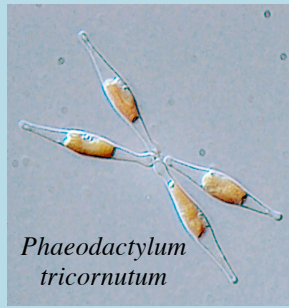
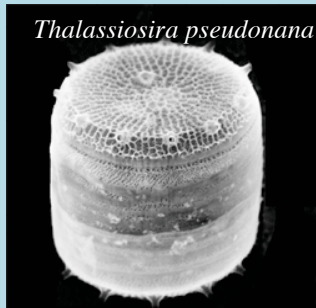


Epigenomics and its Connection to Evolution

To complement the previous
multicellular organism
epigenetic information

Model unicellular organisms

Diatoms



A diverse, widespread
phytoplankton.

Unicellular eukaryotic
plants
(Primary producers)

They evolved about 150-200
million years ago from a
eukaryotic cell that engulfed
bacteria similar to algae.

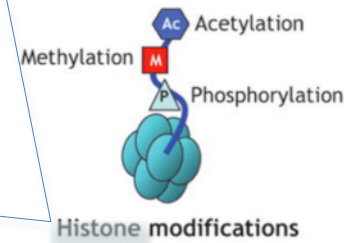
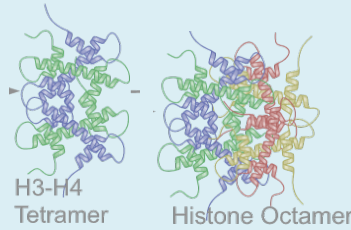
Epigenetic markers are **chemical modifications to DNA** that **do not** affect the sequence, but **do** affect the **expression** of genes.

They are generally **influenced by the environment** and are **inherited by offspring**.

DNA wraps around proteins called histones

- H2A, H2B, H3, H4
- Forms a nucleosome

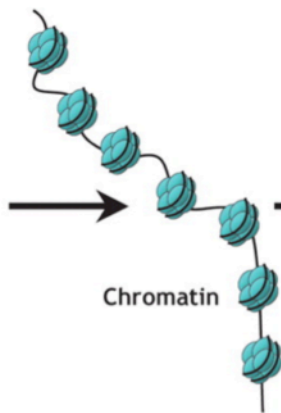
Archaea Eukaryota



Histone modifications

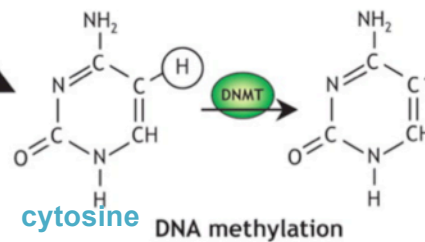


Chromosome

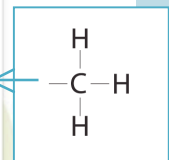


Chromatin

Nucleosome



cytosine DNA methylation



Epigenomic modifications can change
genomes throughout evolution because
organisms are able to modify gene
expression to **survive stressors** as well as
pass down enhanced fitness onto
descendants.

Comparing epigenetic markers can give
Insight into the gene modifications that aided
in **divergent evolutions and formation of
new species**.

