

Epigenetics

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Genetics –

“We are what our parents are”

Continuity

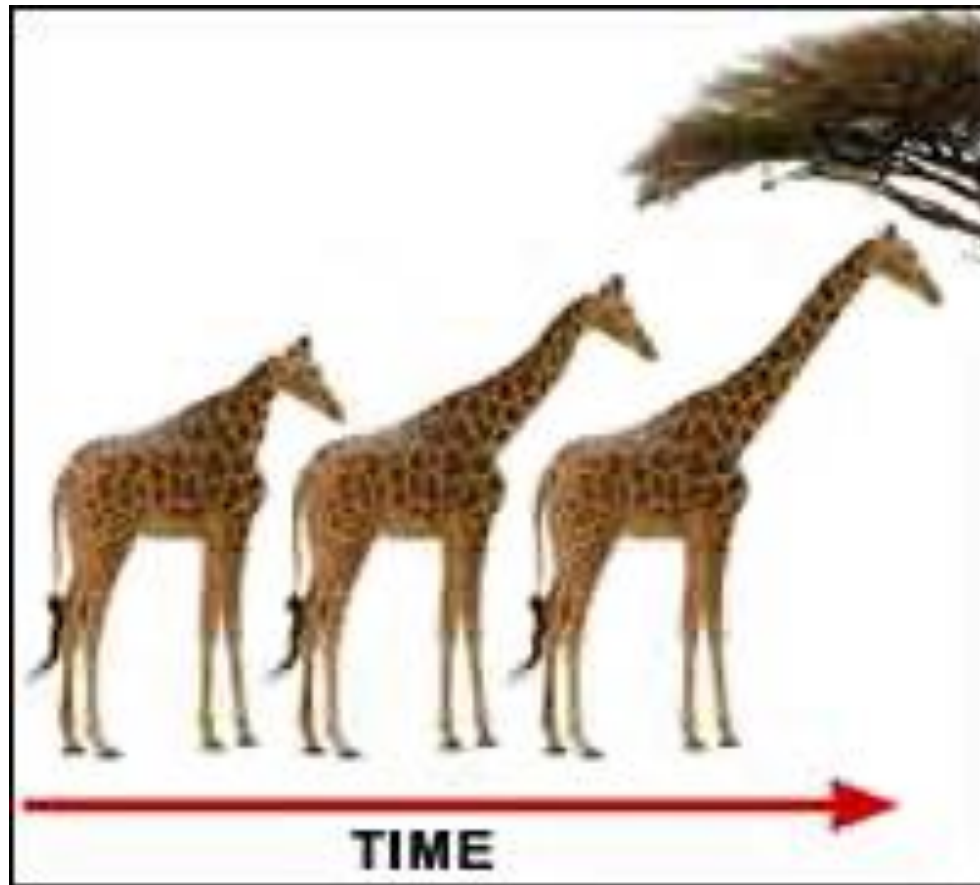
Epigenetics –

“Changes at the cellular level in response to
our environment that we can pass
to our children”

Lamarckian theory (1700's):

Change Through Use and Disuse

“Heritability of acquired characteristics”



Take home message:
Open-mindedness

The “wrong” answers of today
could be
a Nobel prize
of
tomorrow”

GENETICS:

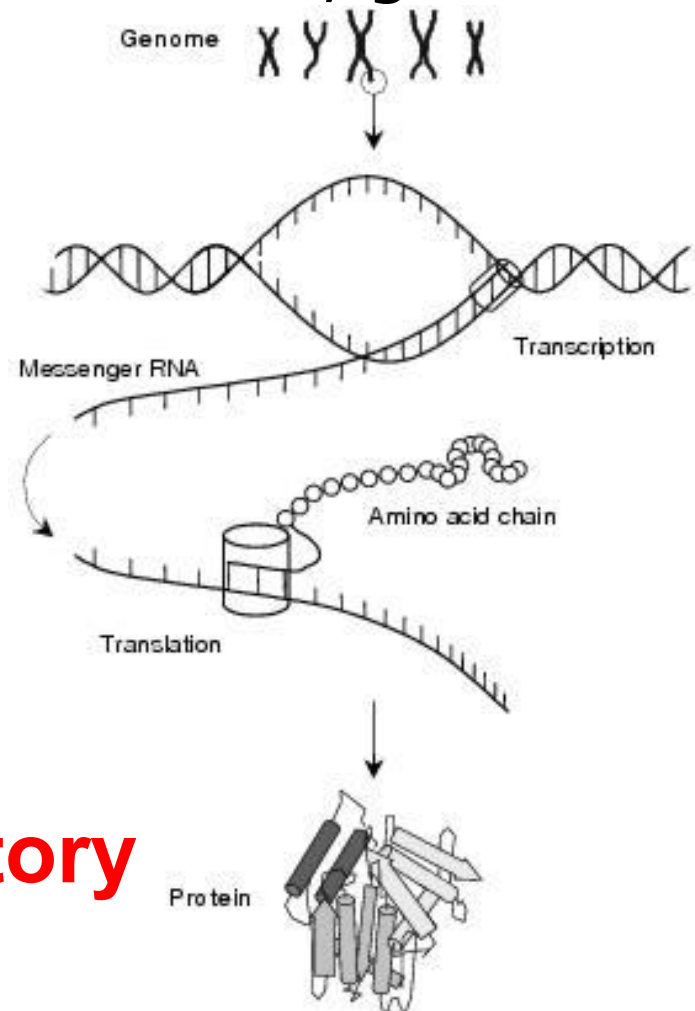
The study of heredity (in the past) of mostly DNA coded information from cell to cell, generation to generation.

CENTRAL DOGMA:

Genes/DNA

- RNA (Transcription)
- Protein (Translation)
- Functions/Traits

DNA is only part of the story



Chromosome-wide regulation:

X inactivation in mammals

Female: XX

Male: XY

Bar body:

- One of the two Xs in the female stays highly condensed.
- Transcription is inactivated on the Bar body.
- The inactivated X contains hypermethylated DNA.

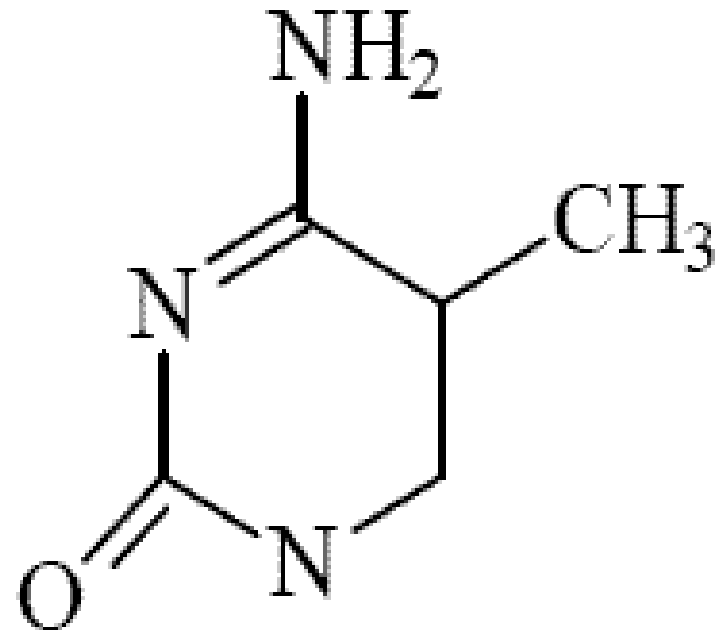
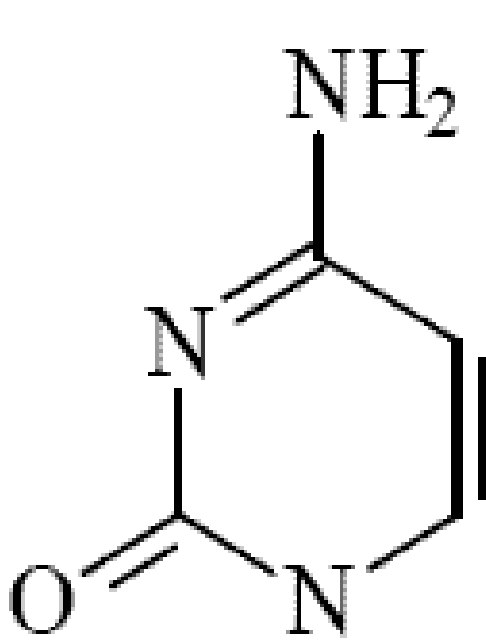


random or skewed
X inactivation



- **Recent concept:** a large number of genes can variably escape silencing on one or both chromosomes (aging).
- X-chromosome vulnerability
- X has enriched immune-related genes → Age-dependent autoimmune diseases

DNA Methylation

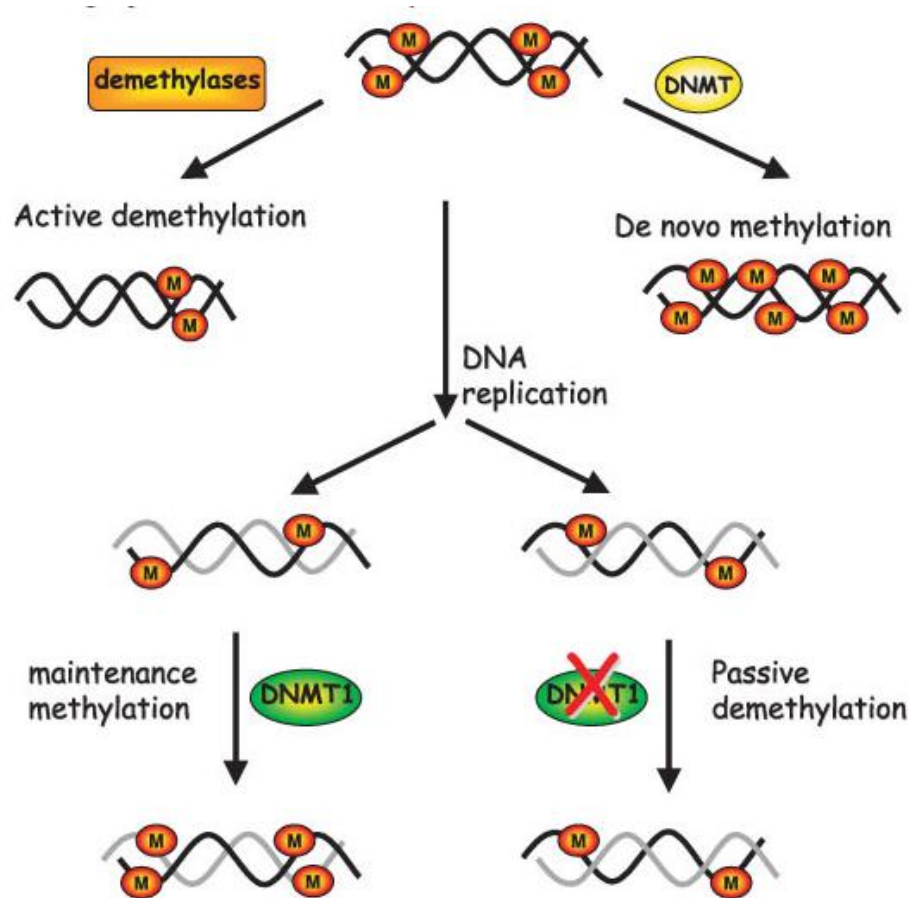


Cytosine $\xrightarrow{\text{DNMT}}$ 5'-methylated cytosine

DNA methyl transferase (DNMT)

De novo vs. maintenance methylation of DNA

Massive demethylation occurs in early embryos.



This is followed by de novo Methylation.

Then, maintenance methylation.

Loss of 5%
per CpG site
per cell division

Maternal nutritional influences on the fetal epigenome

A



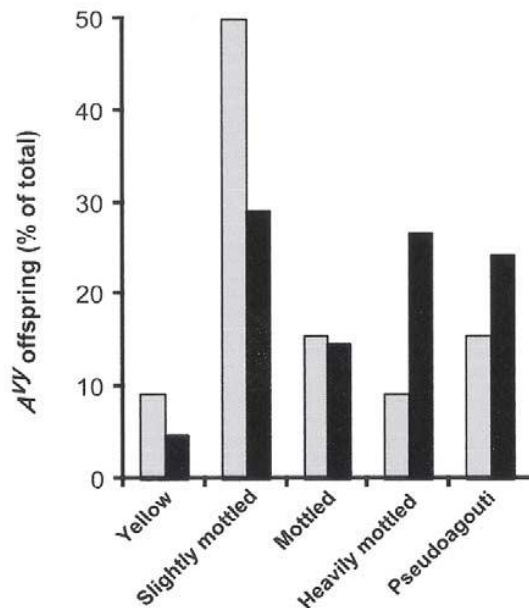
Agouti^{vy} or A^{vy}
a metastable epiallele

Coat colors

Yellow: Hypo-methylation of A^{vy}
increased expression of A^{vy}

Black: hyper-methylation of A^{vy}
decreased expression of A^{vy}

B

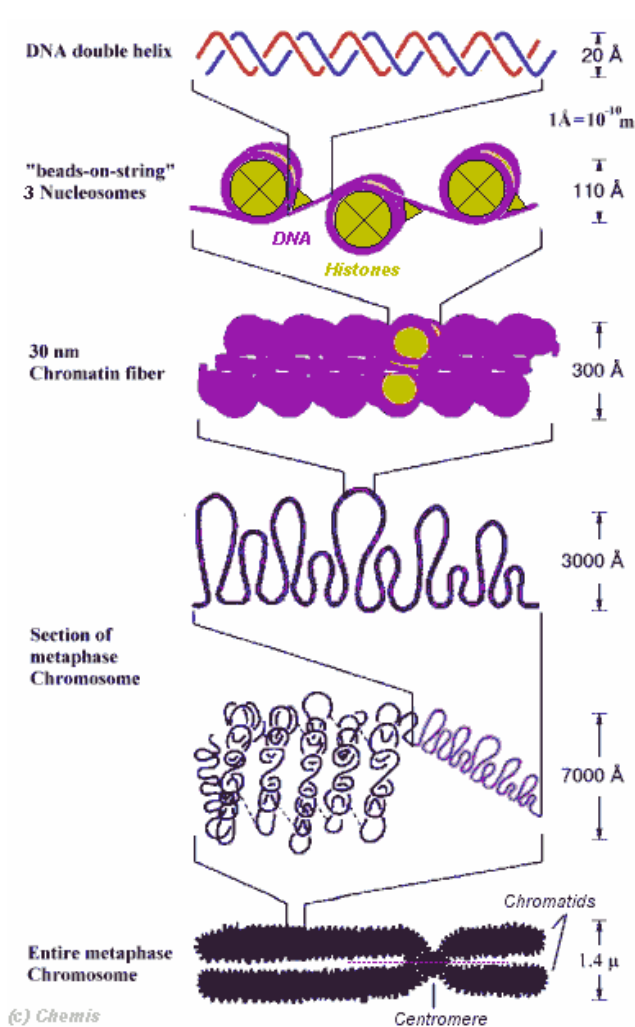


1. Maternal diet supplements affect pups' coat colors

Grey bar: NIH31 diet

Black bar: NIH31 supplemented with methyl donors and cofactors

Epigenetic information is also embedded in Chromatin (DNA + histones)



Chromatin of varying compaction

Bead-on-strings

30 nm fibers

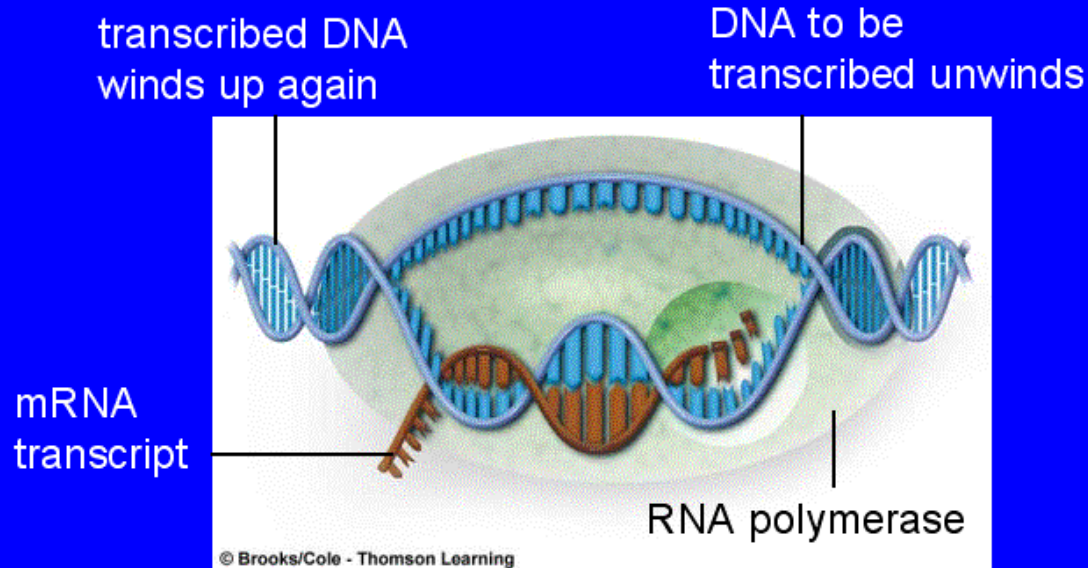
Chromatin loops

Partially condensed chromatin

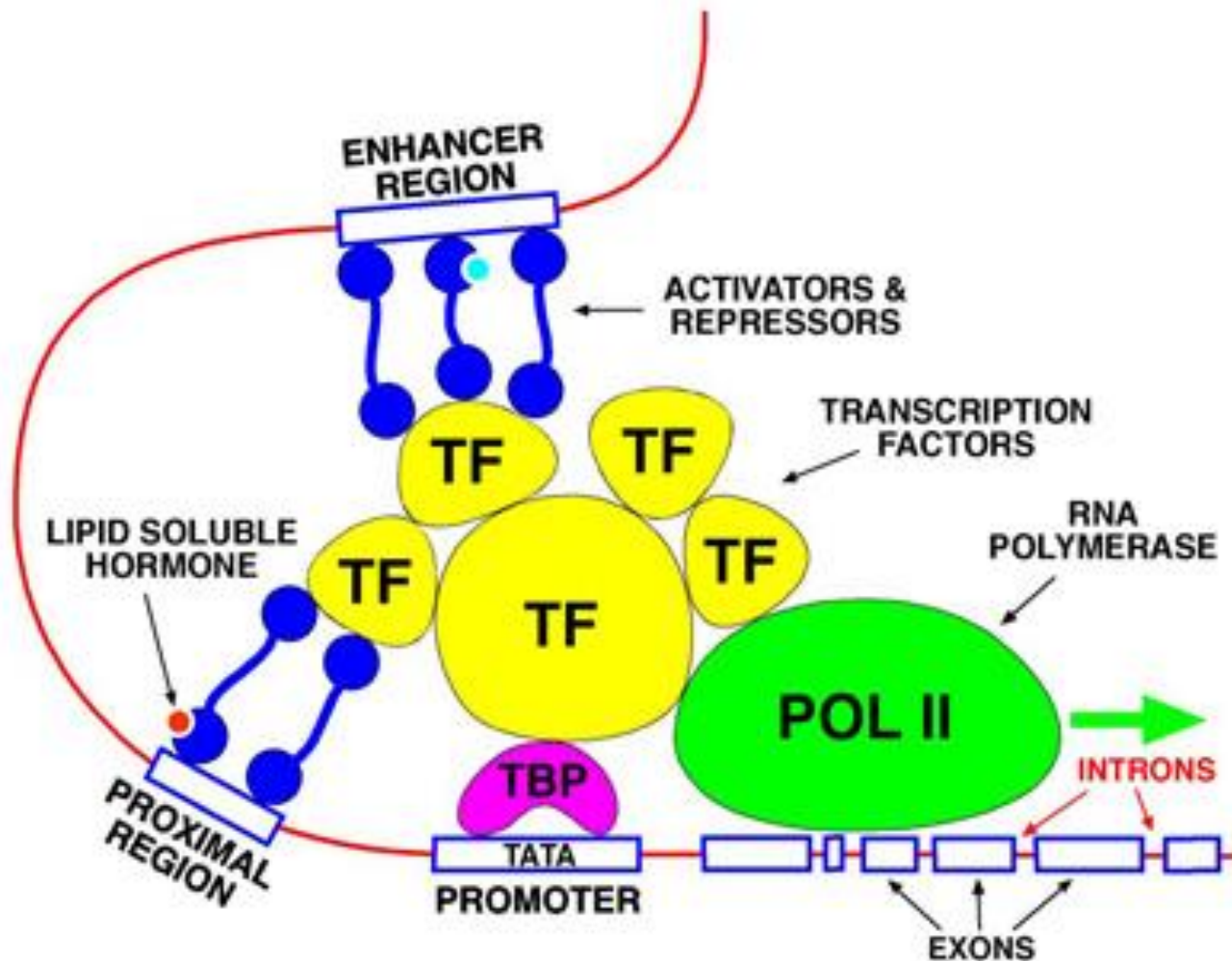
Mitotic chromosome (fully condensed)

Chromatin affects promoter access to proteins involved in transcription

Gene Transcription



Chromatin affects enhancer-promoter interaction



Epigenetic information –

Chemical modifications on DNA & histones

- Direct the formation of chromatin structure
- Influence the transcription of genes

Txn = Transcription

Tln = Translation

EPIGENETICS:

A new branch of genetics. Heredity of non-DNA sequence coded information from cell to cell, generation to generation.

Modified Central Dogma

GENOME = The whole collection of DNA sequence info

+

EPIGENOME = The whole collection of epigenetic information on **chromatin or elsewhere**, including:

Modifications on DNA

Modifications on Histones

Histone variants

Chromatin associated proteins

Non-coding or micro RNAs (carried in sperm)



**CHROMATIN
STRUCTURE** → Txn → RNA → Protein → Cell → Function/Trait

EPIGENETICS:

“The study of heritable alternations that result in gene expression changes without changes in DNA sequences”

“The study of heritable changes in gene function that cannot be explained by changes in DNA sequence”

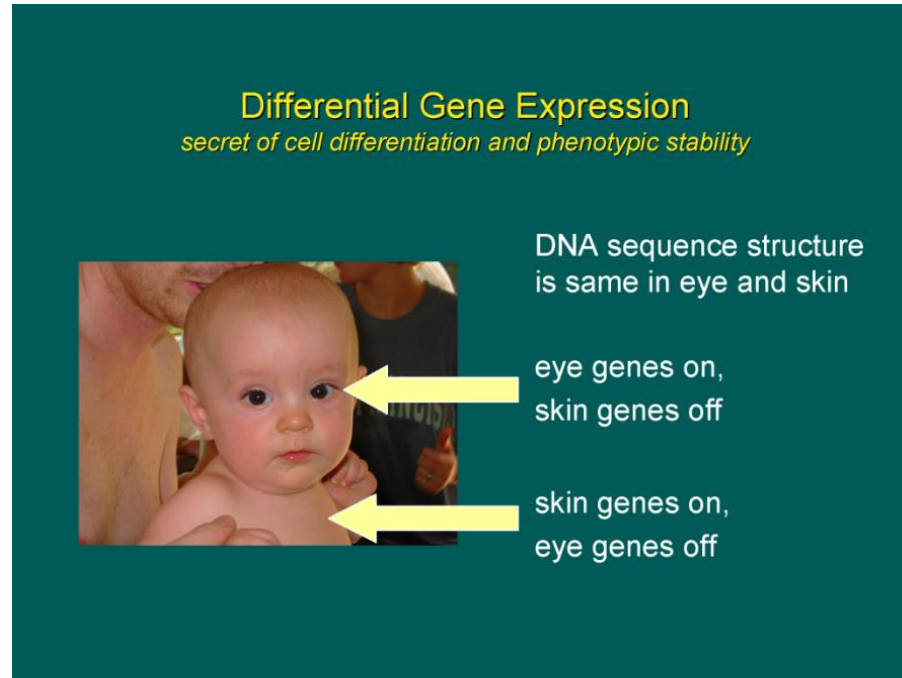
Heredity of non-DNA coded information.

Distinction between

Genetic (DNA-coded) versus Epigenetic (non-DNA coded).

Epigenetic phenomena:

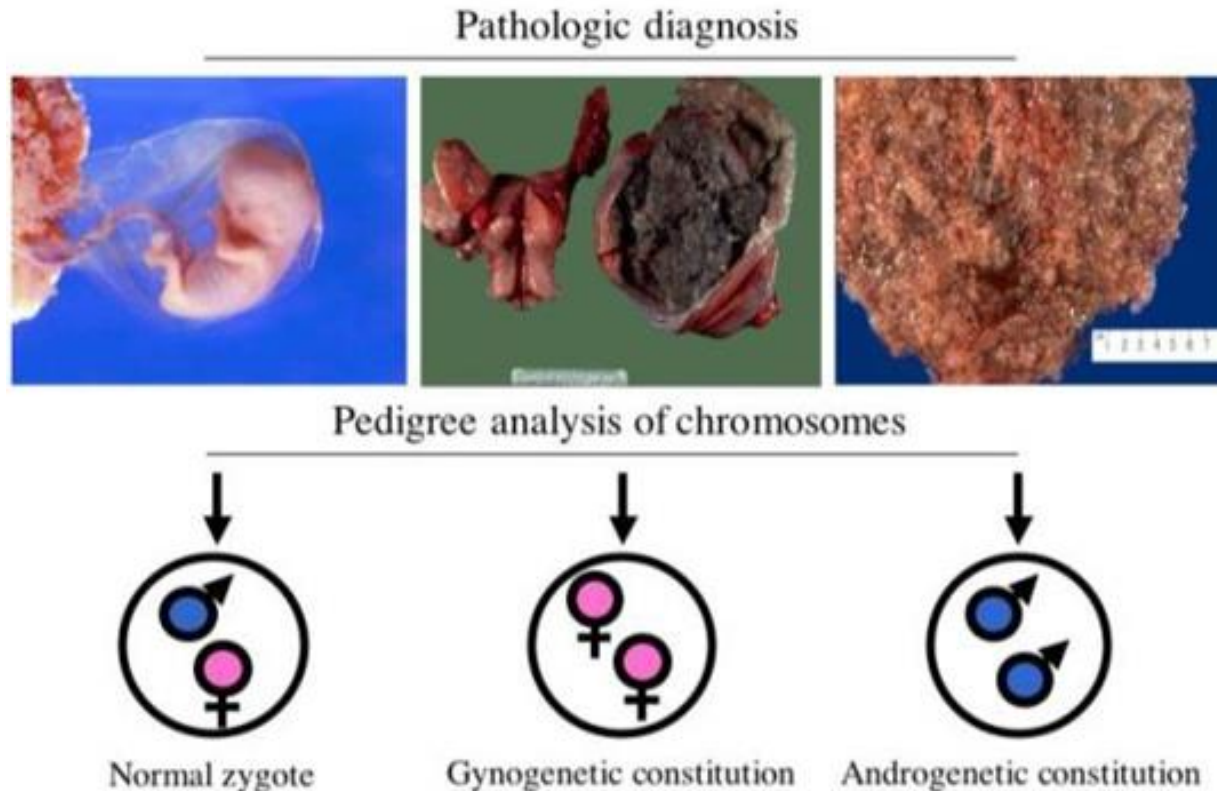
Differential gene expression



Most somatic cells have the same genome,
but different epigenomes.

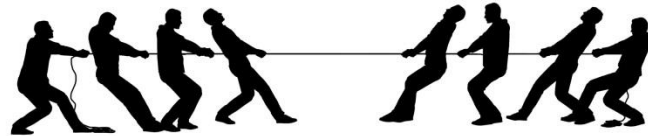
Each cell type has its unique epigenome.

Epigenetic phenomena: Maternal and paternal genomes carry different epigenomes that are **TOGETHER** required for normal development



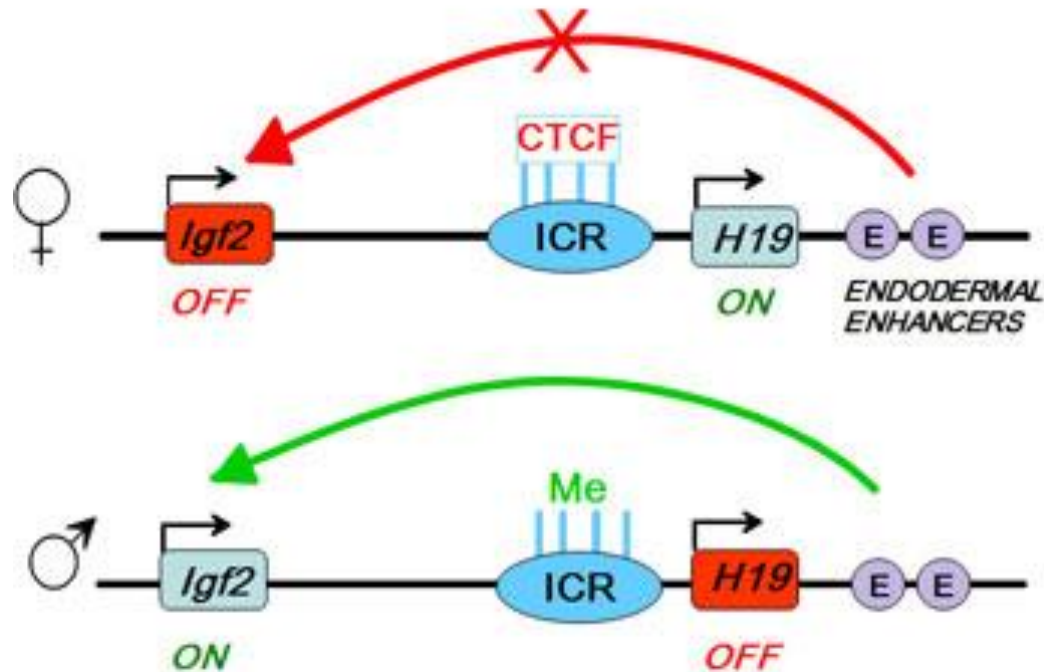
$N_m + N_p = 2N$ normal embryo
 $N_m + N_m = 2N$ abnormal cell mass
 $N_p + N_p = 2N$ abnormal cell mass

1.2. Reason for Imprinting – Haig's Tug-of-War hypothesis



- It is better for the father to have large offspring. These will outcompete the offspring from other fathers.
- It is better for the mother to have many small offspring so that they all survive.

Imprinting Growth Factor Genes



Germlines have their unique epigenetic markers

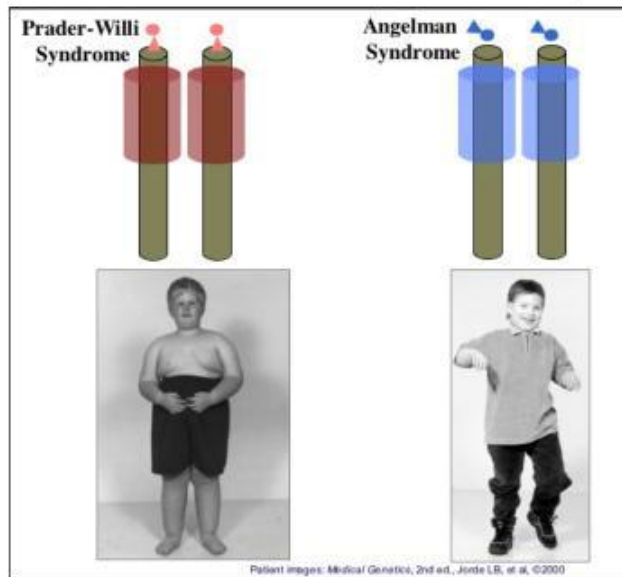
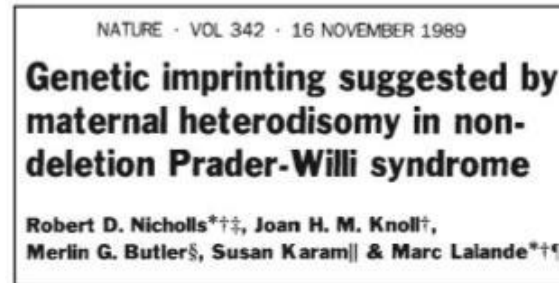
Eggs and sperm carry different epigenetic markers that are complementary in function in supporting the development of full term babies.

Human imprinting diseases

Prader-Willi Syndrome



- Hypotonia (low muscle tone, lack of sucking at birth)
- Small stature
- Small hands & feet
- Chronic hunger
- Late sexual maturity



Imprinting gene(s) in 15q11-13

Angelman Syndrome



- Neurological disorder
- Mental retardation
- Facial anomaly
- Muscular anomaly

Parent-of-origin effects/traits



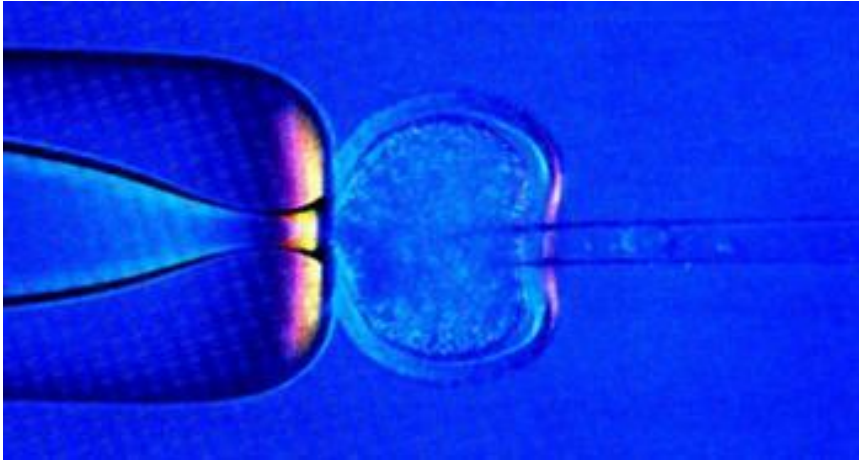
Liger – offspring of female tiger and male lion

Swimming – a tiger characteristics



Tigon – offspring of male tiger and female lion

Epigenetic phenomena: **Animal cloning errors**



Cloned animals are unhealthy. Some are obese, and some have lung defects, neurological problems and premature aging, etc.

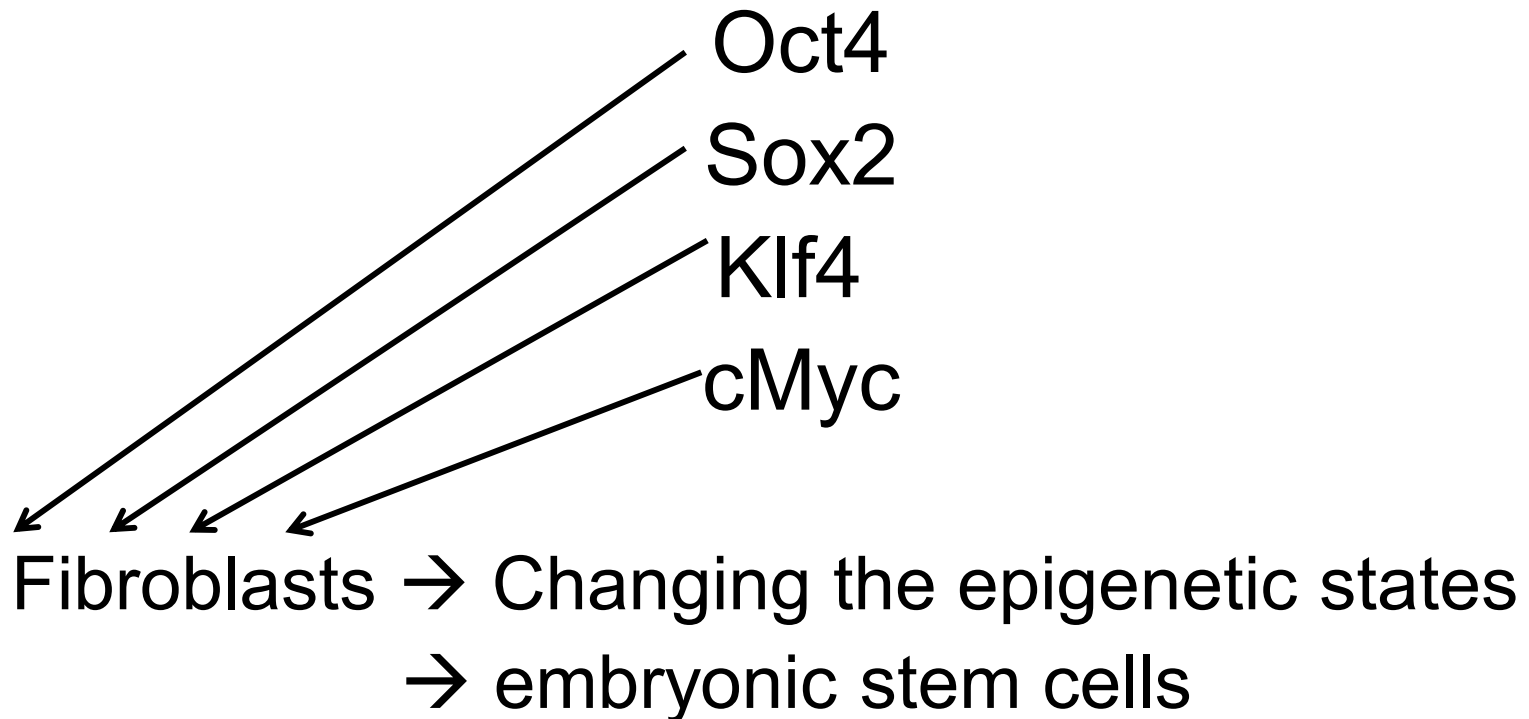
EPIGENETIC REPROGRAMMING:

Epigenetic information of a somatic nucleus/DNA has to be erased and replenished with those characteristic of the germ cells to ensure normal development. Otherwise, normal genes could be mis-regulated, thereby disrupting development or health.

iPS cells: induced pluripotent stem cells

EPIGENETIC REPROGRAMMING

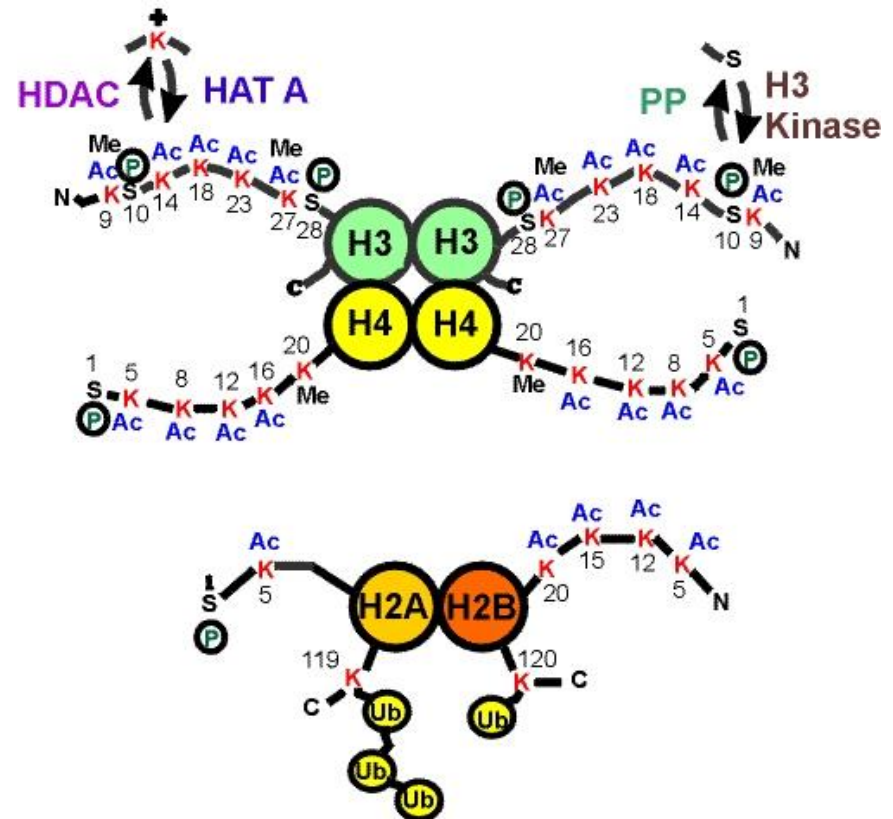
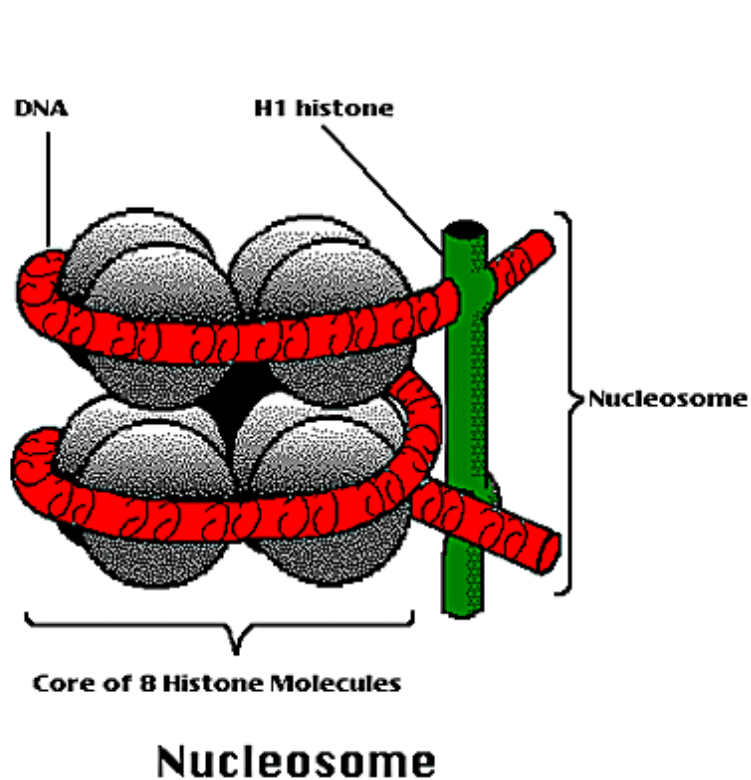
by four transcription factors:



What happened to Dolly

- Cloned from a nucleus of a 6-yr old sheep
- Had 20% shorter telomere length
- **Gave birth** to Bonnie and triplets in the old fashioned way (with normal telomere length)
- Died 14 Feb 2003 at 6 yr of age (**~ 1/2 of normal life span**)
- **Suffering from arthritis, lung disease, etc**
- Preserved and on display at the National Museum of Scotland.

Histone tails: Major sites of histone modifications



Histone tails: 16 – 25 aa
30% of histone mass
Invisible in crystal structures

Genetic code

	U	C	A	G
U	UUU } Phe UUC } UUA } Leu UUG }	UCU } Ser UCC } UCA } UCG }	UAU } Tyr UAC } UAA } Stop UAG }	UGU } Cys UGC } UGA } Stop UGG } Trp
C	CUU } Leu CUC } CUA } CUG }	CCU } Pro CCC } CCA } CCG }	CAU } His CAC } CAA } Gln CAG }	CGU } Arg CGC } CGA } CGG }
A	AUU } Ile AUC } AUA } AUG } Met	ACU } Thr ACC } ACA } ACG }	AAU } Asn AAC } AAA } Lys AAG }	AGU } Ser AGC } AGA } Arg AGG }
G	GUU } Val GUC } GUA } GUG }	GCU } Ala GCC } GCA } GCG }	GAU } Asp GAC } GAA } Glu GAG }	GGU } Gly GGC } GGA } GGG }

Histone code or epigenetic info

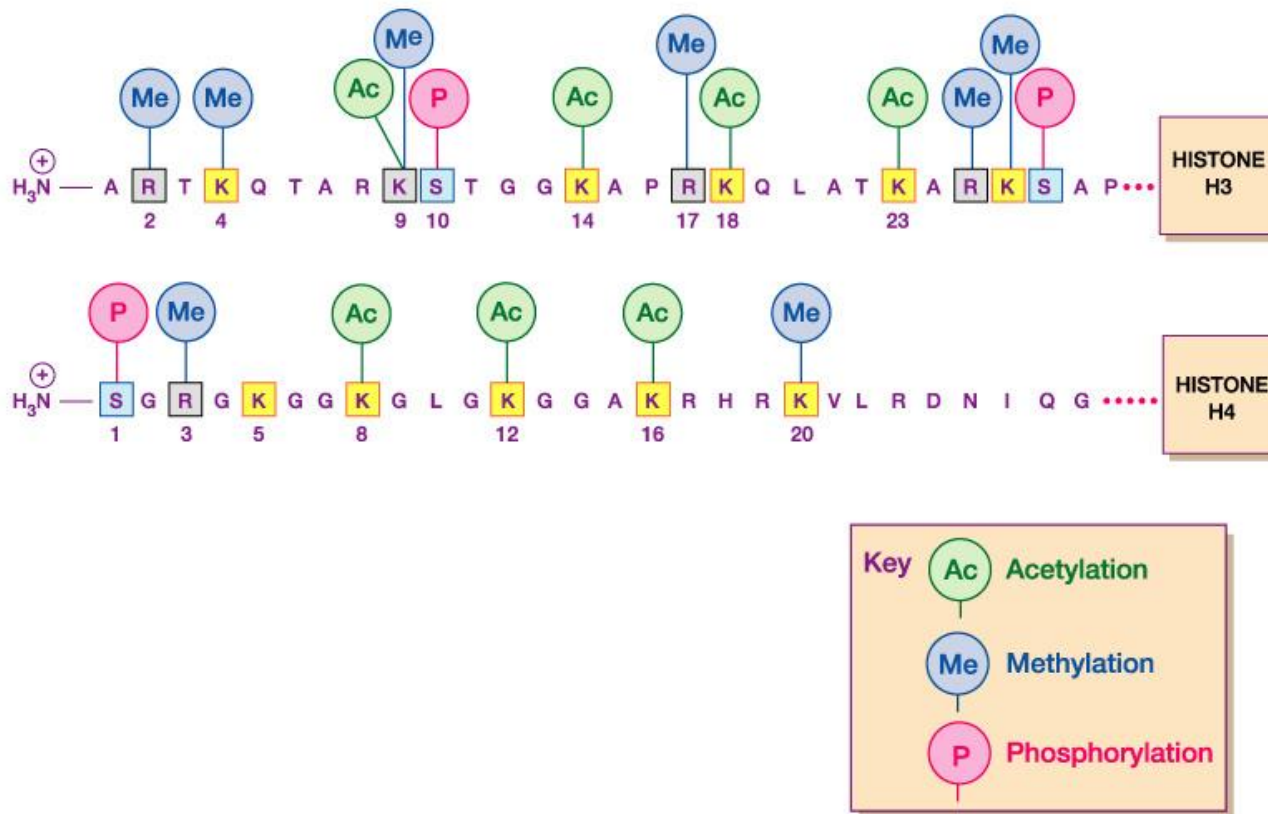


Figure 10-1 Human Molecular Genetics, 3/e. (© Garland Science 2004)

Context dependent meanings

Two key points to remember

- Understand what **heterochromatin** is and how DNA methylation Histone modification affects the heterochromatin
- Understand how imprinting involves parent of origin specific modifications in the DNA in such a manner that males want big offspring and females want little offspring – **tug of war.**