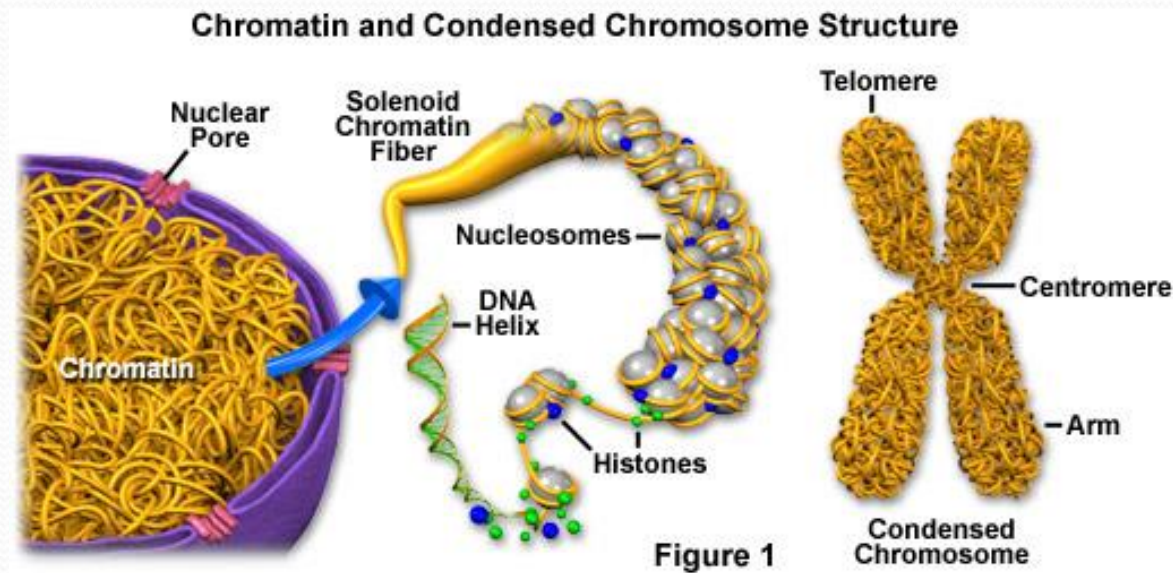


Zoo-342 Molecular biology
Lecture 3

**The eukaryotic chromosome and chromatin
organization**

The eukaryotic chromosome and chromatin organization



The eukaryotic chromosome:

- The eukaryotic chromosome is composed of both **double-stranded DNA** and **proteins**. The combination of DNA and protein is called **chromatin**.
- Two types of chromatins are formed:
 - 1) **Euchromatin**: loosely packaged (less condensed), can be transcribed.
 - 2) **Heterochromatin**: tightly packaged (highly condensed), not transcribed.

Chromatin organization:

- The total extended length of DNA in a single human cell would be nearly 2 m long if stretched from end to end.
- The problem is how to package this DNA inside the nucleus, which is only 2-4 μm (10^{-6} m) in diameter. Therefore, the primary function of chromatin is to package DNA into a smaller volume to fit in the cell.
- The basic structural unit of chromatin is the **nucleosome**.
- Nucleosomes consist of DNA wrapped around the protein core, which is composed of histone protein.
- The nucleosomes can be observed by placing interphase nuclei in a **hypotonic**, which releases the higher order packaging of the chromosomes.
- The major proteins of chromatin are the histones, basic amino acid (arginine and lysine) that facilitate binding to the negatively charged DNA molecule.

- There are five major types of histones called H1, H2A, H2B, H3 and H4 (Table 1).
- The structure of nucleosomes have been studied by treating chromatin with **DNase**, large amount of DNA are degraded, leaving only small particles composed of **DNA and proteins**.
- The DNA in these small DNA-protein complexes consists of fragments of about 210 bp (Figure 1). This DNA is protected from DNase digestion **due to its tight association with the proteins**.
- **Treatment** of the 210-bp DNA-protein complex with **micrococcal nuclease** removes additional DNA. At this point, **the protected DNA is 165 bp** in length.
- Harsh micrococcal nuclease treatment of these smaller DNA-protein complex releases of the basic **H1 histone** protein, and the DNA is digested down to a final size of approximately **145 bp** (Figure 1).
- The protein part of the core consisted of two molecules of each of four different histones: H2A, H2B, H3 and H4.

Class	Amino Acids	Number of Amino Acids	Percentage of Basic Amino Acids
H1	Very lysine-rich	213	30
H2A	Lysine-, arginine-rich	129	23
H2B	Moderately lysine-rich	125	24
H3	Arginine-rich	135	24
H4	Arginine-, glycine-rich	102	27

Table 1: Types and composition of histones

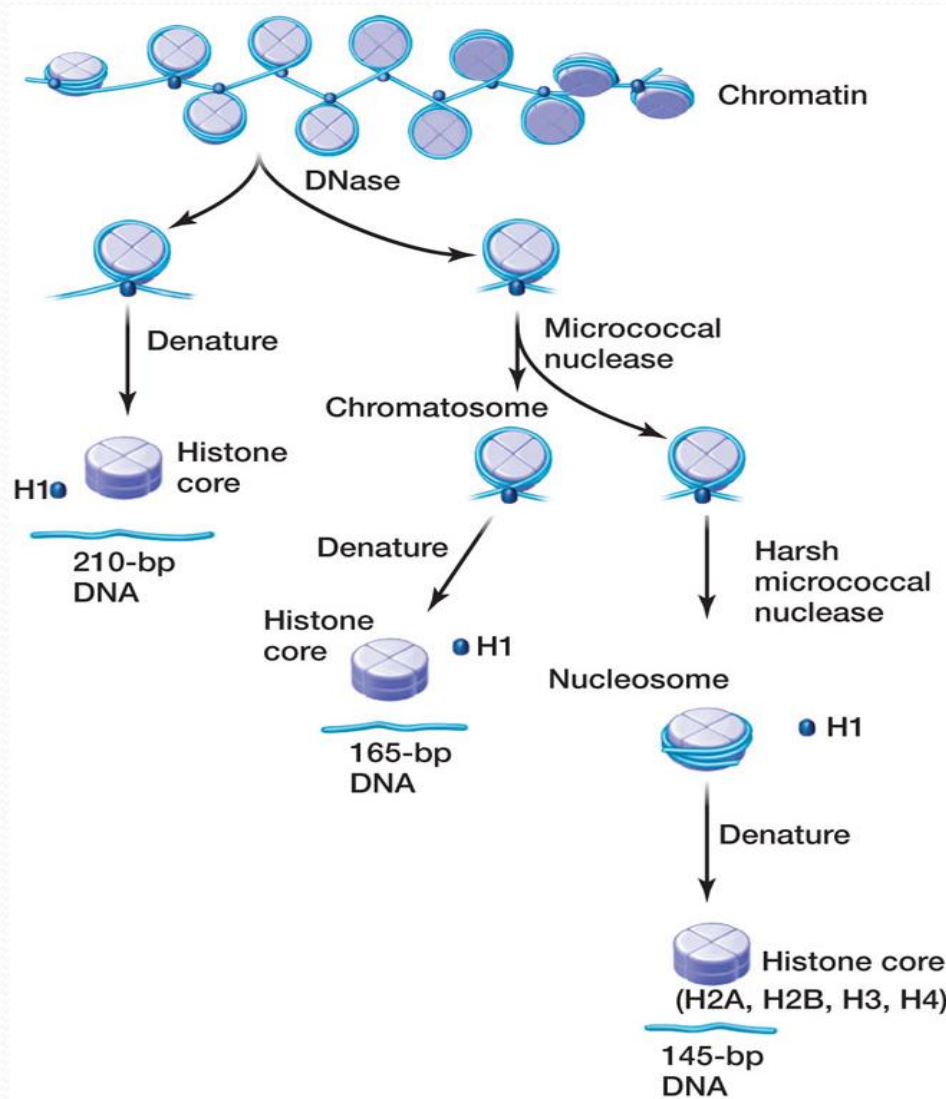


Figure 1: Studying the structure and composition of chromatin

- These data allow us to build a model for the organization of the nucleosome. Approximately 145 bp of genomic DNA is associated with the octamer of four histone proteins (8 proteins).
- The DNA located between the nucleosomes, which amounts to approximately 55-75 bp depending on the species, is termed the **linker DNA** (Figure 2).
- The **H1 histone** interacts with the nucleosome and some of the linker DNA as it enters and exits the nucleosome, which implies that it may serve to cross-link nucleosomes.
- The term **chromatosome** is defined as the core nucleosome plus the **H1** protein, a unit that includes approximately 165 bp of DNA.

- The binding of the histone core by the DNA is a major factor in controlling gene expression.
- The histones and nucleosome can be altered to regulate transcription in two major ways.
 - 1) Some histone proteins can be modified in particular, **phosphate groups**, **methyl groups**, and **acetyl groups** added and removed from specific amino acids in the H3 and H4 histone proteins. These modifications can lead to enhanced or repressed transcription.
 - 2) Histone cores can be removed from the DNA or moved to a different location on the DNA.

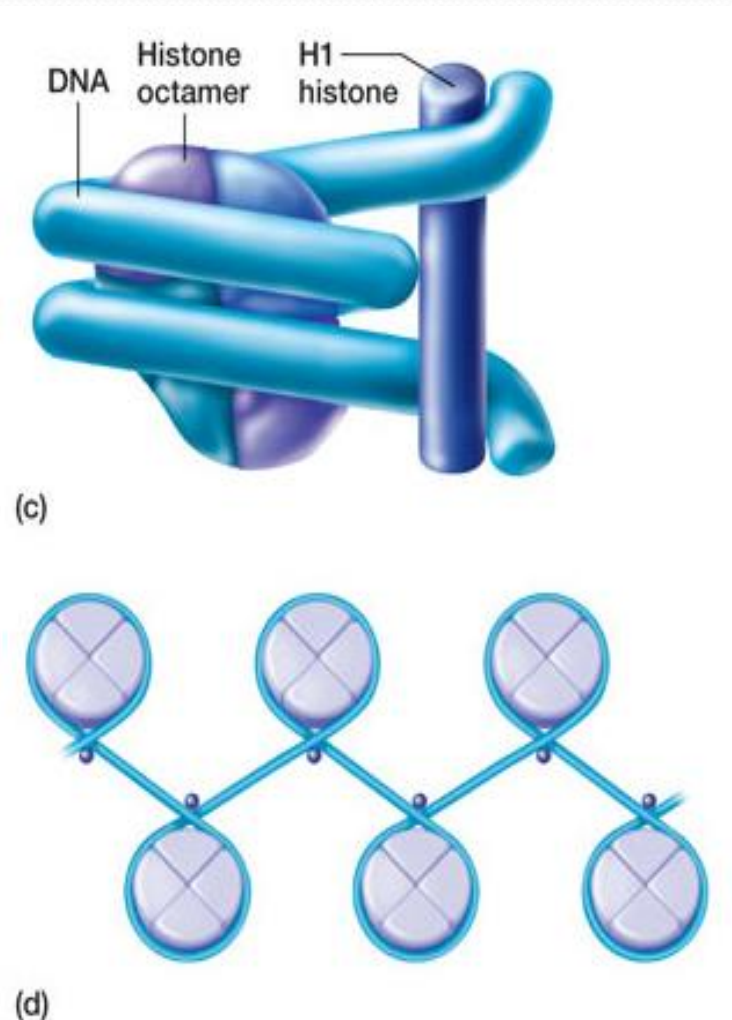
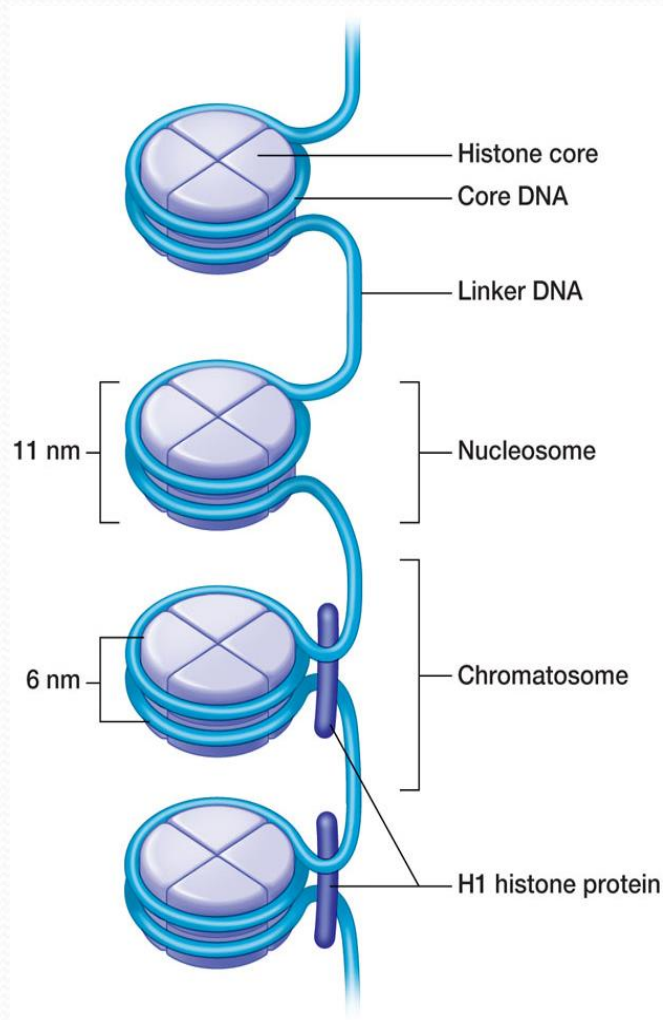


Figure 2: The eukaryotic chromosome is associated with histone proteins to forms nucleosome.