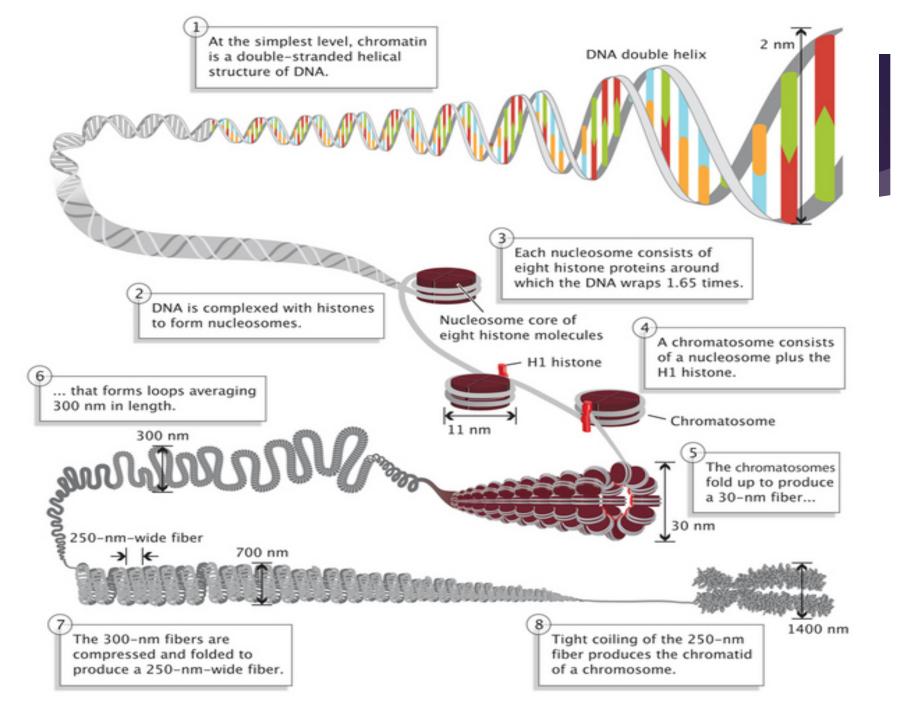
THE ORGANIZATION AND CONTROL OF EUKARYOTIC GENOMES

- BY ANANDI R



Genome includes

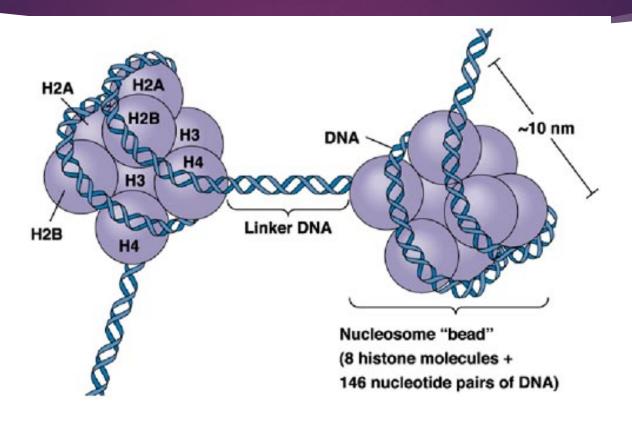
- Double stranded DNA
- Histone proteins
- Nucleosomes
- Chromatosomes
- Chromatids
- Chromosomes

Chromatin structure

- ► Eukaryotic DNA is precisely combined with large amounts of protein.
- During interphase, chromatin fibers are highly extended.
- ► If extended, each DNA molecule would be about 6 cm long

- ► First level Histone proteins
- ► Their positively charged amino acids bind tightly to negatively charged DNA.
- ► The five types of histones are very similar from one eukaryote to another .
- Unfolded chromatin has the appearance of beads on a string, a nucleosome, in which DNA winds around a core of histone proteins.

Nucleosomes



- ► 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-synthesizing polymerases to move along the DNA.

- ► **Level two** As chromosomes enter mitosis the beaded string coils to form the 30-*nm chromatin fiber*.
- ► Level three This fiber forms *looped domains* attached to a scaffold of non histone proteins.
- Level four the looped domains coil and fold to produce the characteristic metaphase chromosome.

- ► Interphase chromatin is generally much less condensed than the chromatin of mitosis with the 30-nm fibers and looped domains remaining intact.
- ► The chromatin of each chromosome occupies a restricted area within the interphase nucleus.
- Interphase chromosomes have areas that remain highly condensed, **heterochromatin**, and less compacted areas, **euchromatin**.

Assignment

Differentiate between heterochromatin and euchromatin.

Genome Organization at the DNA Level

- ► In eukaryotes, most of the DNA (about 97% in humans) does *not* code for protein or RNA.
 - 1. noncoding regions are regulatory sequences.
 - 2. introns.
 - 3. Repetitive DNA, present in many copies in the genome. (Three Types)

Types of Repeated DNA Sequences

1. Long terminal repeats (LTRs)

- Are identical sequences of DNA that repeat hundreds or thousands of times found at either end of retrotransposons or proviral DNA formed by reverse transcription of retroviral RNA.
- They are used by viruses to insert their genetic material into the host genomes.

Types of Repeated DNA Sequences

2. Tandem repeats

occur in DNA when a pattern of one or more nucleotides is repeated and the repetitions are directly adjacent to each other.

- Satellite DNA typically found in centromeres and heterochromatin
- ► Minisatellite repeat units from about 10 to 60 base pairs, found in many places in the genome, including the centromeres
- Microsatellite repeat units of less than 10 base pairs; this includes telomeres, which typically have 6 to 8 base pair repeat units

Types of Repeated DNA Sequences

3. Interspersed repeat

- These sequences propagate themselves by RNA mediated transposition, they have been called retrotransposons.
- Some types of interspersed repetitive DNA elements allow new genes to evolve by uncoupling similar DNA sequences from gene conversion during meiosis.
- ► Types : Transposable elements, DNA transposons, retrotransposons