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CHAPTER 19 THE ORGANIZATION AND CONTROL OF EUKARYOTIC GENOMES

I. Introduction

- Gene expression in eukaryotes has two main differences from the same process in prokaryotes.
 - First, the typical multicellular eukaryotic genome is much larger than that of bacteria.
 - Second, cell specialization limits the expression of many genes to specific cells.
- The estimated 35,000 genes in the human genome includes an enormous amount of DNA that does not program the synthesis of DNA or proteins.
- This DNA is elaborately organized.
 - Not only is the DNA associated with proteins to form chromatin, but the chromatin is organized into higher organizational levels.
- Level of packing is one way that gene expression is regulated.
 - Densely packed areas are repressed.
 - Loosely packed areas are being actively transcribed.

II. Eukaryotic Chromatin Structure

- Chromatin structure is based on successive levels of DNA packing.
 - While the single circular chromosome of bacteria is coiled and looped in a complex, but orderly manner, eukaryotic chromatin is far more complex.
 - Eukaryotic DNA is precisely combined with large amounts of protein.
 - During interphase of the cell cycle, chromatin fibers are usually highly extended within the nucleus.
 - Eukaryotic chromosomes contain an enormous amount of DNA relative to their condensed length.
 - Each human chromosome averages about 2×10^8 nucleotide pairs.
 - If stretched out, DNA molecules would be about 1 m long. Thousands of times longer than the cell diameter.
 - The chromosome and all other human chromosomes fit into the nucleus.
 - The success through an elaborate, multistep system of DNA packing.
- Histone proteins are responsible for the first level of DNA packaging.
 - They package chromatin into nucleosomes, which are the basic units of chromatin.
 - The five types of histones are very similar from one eukaryote to another and are even present in bacteria.
 - Linker histon has the appearance of beads on a string, a **nucleosome**, in which DNA winds around a core of histone proteins.
 - The beaded string seems to remain essentially intact throughout the cell cycle.
 - Histones leave the DNA only transiently during DNA replication.
 - They stay with the DNA during transcription.
 - By changing shape and position, nucleosomes allow RNA polymerase to move along the DNA.
 - As chromosomes enter mitosis the beaded string undergoes higher-order packing.
 - The beaded string coils to form the 30-nm chromatin fiber.
 - The fiber forms looped domains attached to a scaffold of nonhistone proteins.
 - In a mitotic chromosome, the looped domains coil and fold to produce the characteristic metaphase chromosome.
 - These packing steps are highly specific and precise with particular genes located in the same places.

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