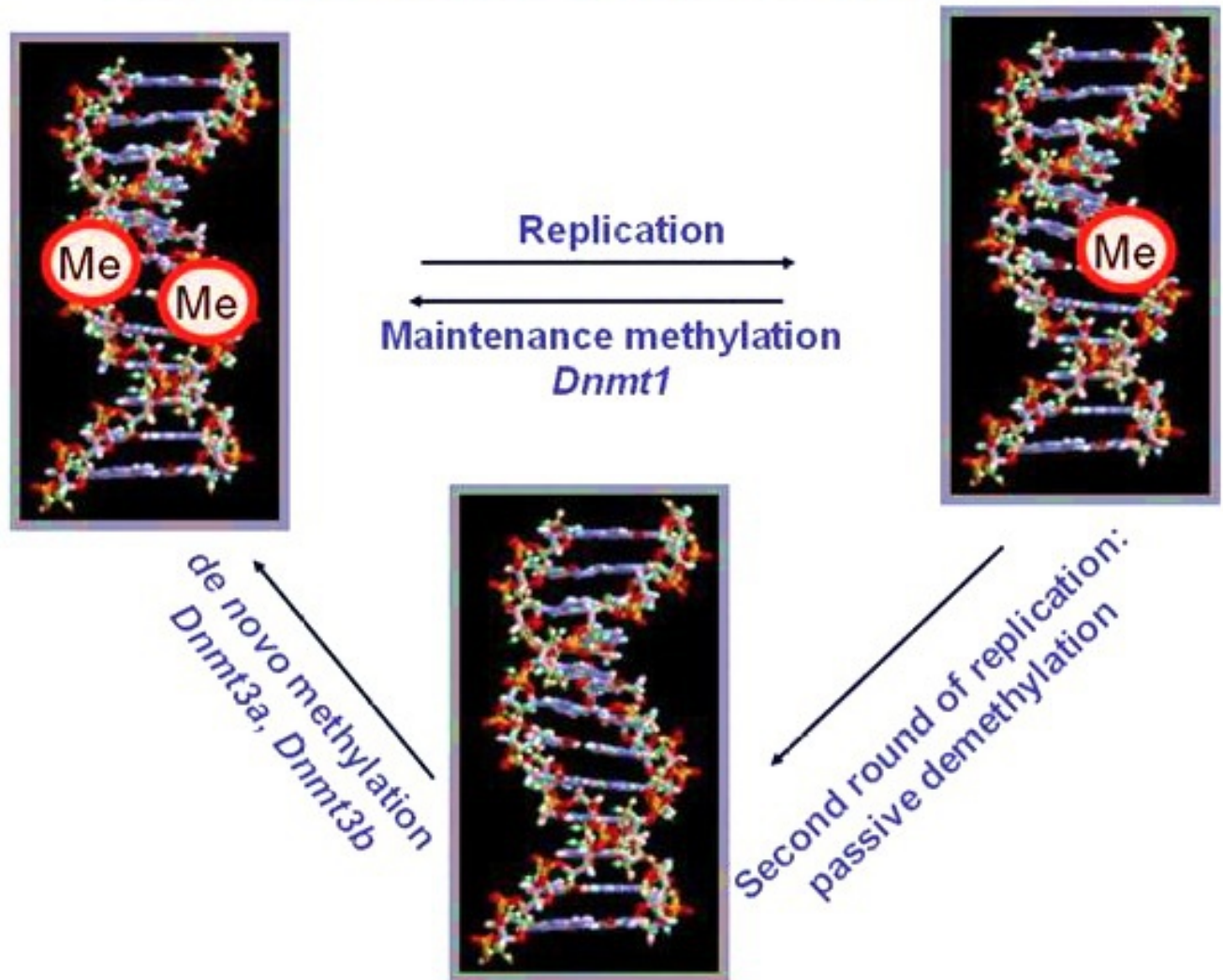


Passive Demethylation of 5-Methyl-Cytosine

Establishment and maintenance

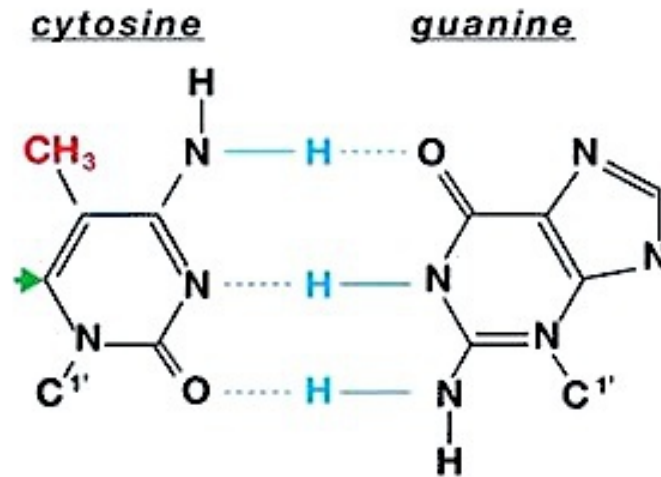


Establishment and Maintenance of Cytosine Methylation



5-Methyl Cytosine in DNA

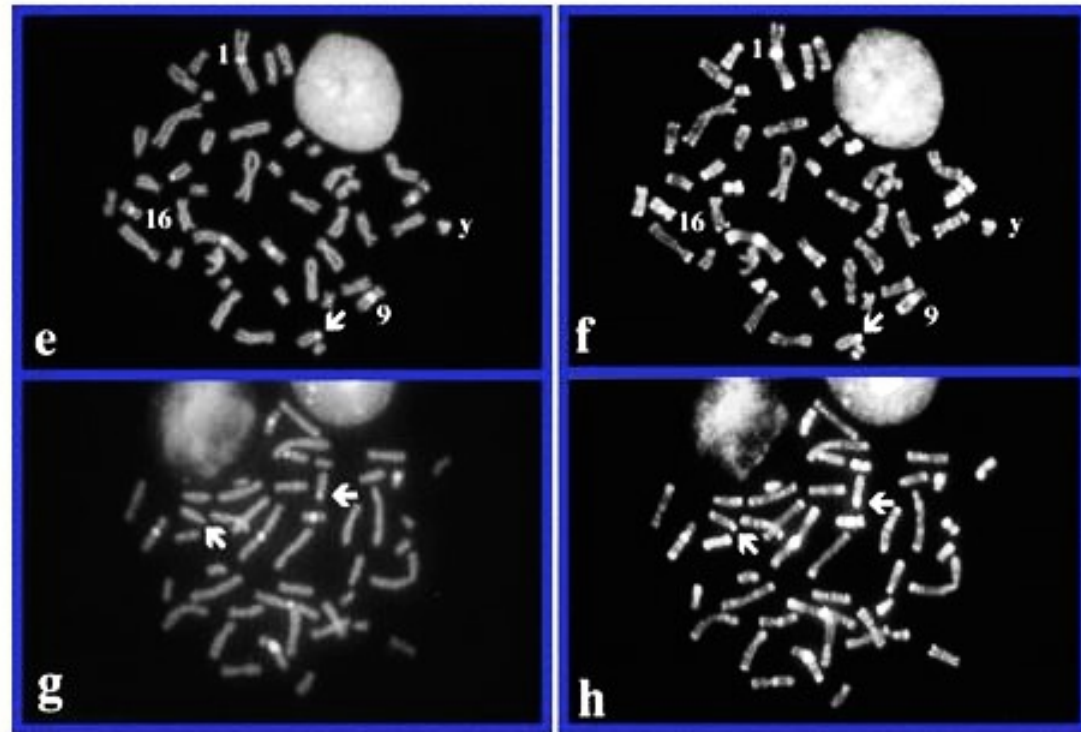
Cytosine methylation



5-Methyl Cytosine is Found in Heterochromatic Regions

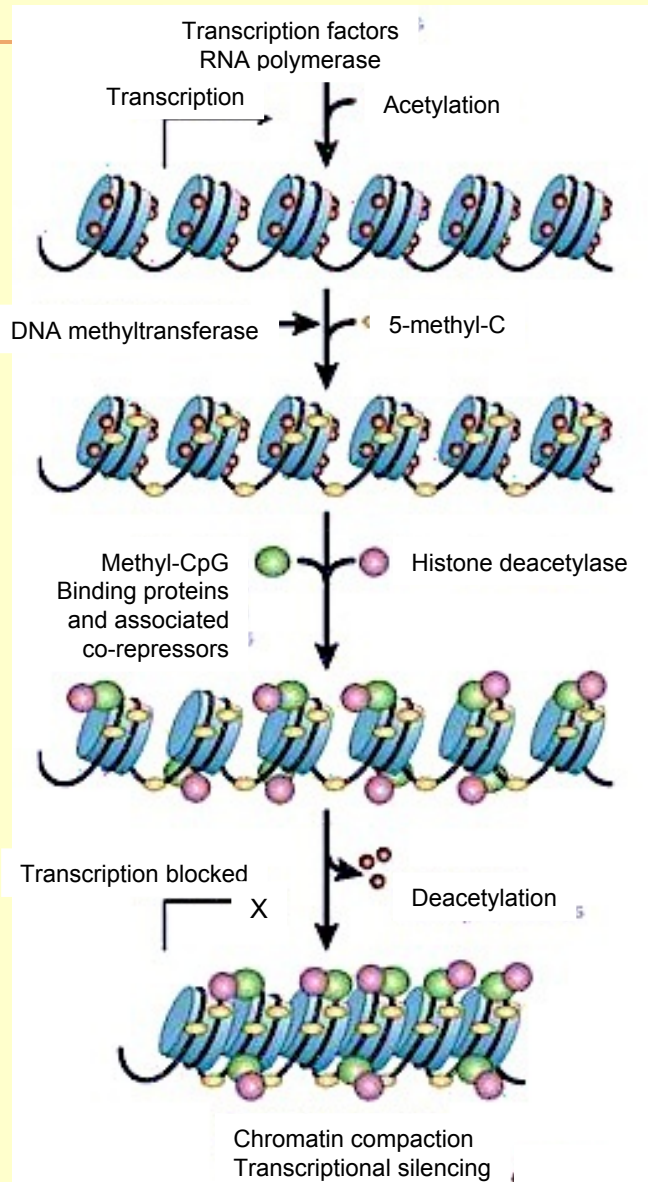
The distribution of cytosine methylation in mammals

- Heterogeneity visible at cytogenetic scale
- Associated with heterochromatic regions



PMID 9609658

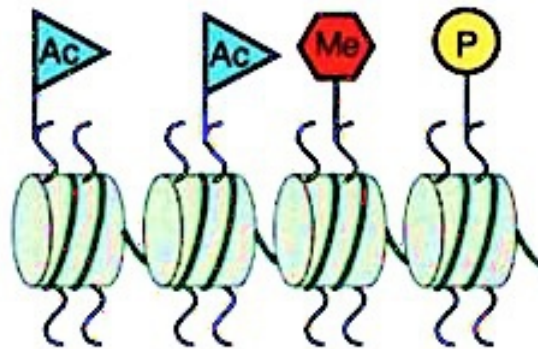
Cytosine Methylation Maintains Inactive-Condensed Chromatin State



Structure & Epigenetics of Euchromatin versus Heterochromatin

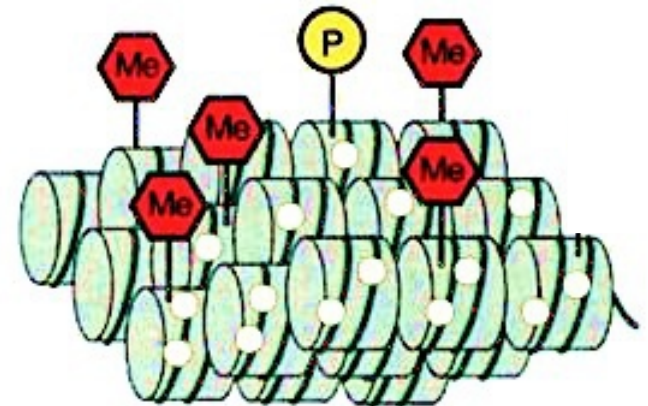
DNA methylation and histone modifications help to compartmentalize the genome into domains of different transcriptional potentials

Euchromatin



- High histone acetylation
- Low DNA methylation
- H3-K4 methylation

Heterochromatin



- Low histone acetylation
- Dense DNA methylation
- H3-K9 methylation

Histone Code

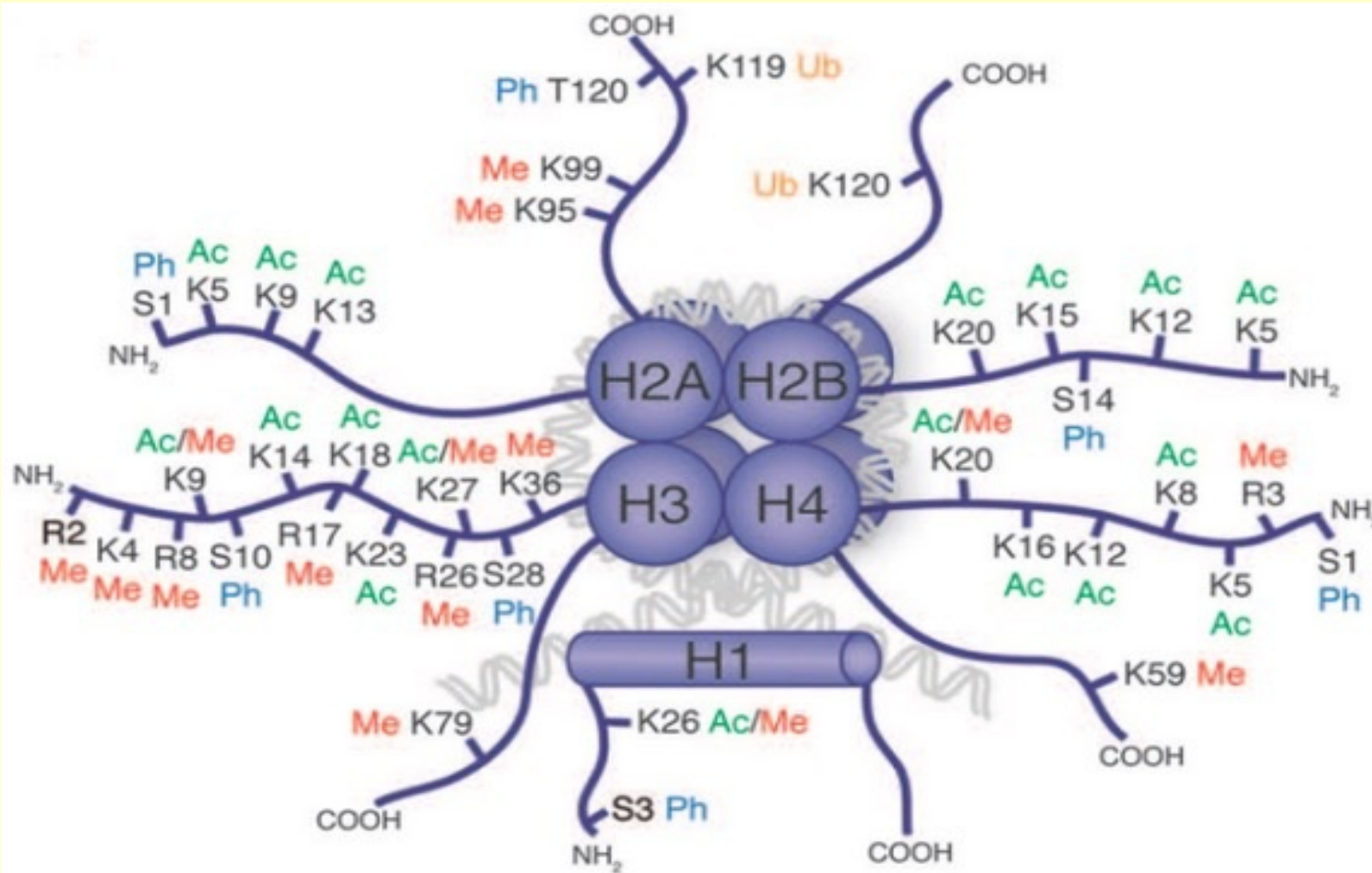
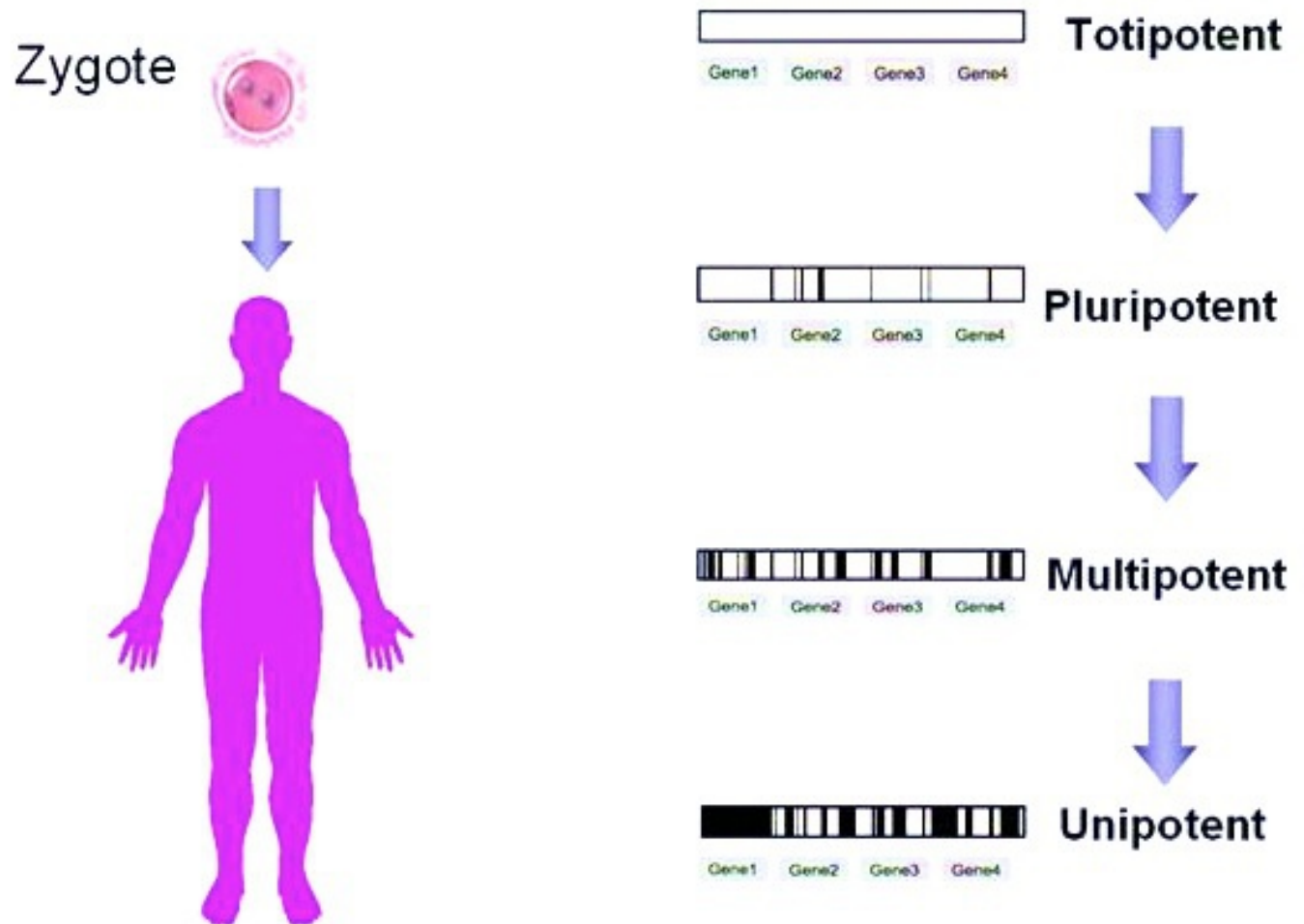


FIGURE 4.

Nucleosome with histone posttranslational modifications (Adapted from 1)

Methylated DNA from Zygote to Adult

Differentiated cells become more restricted in their potential



Methylated DNA from Zygote to Adult

Zygote



```
ACATAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCATGAT
GTTTTTATTACCAGGATGATCACCATTGGGTACCATTTACCAGGATTACACAGTTTTAGATGACC
AGTAGCTATTAGAGGATTTTAAATTTATTTAGGATTTTATGGGATTGATAAAGGGAGATTTAAC
TAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCATGATGTT
TTTATTACCAGGATGATCACCATTGGGTACCATTACCAGGATTACACAGTTTTAGATGACCAGT
AGCTATTAGAGGATTTTAAATTTATTTAGGATTTTATGGGATTGATAAAGGGAGATTTTATTAT
AGGACATAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCAT
GATGTTTTTATTACCAGGATGATCACCATTGGGTACCATTTACCAGGATTACACAGTTTTAGATG
ACCAGTAGCTATTAGAGGATTTTAAATTTATTTAGGATTTTATGGGATTGATAAAGGGAGATTTA
ACATAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCATGAT
```

**How is the diversity of cell types
created and maintained
in multi-cellular organisms?**

```
ACATAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCATGAT
GTTTTTATTACCAGGATGATCACCATTGGGTACCATTTACCAGGATTACACAGTTTTAGATGACC
AGTAGCTATTAGAGGATTTTAAATTTATTTAGGATTTTATGGGATTGATAAAGGGAGATTTAAC
TAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCATGATGTT
TTTATTACCAGGATGATCACCATTGGGTACCATTACCAGGATTACACAGTTTTAGATGACCAGT
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AGGACATAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCAT
GATGTTTTTATTACCAGGATGATCACCATTGGGTACCATTTACCAGGATTACACAGTTTTAGATG
ACCAGTAGCTATTAGAGGATTTTAAATTTATTTAGGATTTTATGGGATTGATAAAGGGAGATTTA
ACATAGACATACACACTGTTGATTAGGGAGATAGTGACAGATCCATTACAGCACCATACCATGAT
```

DNA Methylation Differentiates Totipotent Embryonic Stem Cells from Unipotent Adult Stem Cells

Pluripotent cell



ctggaggtgcaatggctgtcttgtcctggcctt
ggacatgggctgaaatactgggttcacccatat
ctaggactctagacgggtgggtaagcaagaact
gaggagtggccccagaaataattggcacacgaa
catteaatggatgttttaggctctccagaggat
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ttctcccacccccacagctctgctcctccacc
accagggggcggggcccagaggtcaaggctaga
gggtgggattggggagggagaggtgaaaccgt
cctaggtgagccgtctttccaccaggcccccg
ctcggggtgccaccttcccattggctggacac

Unipotent cell



Ctggaggtgcaatggctgtcttgtcctggcctt
ggacatgggctgaaatactgggttcacccatat
ctaggactctagacgggtgggtaagcaagaact
gaggagtggccccagaaataattggcacacgaa
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gggtgggattggggagggagaggtgaaaccgt
cctaggtgagccgtctttccaccaggcccccg
ctcggggtgccaccttcccattggctggacac

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Pluripotent cell



ctggaggtgcaatggctgtcttgtcctggcctt
ggacatgggctgaaatactgggttcacccatat
ctaggactctagacggggtgggtaagcaagaact
gaggagtggccccagaaaataattggcacacgaa
cattcaatggatgttttaggctctccagaggat
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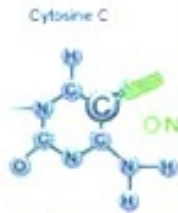
Unipotent cell



Ctggaggtgcaatggctgtcttgtcctggcctt
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ctaggactctagacggggtgggtaagcaagaact
gaggagtggccccagaaaataattggcacacgaa
cattcaatggatgttttaggctctccagaggat
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ttctccccccccacagctctgctcctccacc
accagggggcggggcccagaggtcaaggctaga
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cctaggtgagccgtctttccaccaggccccgg
ctcggggtgccaccttccccatggctggacac

DNA Methylation Differentiates Totipotent Embryonic Stem Cells from Unipotent Adult Stem Cells

DNA methylation



Pluripotent cell

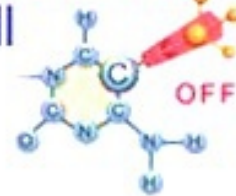


≠

Unipotent cell



Methyl-Cytosine 5mC

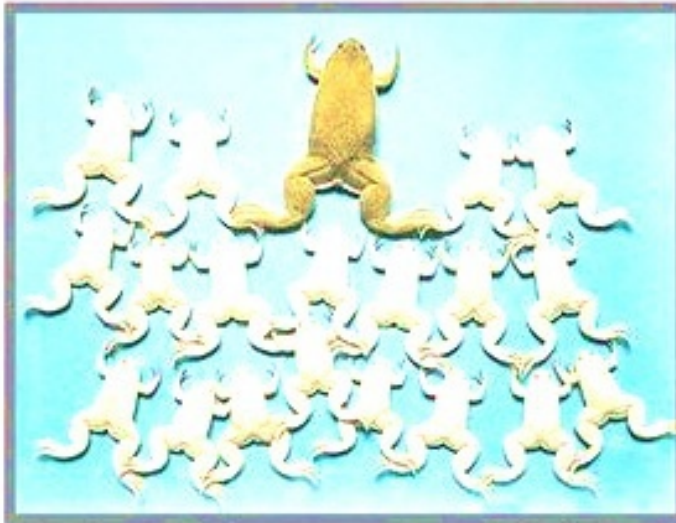


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Differentiated Cells can Become Totipotent

Nuclear transplantation demonstrates nuclear equivalence

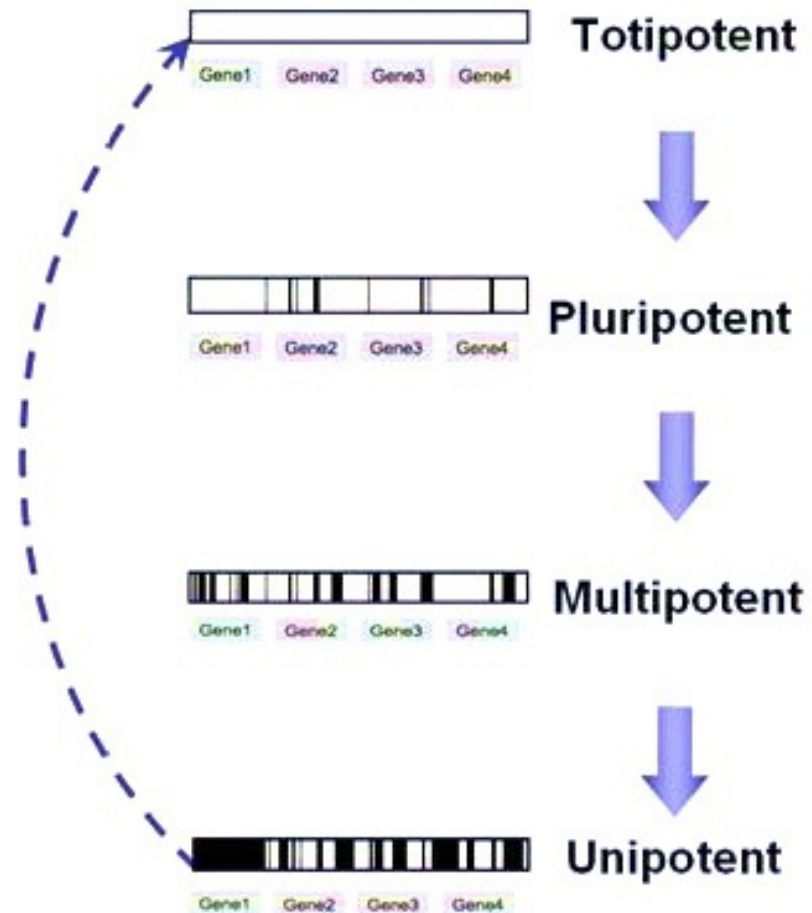


Briggs and King, 1952

Gurdon, 1960s

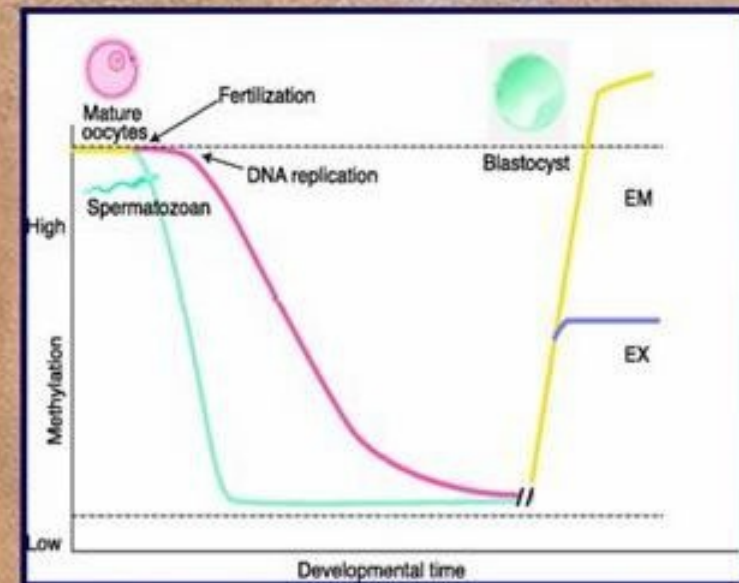
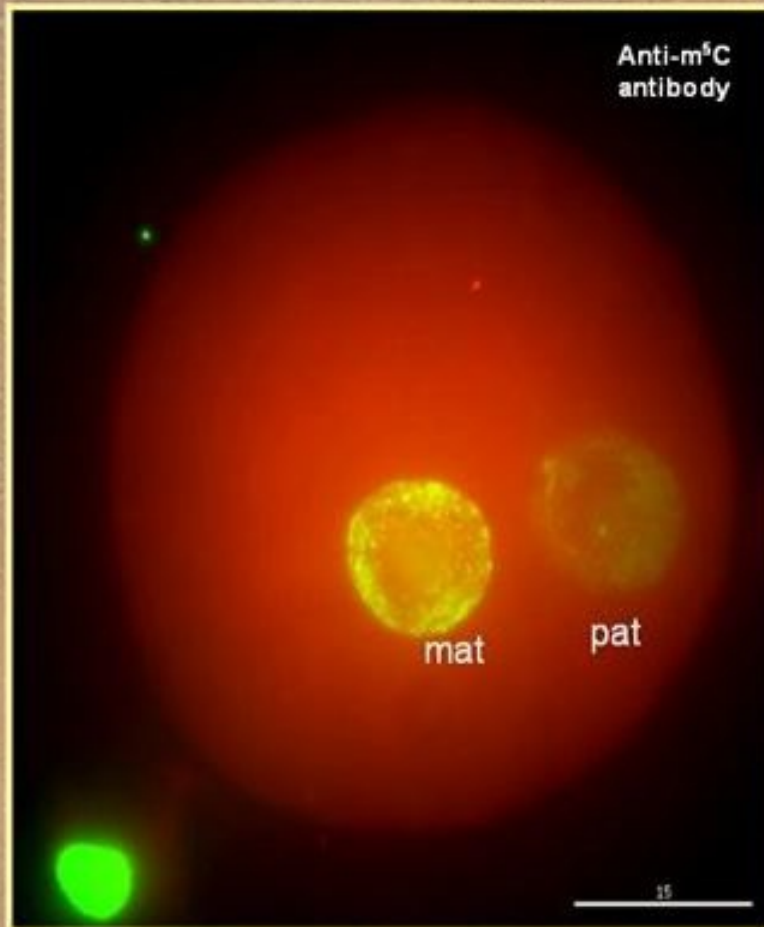
“Dolly”

Differentiated cells maintain the potential to generate an entire organism



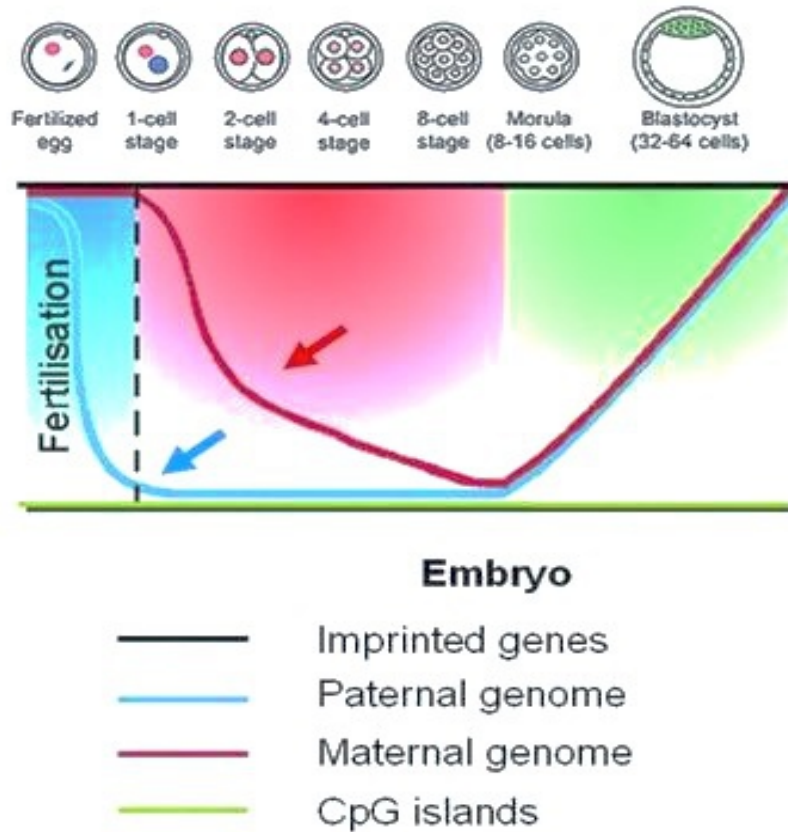
Demethylation of the Paternal Genome

De-methylation of the paternal pronucleus in the one-cell embryo of mouse



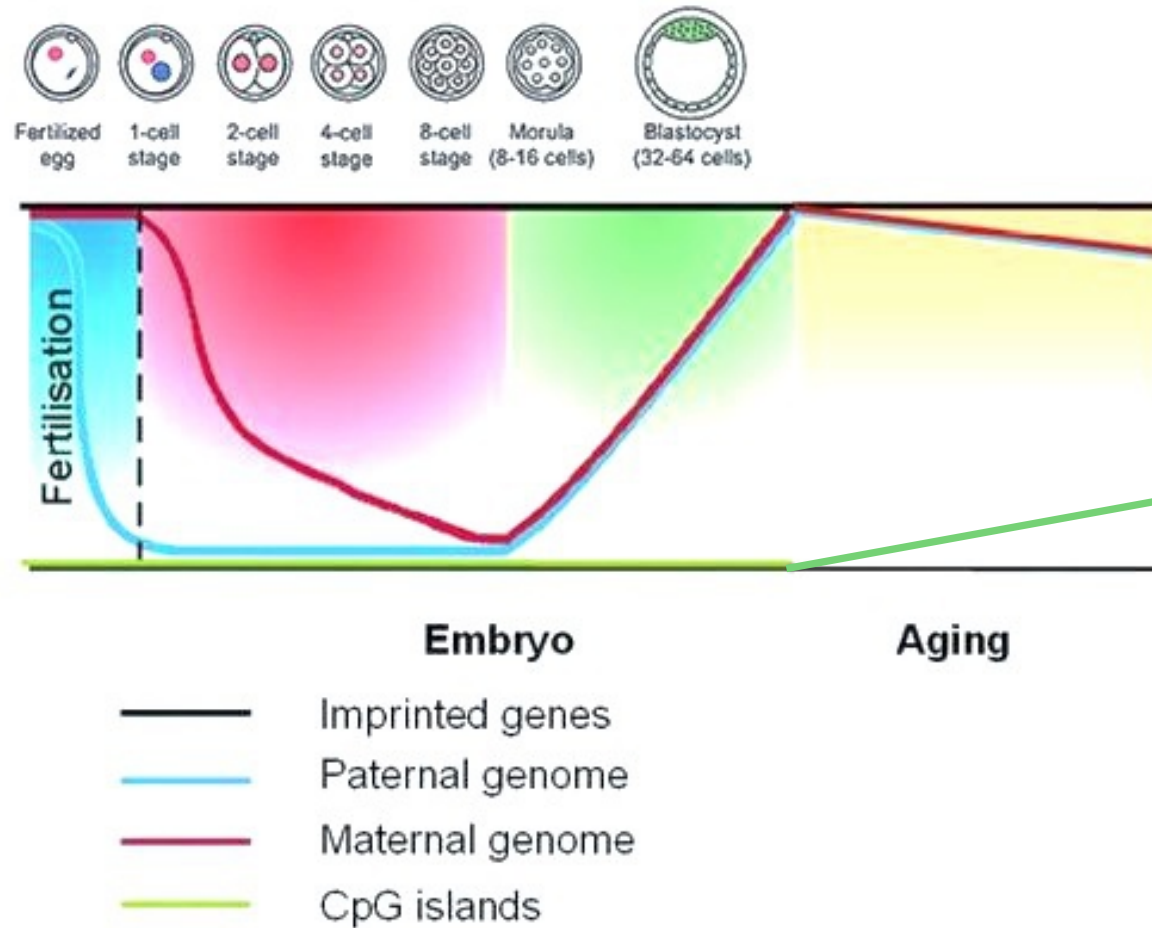
Methylation Changes During Development

Reprogramming the DNA methylome



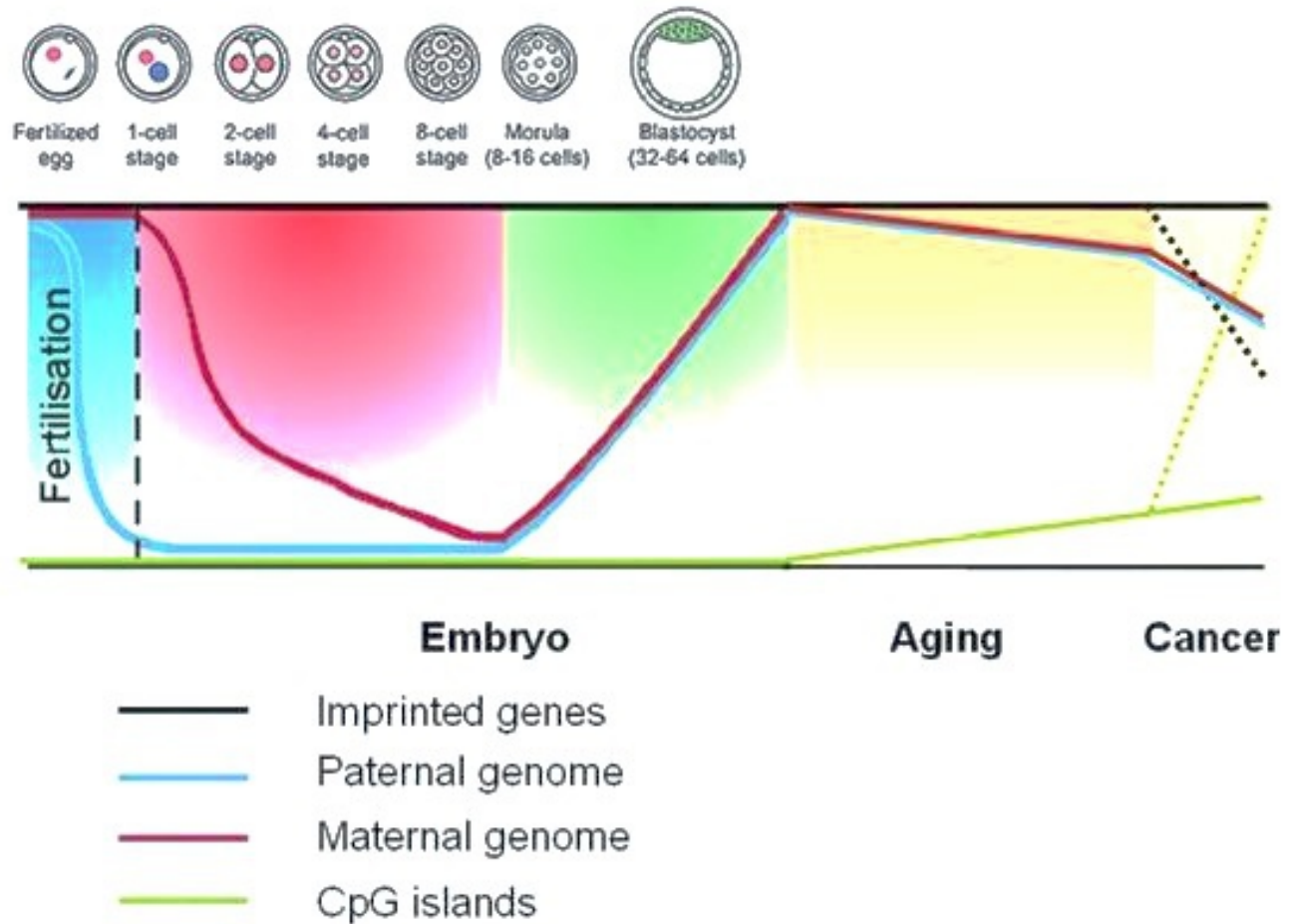
Methylation Changes During Development

Reprogramming the DNA methylome

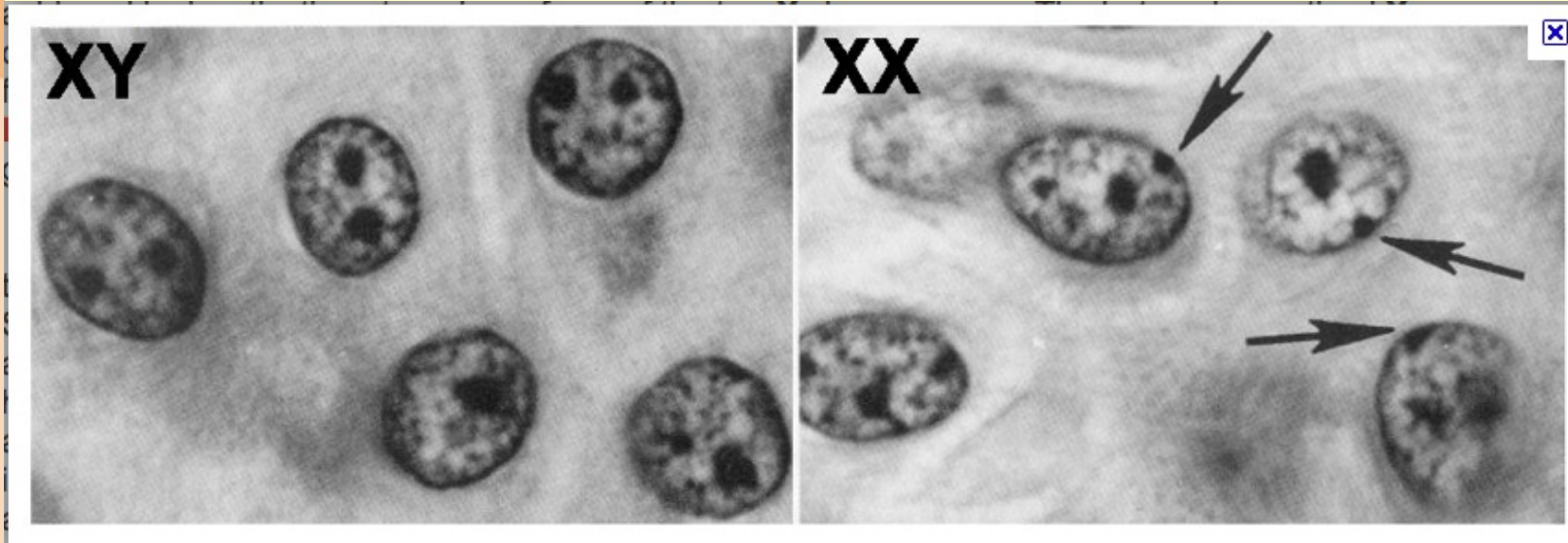


Methylation Changes During Development

Reprogramming the DNA methylome



X Chromosome Inactivation: Barr Bodies

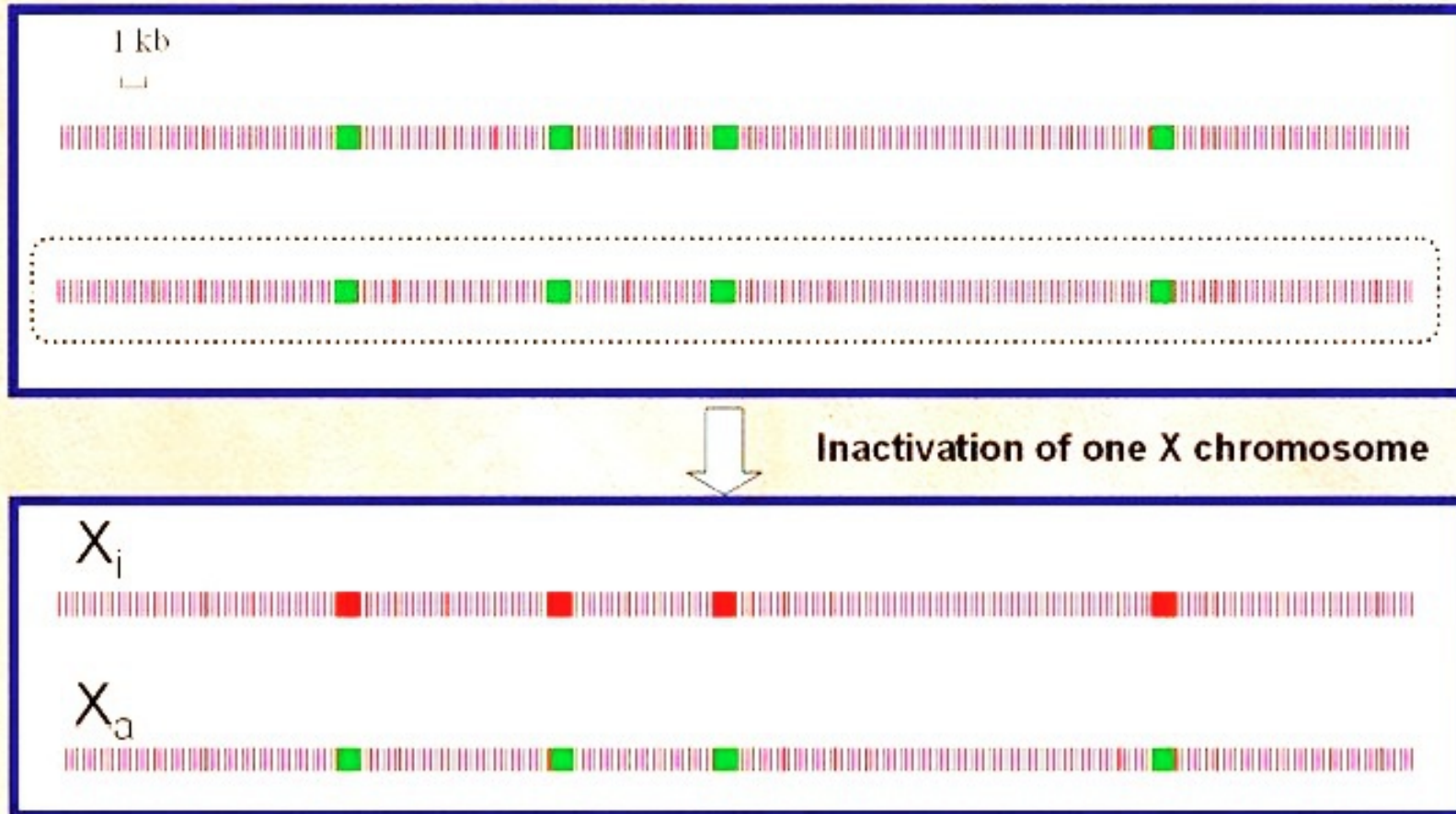


Barr, M. L., Bertram, E. G., (1949), A Morphological Distinction between Neurones of the Male and Female, and the Behaviour of the Nucleolar Satellite. *Nature*. **163** (4148): 676-7.

Lyon, M. F., (2003), The Lyon and the LINE hypothesis. *j.semcd* 14, 313-318. (Abstract)

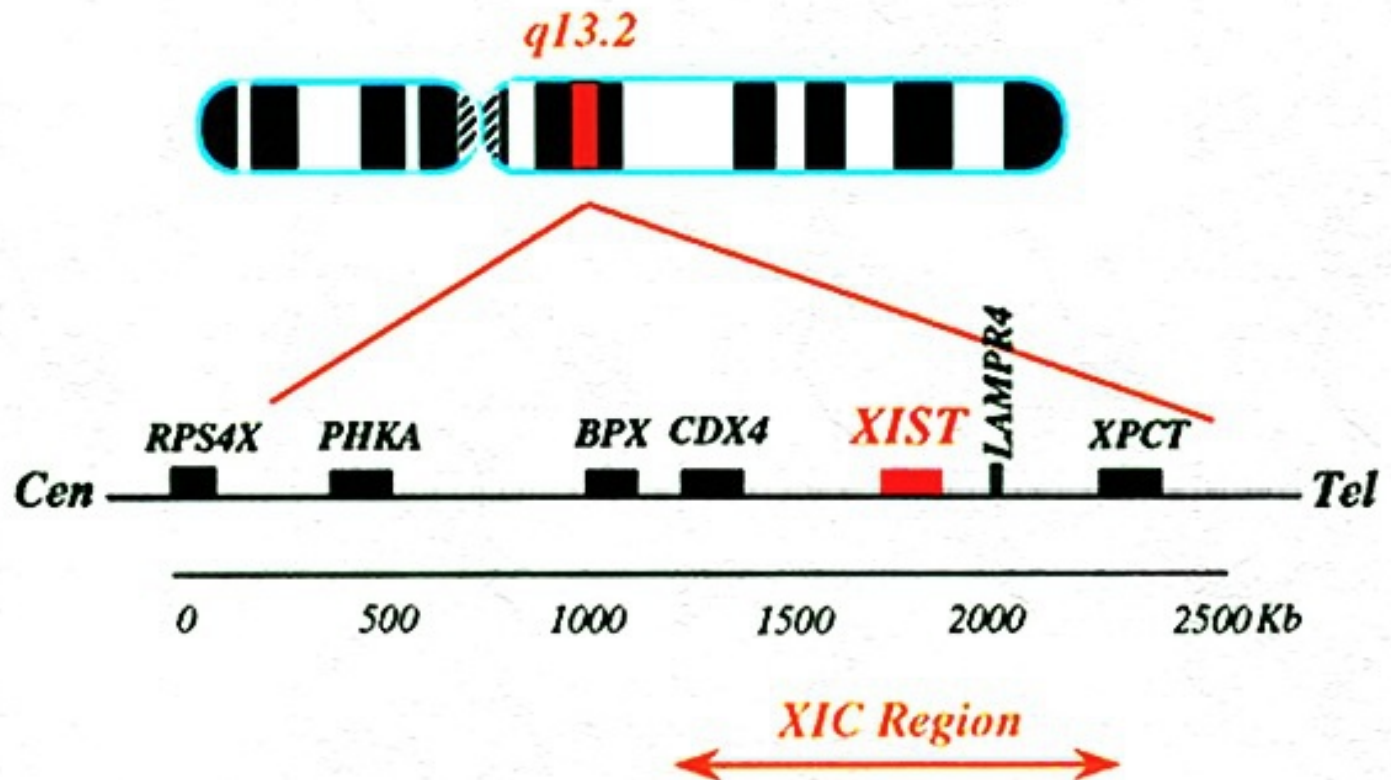
X Chromosome Inactivation: CG Island Methylation

De novo methylation of CpG islands on the inactive X chromosome



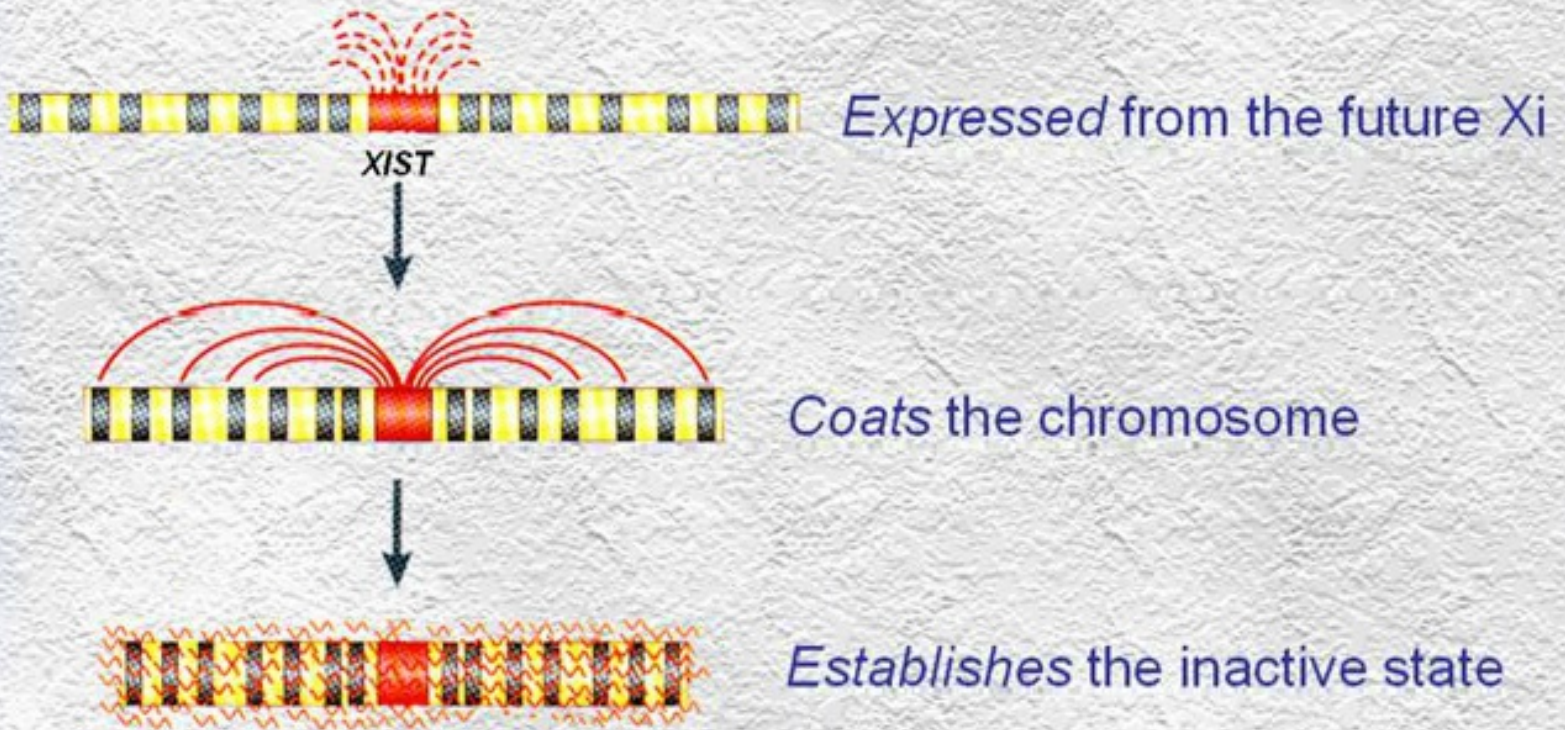
XIC Region

The XIC region on the human X chromosome



Xist Works in Cis

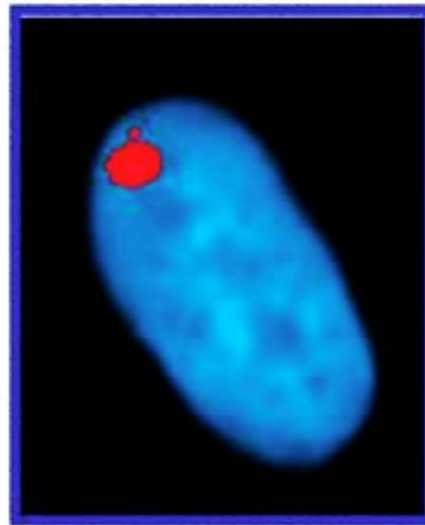
How *XIST* silences the future inactive X



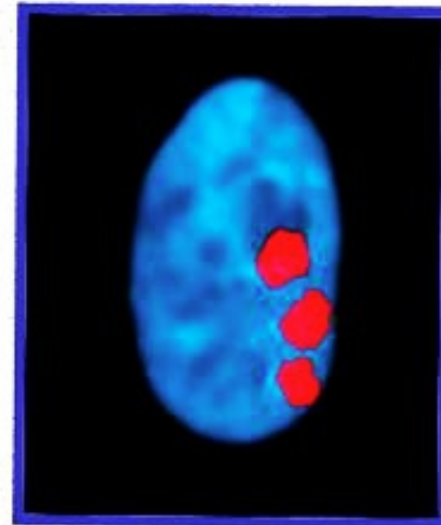
After Avner

Only one X is active

46, XX female

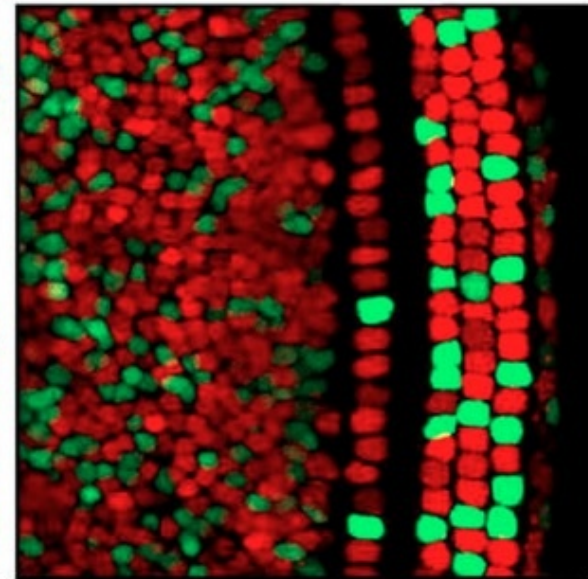
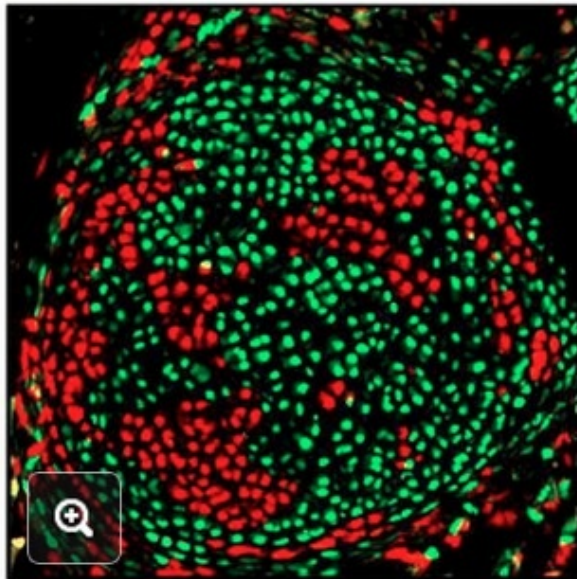
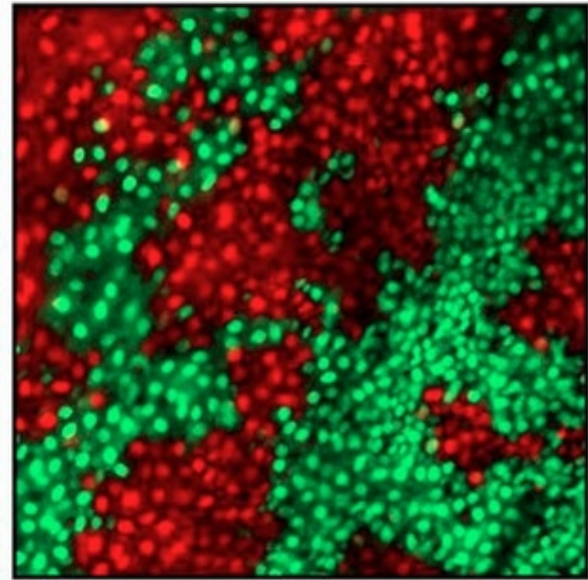
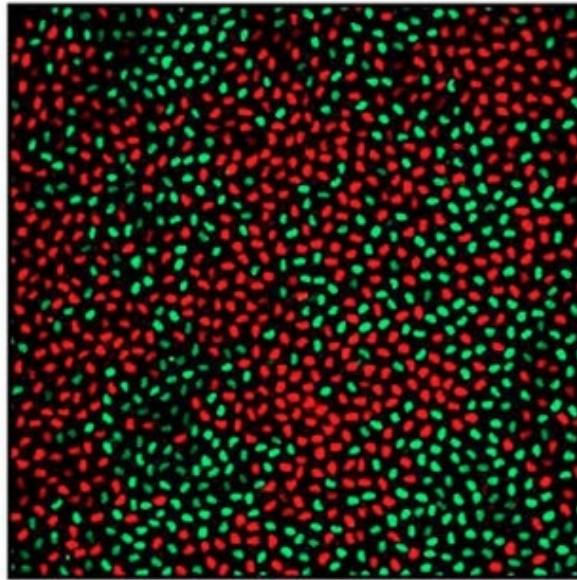


49, XXXXY male



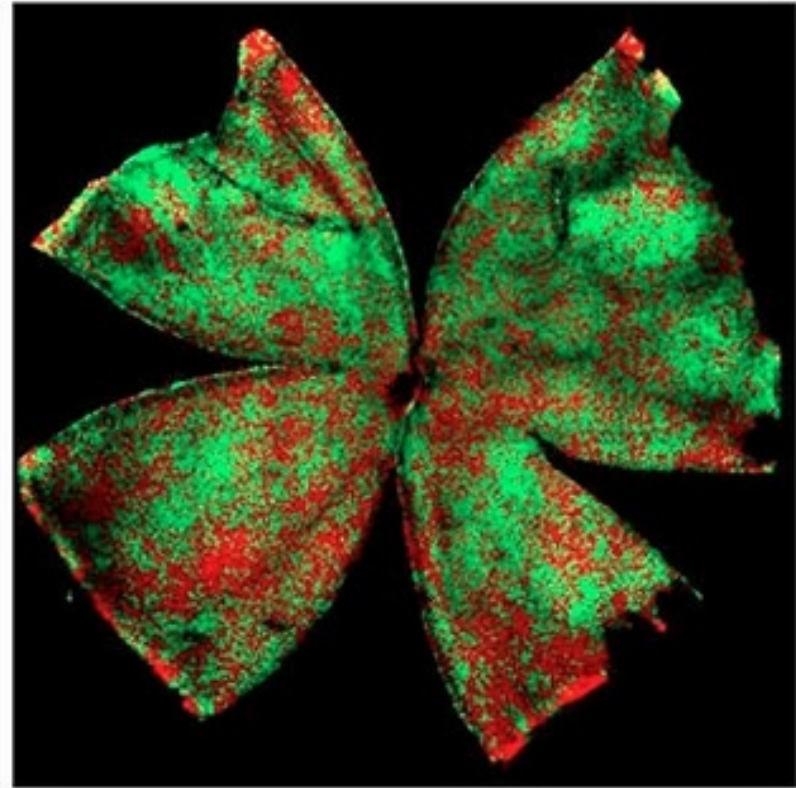
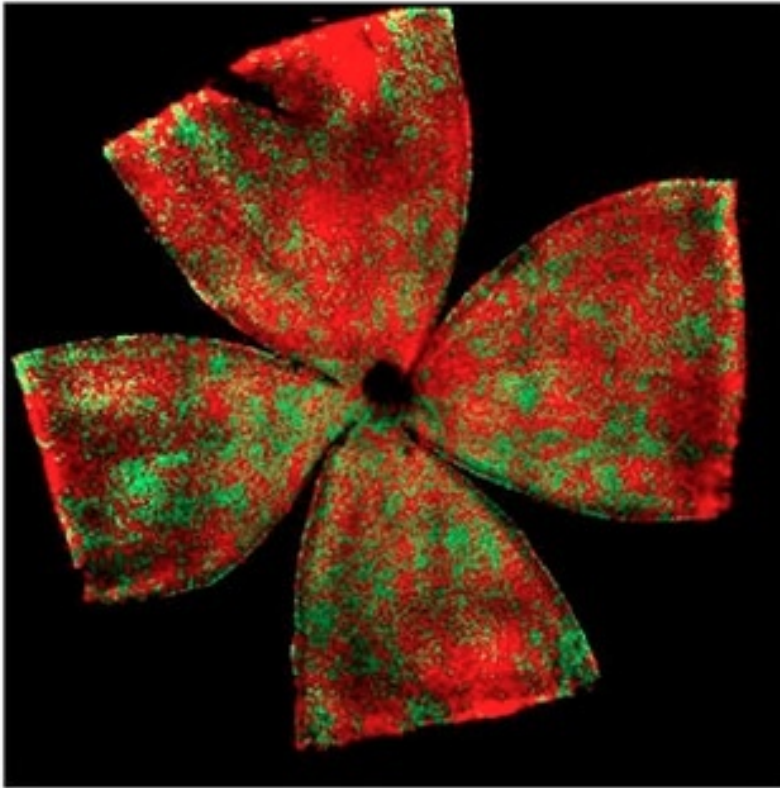
Barr bodies visualized by XIST RNA FISH

Female X chromosome Mosaicism (cornea, skin, cartilage & inner ear)



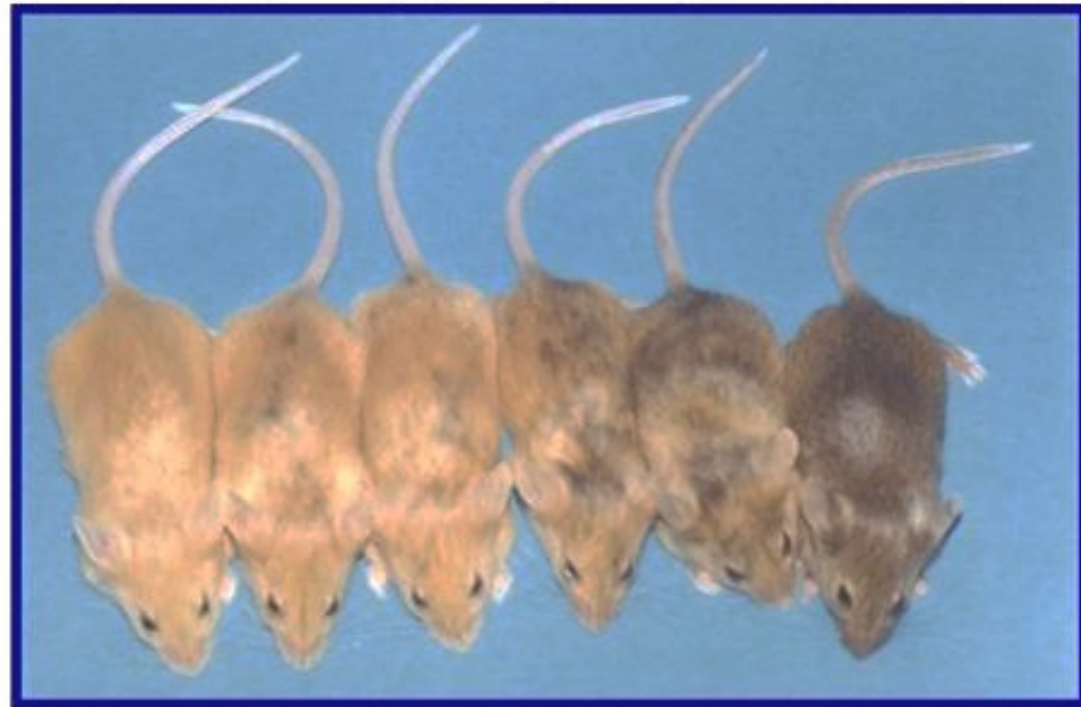
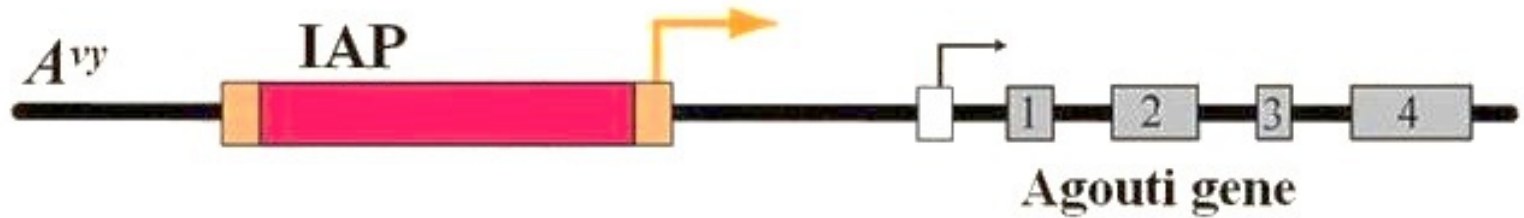
Female X chromosome Mosaicism

Left and Right Retina

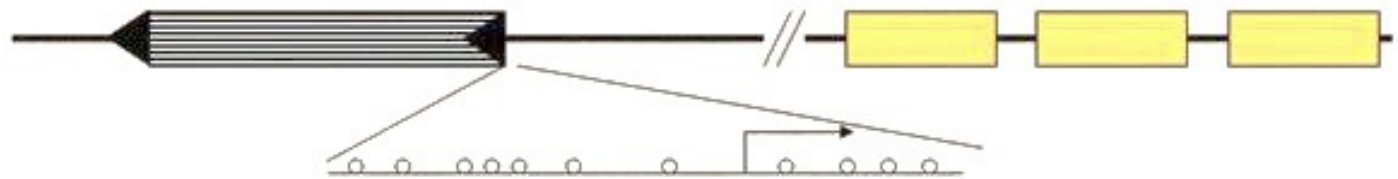


Agouti Genes in Mice

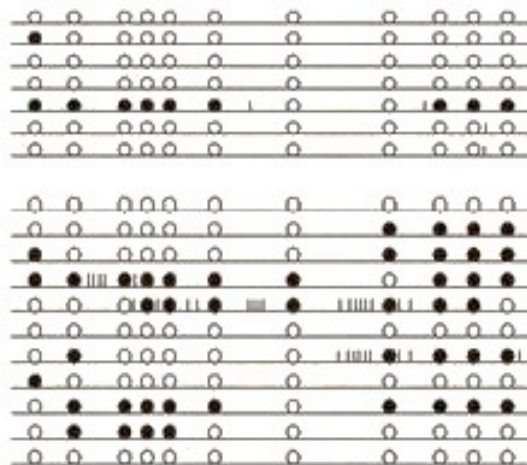
Agouti viable yellow (A^{vy})



Methylation of Agouti Genes in Mice

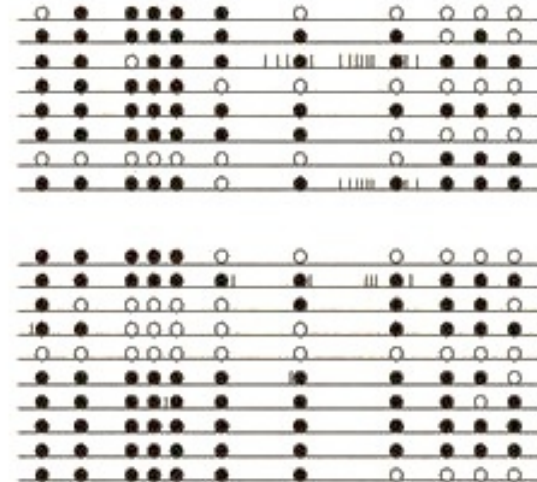


Yellow



27%
mCpG

Pseudoagouti



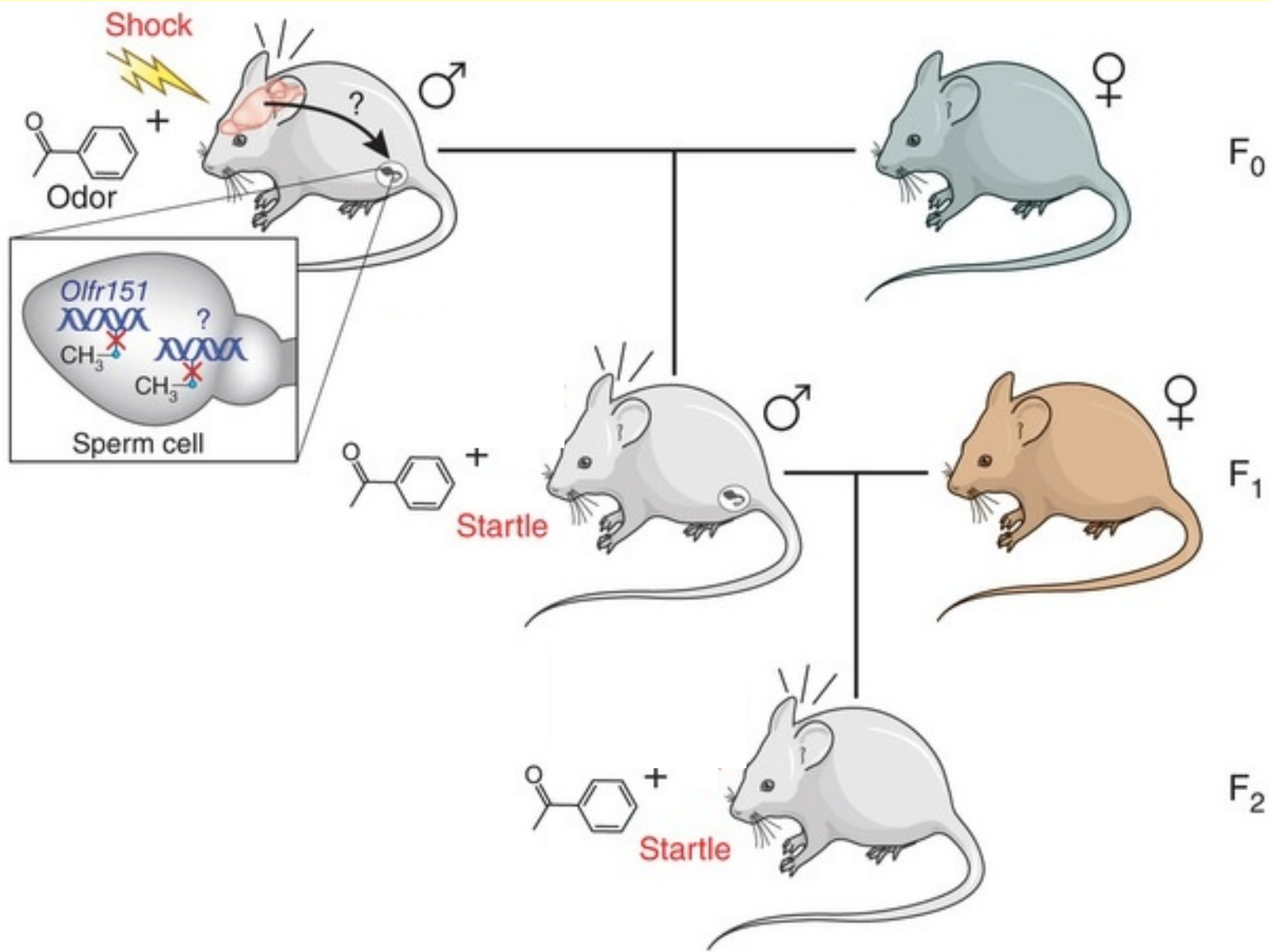
69%
mCpG

Hongerwinter 1944

- German's blocked food to the Dutch in the winter of 1944.
- Calorie consumption dropped from 2,000 to 500 per day for 4.5 million.
- Children born or raised in this time were small, short in stature and had many diseases including, edema, anemia, diabetes and depression.
- The Dutch Famine Birth Cohort study showed that women living during this time had children 20-30 years later with the same problems despite being conceived and born during a normal dietary state.



Epigenetic Inheritance of Fear



Summary of Epigenetic Gene Regulation

- Patterns of DNA methylation in adult cells parallels cell fate, chromatin structure and gene activation.
- Most DNA methylation is removed at fertilization and re-established during embryogenesis.
- Imprinted genes keep their parental pattern of methylation giving rise to parental patterns of expression.
- Patterns of histone modifications parallel DNA methylation.
- Methylated gene regions are genetically inactive, highly condensed and special histone modifications.
- Active gene regions have little DNA methylation and distinctive histone modifications (acetyl groups and H3K4methyl).
- X chromosome inactivation in females is correlated with extensive CG island methylation on one chromosome, condensation, inactivation and Barr body formation.
- Alterations in gene and CG island methylation patterns are seen in aging and in cancer.
- Most CG islands are not methylated except for X chromosome inactivation and tumor suppressors in cancer.