

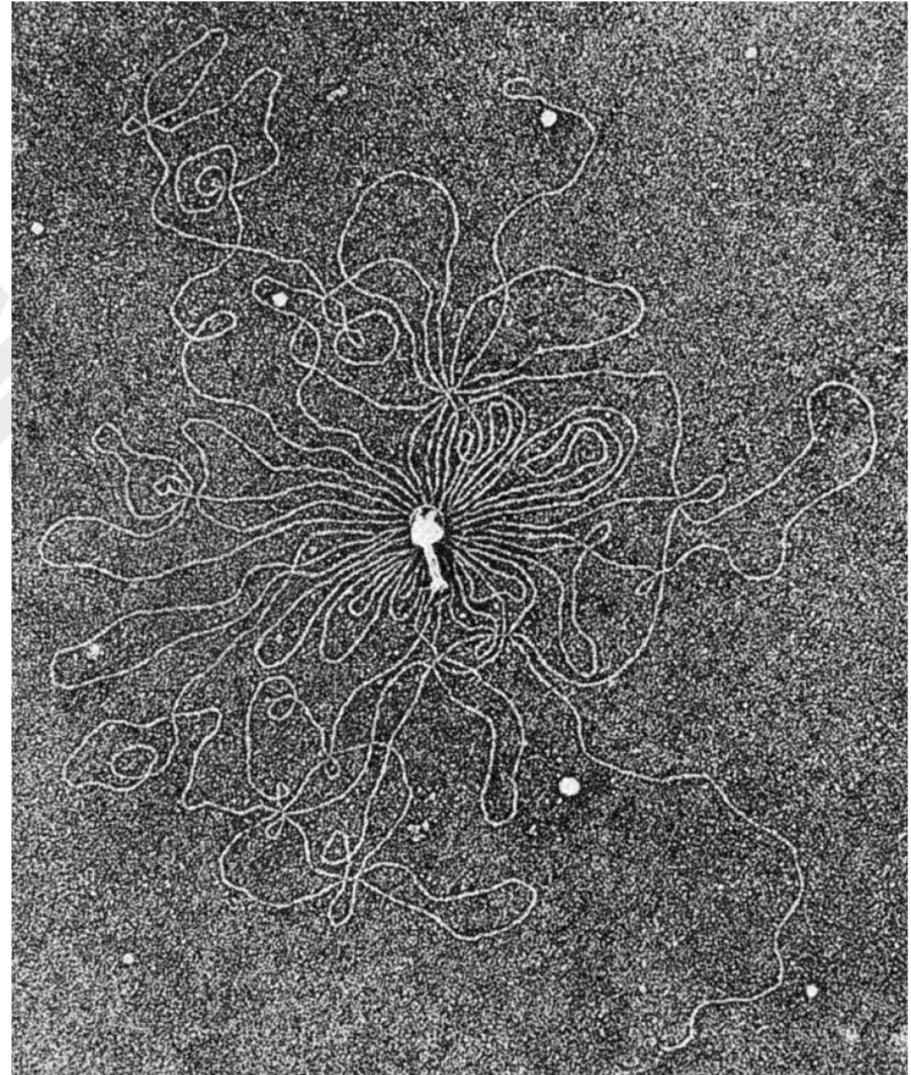
# 24 Genes and Chromosomes

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# Chromosomes

- **chromosomes** = tertiary packaging of DNA
  - serve as repositories of genetic information



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# Principle 1 (1 of 4)

**Chromosomes include dedicated sequences that ensure their replication, transcription, packaging, and transmission from one generation to the next. They are more than a long stretch of protein-encoding genes.**

## Principle 2 (1 of 6)

**Chromosomes are large.** To constrain them in a small space can require multiple layers and multiple modes of tertiary structure.

## Principle 3 (1 of 7)

**Chromosomes in all cells are maintained in a state of torsional stress.** DNA is underwound relative to the stable B-form structure, facilitating both the packaging of DNA and access to the genetic information contained within it.

## **Principle 4** (1 of 7)

**Specialized proteins and RNAs maintain chromosome structure.** Topoisomerases control DNA underwinding. Histones, condensins, cohesins, and other DNA-binding proteins provide scaffolds to organize chromosome structure. Certain long, noncoding RNAs also play important roles in chromosome structure and function.

# 24.1 Chromosomal Elements

# Genes Are Segments of DNA That Code for Polypeptide Chains and RNAs

- **phenotype** = visible property
- classically, a gene was defined as a portion of a chromosome that determines or affects a single phenotype

# Molecular Definitions of a Gene

- **mutations** = alterations in DNA sequence
- experiments by Beadle and Tatum found that mutant fungal strains lacked one or another specific enzyme
- the **one gene-one enzyme** hypothesis = a gene is a segment of genetic material that determines, or codes for, one enzyme
  - later broadened to **one gene-one polypeptide** hypothesis

# The Modern Biochemical Definition of a Gene

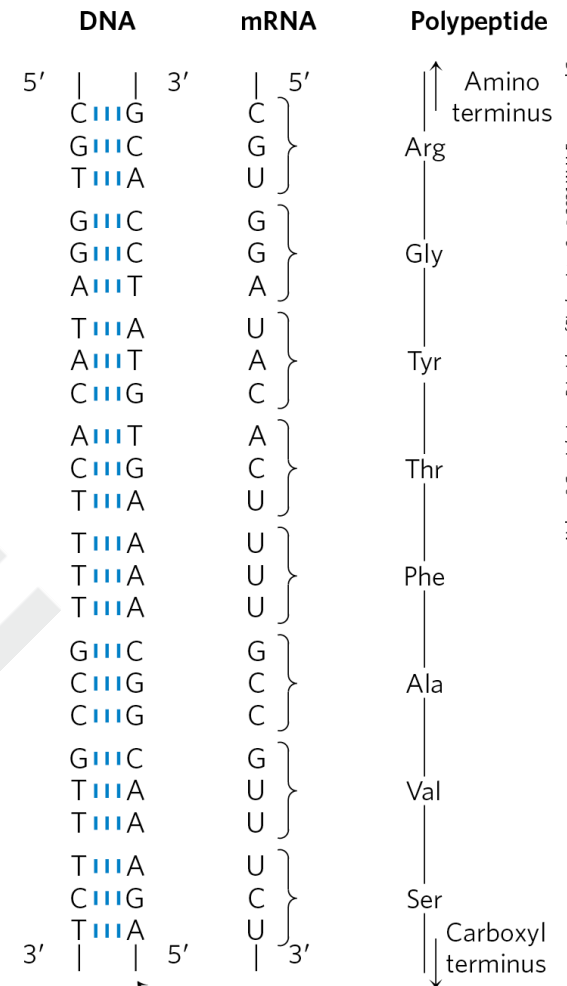
- **gene** = all the DNA that encodes the primary sequence of some final gene product (polypeptide or RNA with a structural or catalytic function)

# Regulatory Sequences

- **regulatory sequences** = segments of sequences of DNA that have a purely regulatory function
- functions of regulatory sequences:
  - denote the beginning or the end of genes
  - influence the transcription of genes
  - initiation points for replication or recombination

# Colinearity of DNA, mRNA, and Protein

- each amino acid of a polypeptide chain is coded for by three consecutive nucleotides in a single strand of DNA (“codon”)
- a polypeptide chain of 350 amino acid residues (an average-size chain) corresponds to 1,050 base pairs (bp) of coding DNA



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# Number of Genes in a Single Chromosome

- the *Escherichia coli* chromosome is a circular DNA molecule with 4,641,652 bp
  - encodes ~4,300 genes for proteins and >200 genes for structural or catalytic RNAs
- the human genome is ~3.1 billion bp
  - encodes approximately 20,000 genes across 24 different chromosomes

# DNA Molecules Are Much Longer than the Cellular or Viral Packages That Contain Them

- chromosomal DNAs are many orders of magnitude longer than the cells or viruses in which they are located

**Table 24-1 The Sizes of DNA and Viral Particles for Some Bacteria Viruses (Bacteriophages)**

Virus	Size of viral DNA (bp)	Length of viral DNA (nm)	Long dimension of viral particles (nm)
$\phi$ X174	5,386	1,939	25
T7	39,936	14,377	78
$\lambda$ (lambda)	48,502	17,460	190
T4	168,889	60,800	210

# Viruses

- viruses = infectious parasites that use resources of a host cell to carry out processes required to propagate
  - many consist of just a single RNA or DNA molecule and a protein coat

## Principle 2 (2 of 6)

**Chromosomes are large.** To constrain them in a small space can require multiple layers and multiple modes of tertiary structure.

# RNA and DNA Viruses

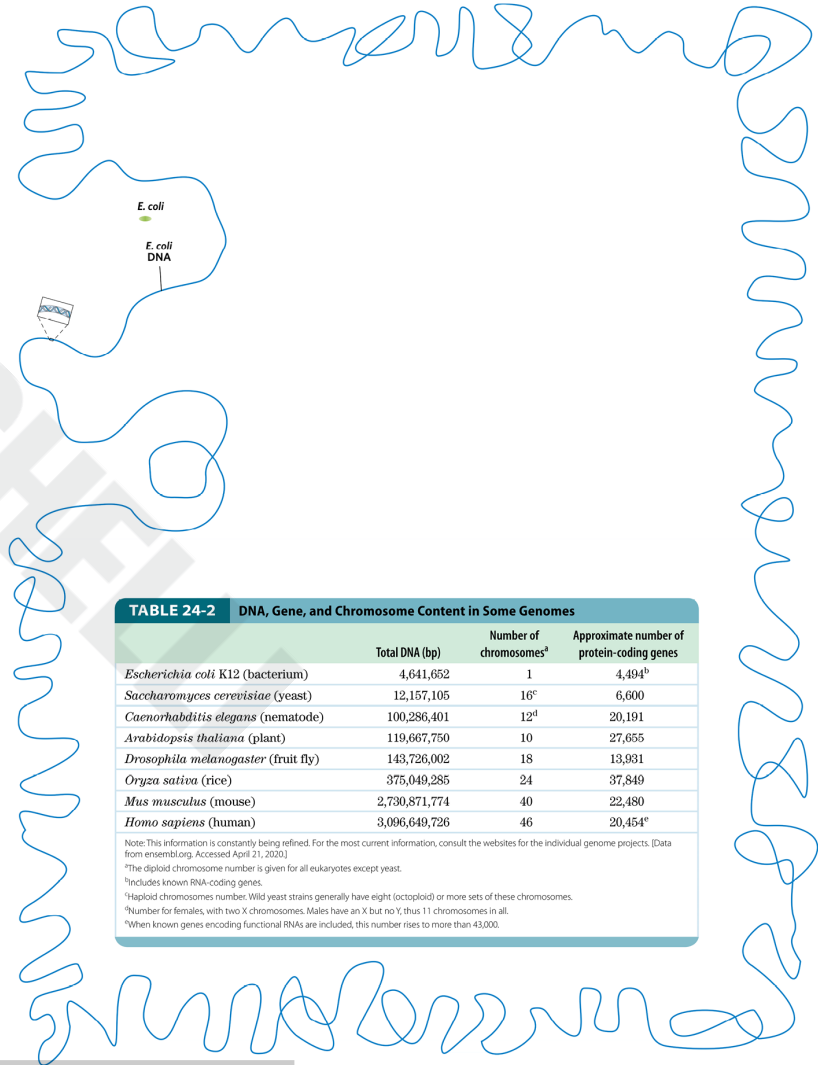
- almost all plant viruses and some bacterial and animal viruses have RNA genomes
  - tend to be particularly small
- DNA viral genomes vary greatly in size and tend to be circular for at least part of their life
- the contour lengths of their DNA are typically hundreds of times longer than the long dimensions of the viral particles that contain them

# Replicative Forms Appear During Viral Replication

- **replicative forms** = specific types of viral DNA that appear during viral replication within a host cell
- examples of replicative forms:
  - many linear DNAs become circular
  - all single-stranded DNAs become double-stranded

# Bacteria

- an *E. coli* cell contains ~100x as much DNA as a bacteriophage  $\lambda$  particle
- the *E. coli* chromosome is a single, double-stranded circular DNA molecule of 4,641,652 bp
  - the genome is ~850 times the length of the cell



**TABLE 24-2 DNA, Gene, and Chromosome Content in Some Genomes**

	Total DNA (bp)	Number of chromosomes <sup>a</sup>	Approximate number of protein-coding genes
<i>Escherichia coli</i> K12 (bacterium)	4,641,652	1	4,494 <sup>b</sup>
<i>Saccharomyces cerevisiae</i> (yeast)	12,157,105	16 <sup>c</sup>	6,600
<i>Caenorhabditis elegans</i> (nematode)	100,286,401	12 <sup>d</sup>	20,191
<i>Arabidopsis thaliana</i> (plant)	119,667,750	10	27,655
<i>Drosophila melanogaster</i> (fruit fly)	143,726,002	18	13,931
<i>Oryza sativa</i> (rice)	375,049,285	24	37,849
<i>Mus musculus</i> (mouse)	2,730,871,774	40	22,480
<i>Homo sapiens</i> (human)	3,096,649,726	46	20,454 <sup>e</sup>

Note: This information is constantly being refined. For the most current information, consult the websites for the individual genome projects. (Data from ensembl.org. Accessed April 21, 2020.)

<sup>a</sup>The diploid chromosome number is given for all eukaryotes except yeast.

<sup>b</sup>Includes known RNA-coding genes.

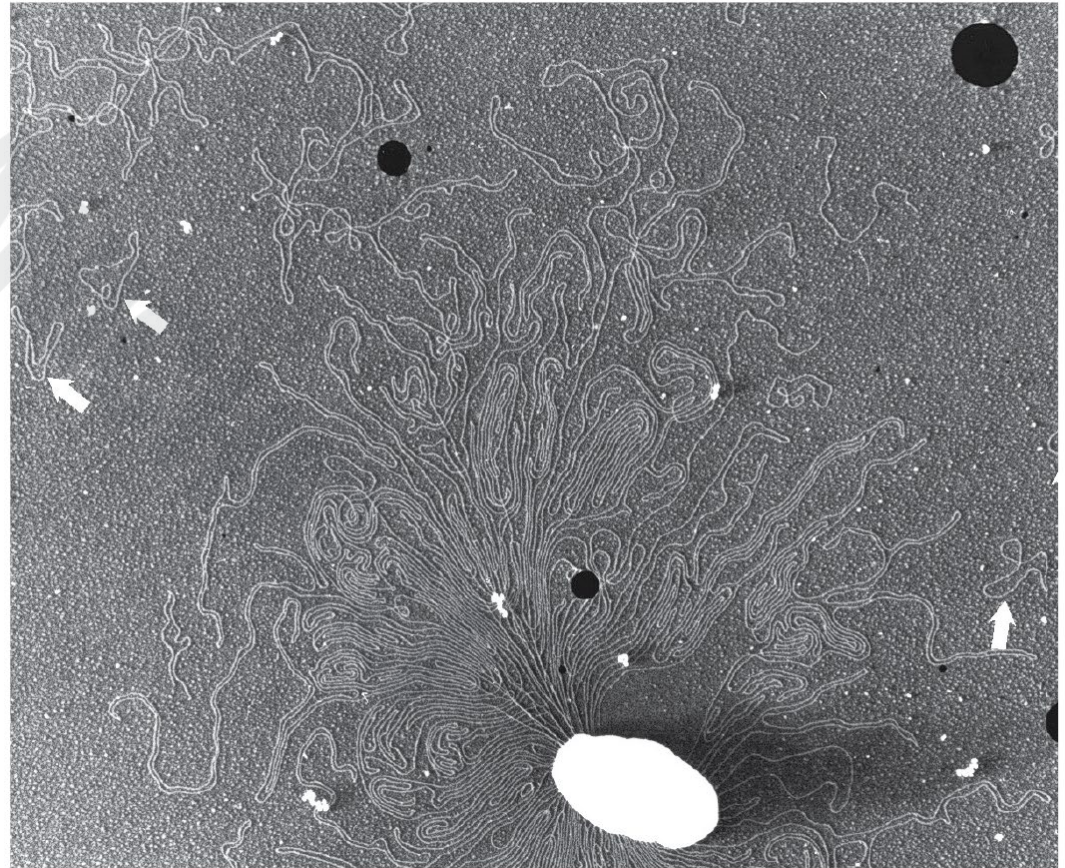
<sup>c</sup>Haploid chromosomes number. Wild yeast strains generally have eight (octoploid) or more sets of these chromosomes.

<sup>d</sup>Number for females, with two X chromosomes. Males have an X but no Y, thus 11 chromosomes in all.

<sup>e</sup>When known genes encoding functional RNAs are included, this number rises to more than 43,000.

# Bacterial Plasmids

- **plasmids** = small, circular DNA molecules that are in the cytosol of many bacteria
  - undergo replication
  - some carry useful genes, such as antibiotic resistance genes



# Eukaryotes

- a yeast cell contains 2.6x as much DNA as an *E. coli* cell.
- a *Drosophila* (fruit fly) cell contains >35x as much DNA as an *E. coli* cell
- a human cell contains ~700x as much DNA as an *E. coli* cell
  - cells of many plants and amphibians contain even more

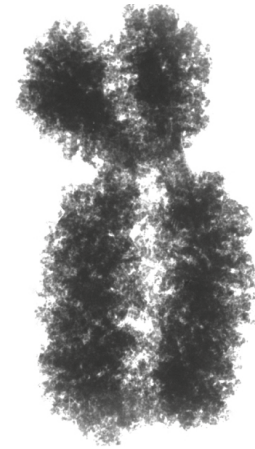
# DNA, Gene, and Chromosome Content in Some Genomes

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# Eukaryotic Chromosomes

- genetic material of eukaryotic cells is distributed into chromosomes
  - the diploid ( $2n$ ) number depends on the species
  - each chromosome contains a single, very large, duplex DNA molecule that carries a characteristic set of genes
- human somatic cells have 46 chromosomes



(a)

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(b)

## Principle 2 (3 of 6)

**Chromosomes are large.** To constrain them in a small space can require multiple layers and multiple modes of tertiary structure.

# The Length of Human DNA Molecules

- placed end to end, DNA molecules of one human genome would extend for about a meter
  - human cells are diploid, so each cell has 2 m of DNA
- the adult human body contains  $\sim 10^{14}$  cells and a total DNA length of  $2 \times 10^{11}$  km
  - greater than the the circumference of the earth ( $4 \times 10^4$  km)
  - greater than the distance between the earth and the sun ( $1.5 \times 10^8$  km)

# Mitochondria Contain DNA

- mitochondrial DNA (mtDNA) molecules are much smaller than nuclear chromosomes
  - codes for mitochondrial tRNAs and rRNAs and for a few mitochondrial proteins
- animal mtDNA contains <20,000 bp in a circular duplex
  - each mitochondrion has 2-10 copies
  - number of copies rises in some cells in an embryo during cell differentiation
- plant mtDNA ranges from 200,000 to 2,500,000 bp

# Chloroplasts Contain DNA

- chloroplast DNA (cpDNA) exists as circular duplexes from 120,000 to 160,000 bp

# Principle 1 (2 of 4)

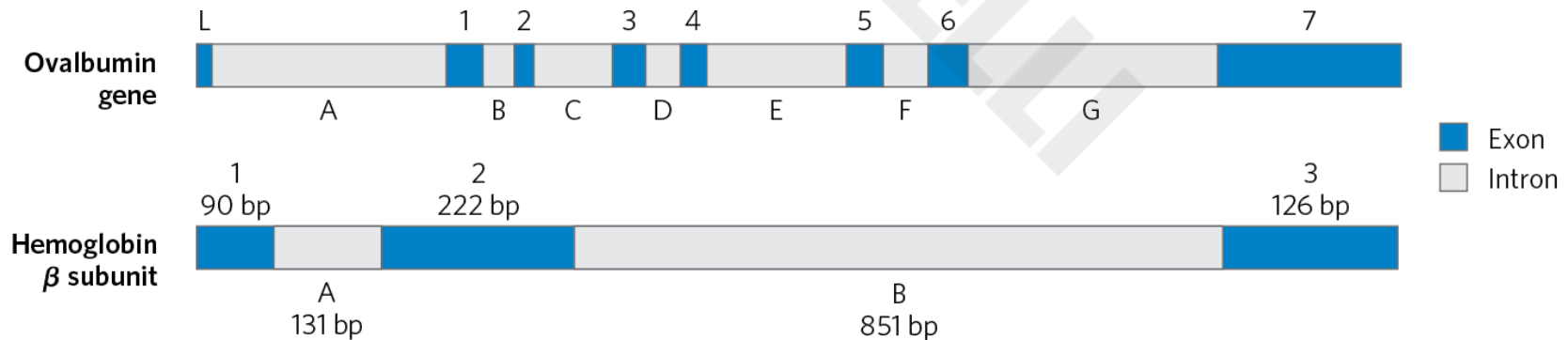
**Chromosomes include dedicated sequences that ensure their replication, transcription, packaging, and transmission from one generation to the next. They are more than a long stretch of protein-encoding genes.**

# Eukaryotic Genes and Chromosomes Are Very Complex

- **introns** = nontranslated, intervening DNA segments that do not code for the amino acid sequence of the polypeptide product
- **exons** = coding DNA segments
  - makes up only ~1.5% of human DNA
- the colinearity of the eukaryotic DNA and amino acid sequence is broken by introns
  - few bacterial genes contain introns

# Introns in Eukaryotic Genes

- in higher eukaryotes, the typical gene has more intron sequences than exon sequences
- together, introns and exons make up as much as 30% of the human genome



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# Highly Repetitive Sequences

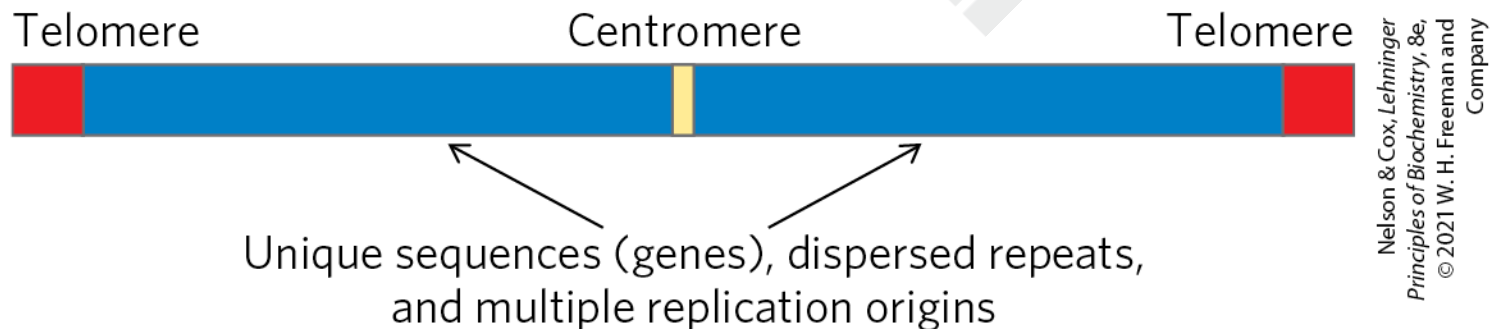
- **highly repetitive sequences (simple-sequence DNA or simple sequence repeats (SSR))** = short sequences, generally less than 10 bp long, that are sometimes repeated millions of times per cell
  - make up ~3% of the human genome
  - do not encode proteins or RNAs
  - associated with centromeres and telomeres
- **satellite DNA** = another name for simple-sequence DNA
  - named as such because it migrates as “satellite” bands in a cesium chloride density gradient

# Principle 1 (3 of 4)

**Chromosomes include dedicated sequences that ensure their replication, transcription, packaging, and transmission from one generation to the next. They are more than a long stretch of protein-encoding genes.**

# Centromeres

- **centromere** = a DNA sequence that functions during cell division as an attachment point for proteins that link the chromosome to the mitotic spindle
  - essential sequences in yeast are ~130 bp long and are very rich in A=T pairs
  - sequences of higher eukaryotes are longer and generally consist of thousands of tandem copies of one or several sequences of 5 to 10 bp

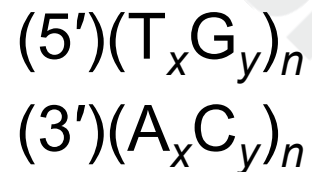


# Principle 1 (4 of 4)

**Chromosomes include dedicated sequences that ensure their replication, transcription, packaging, and transmission from one generation to the next. They are more than a long stretch of protein-encoding genes.**

# Telomeres

- **telomeres** = sequences at the ends of eukaryotic chromosomes that help stabilize the chromosome
  - shortened after each round of replication
  - end with multiple repeated sequences of the form



where  $x$  and  $y$  are generally between 1 and 4 and  $n$  is in the range of 20 to 100 for single-celled eukaryotes and  $>1,500$  in mammals

# Telomere Sequences

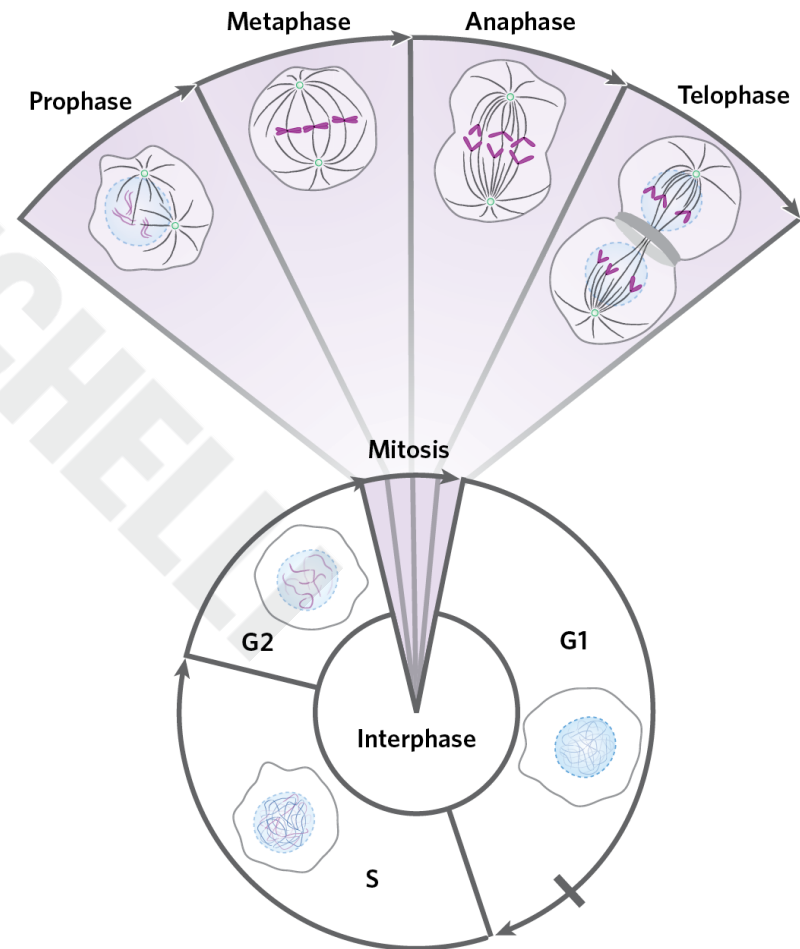
**Table 24-3 Telomere Sequences**

Organism	Telomere repeat sequence
<i>Homo sapiens</i> (human)	(TTAGGG) <sub>n</sub>
<i>Tetrahymena thermophile</i> (ciliated protozoan)	(TTGGGG) <sub>n</sub>
<i>Saccharomyces cerevisiae</i> (yeast)	(T(G) <sub>1-3</sub> (TG) <sub>2-3</sub> ) <sub>n</sub>
<i>Arabidopsis thaliana</i> (plant)	(TTTAGGG) <sub>n</sub>

# 24.3 The Structure of Chromosomes

# Chromatin Consists of DNA, Proteins, and RNA

- **chromatin** = eukaryotic chromosomal material composed of DNA, RNA, and proteins
  - amorphous in G<sub>0</sub> and interphase (G<sub>1</sub>, S, and G<sub>2</sub>) phases
- in the S phase of interphase, the amorphous DNA replicates to produce sister chromatids

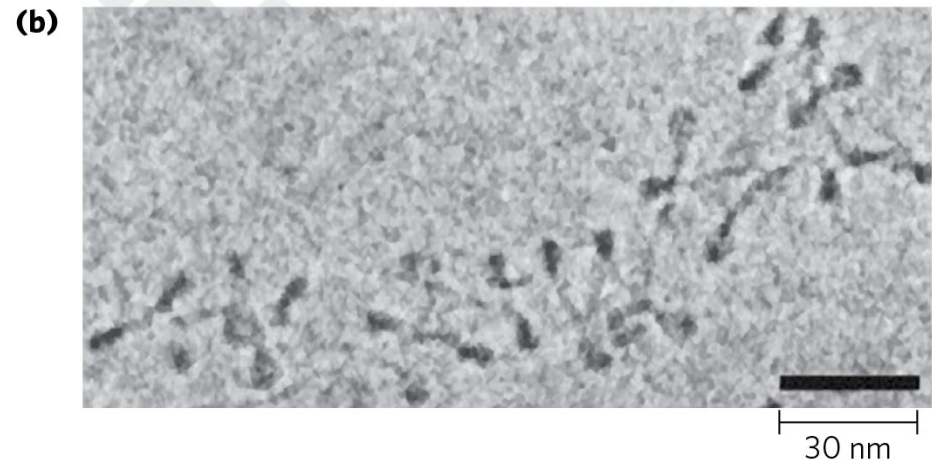
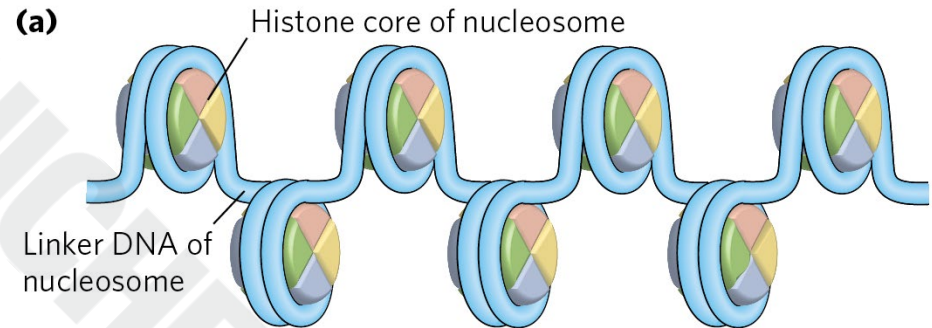


## Principle 4 (2 of 7)

**Specialized proteins and RNAs maintain chromosome structure.** Topoisomerases control DNA underwinding. Histones, condensins, cohesins, and other DNA-binding proteins provide scaffolds to organize chromosome structure. Certain long, noncoding RNAs also play important roles in chromosome structure and function.

# Histones Package and Order DNA into Nucleosomes

- **histones** = proteins that are tightly associated with chromatin and function to package and order the DNA
- **nucleosomes** = the fundamental structural unit of chromatin
  - composed of core histone proteins bound to DNA



# Histones Are Small, Basic Proteins

- all eukaryotic cells have five major classes of histones
  - H3 and H4 are nearly identical in all eukaryotes
  - H1, H2A, and H2B show less sequence similarity across eukaryotic species

**Table 24-5 Types and Properties of the Common Histones**

Histones	Molecular weight	Number of amino acid residues	Content of basic amino acids (% of total): Lys	Content of basic amino acids (% of total): Arg
H1 <sup>a</sup>	21,130	223	29.5	11.3
H2A <sup>a</sup>	13,960	129	10.9	19.3
H2B <sup>a</sup>	13,774	125	16.0	16.4
H3	15,273	135	19.6	13.3
H4	11,236	102	10.8	13.7

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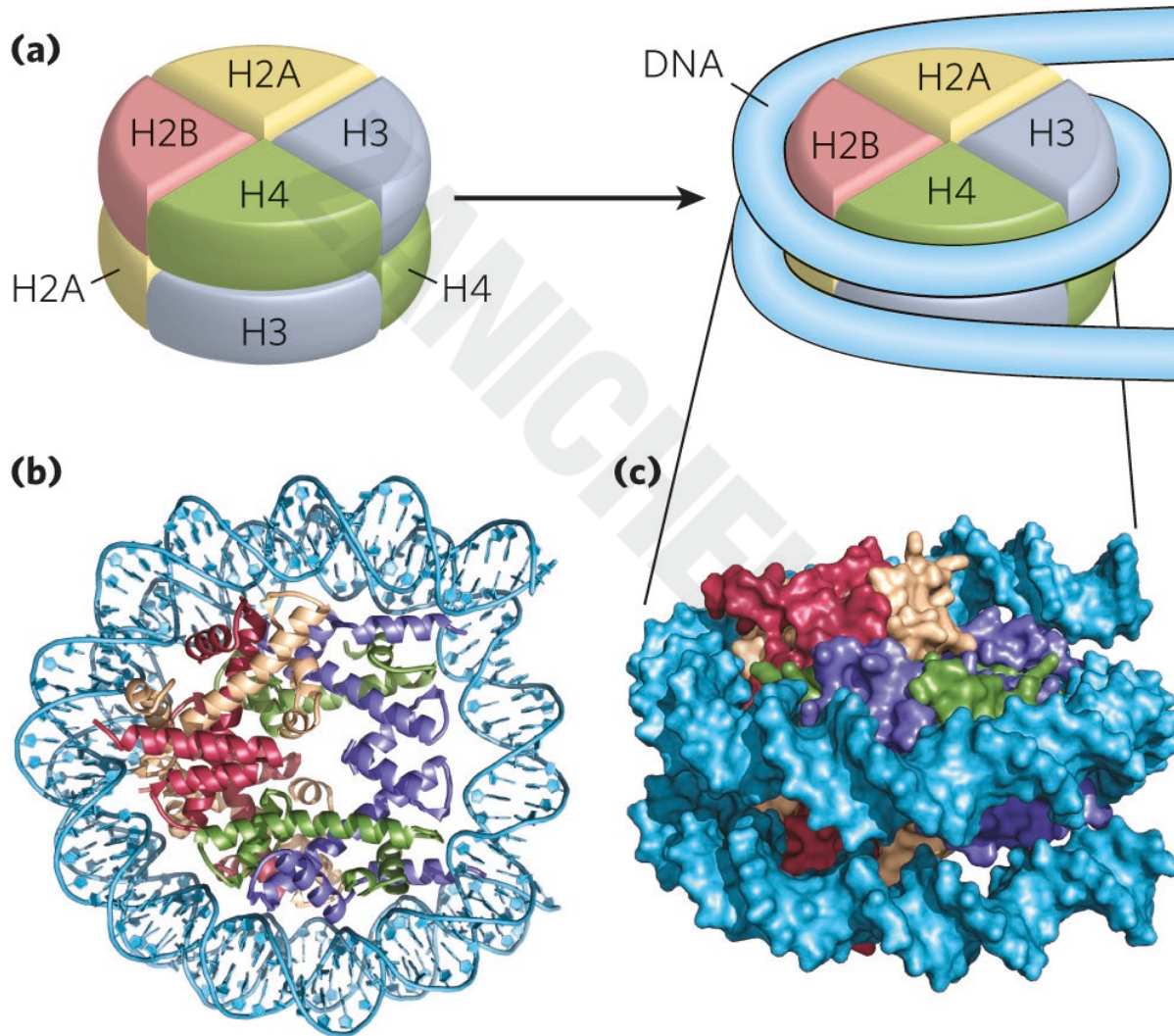
## **Principle 4** (3 of 7)

**Specialized proteins and RNAs maintain chromosome structure.** Topoisomerases control DNA underwinding. Histones, condensins, cohesins, and other DNA-binding proteins provide scaffolds to organize chromosome structure. Certain long, noncoding RNAs also play important roles in chromosome structure and function.

# Nucleosomes Are the Fundamental Organizational Units of Chromatin

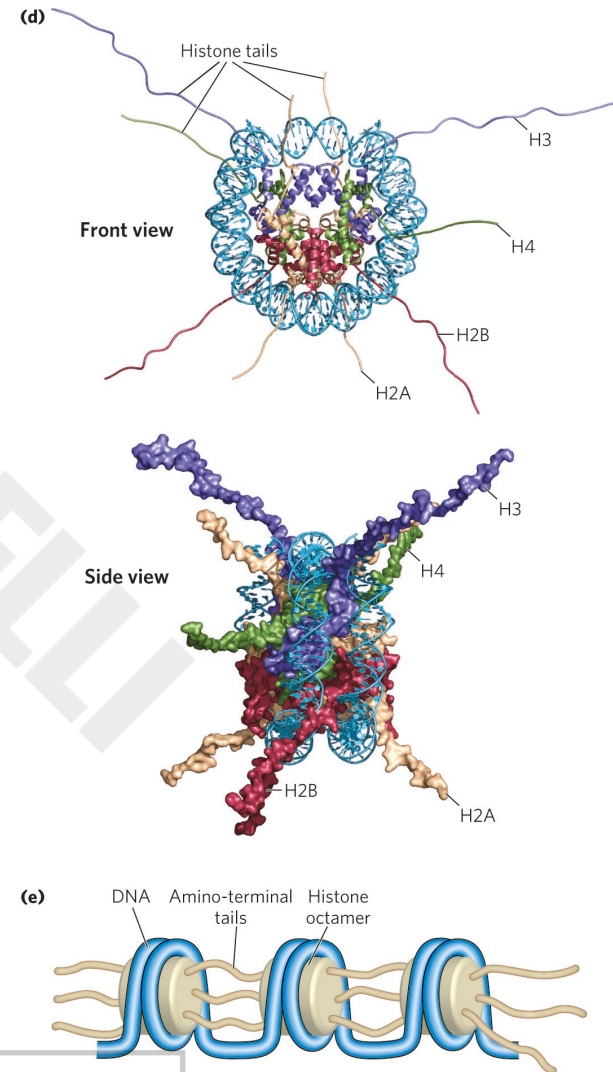
- chromosomes are arranged as “beads-on-a-string”
- the beads are complexes of histones and DNA
  - nucleosome contains eight histone molecules: two copies each of H2A, H2B, H3, and H4
  - contains ~200 bp of DNA, of which 146 bp are bound tightly around the histone core in a left-handed solenoidal supercoil
- the nucleosome is the bead plus the connecting DNA that leads to the next bead

# DNA Wrapped Around a Histone Core



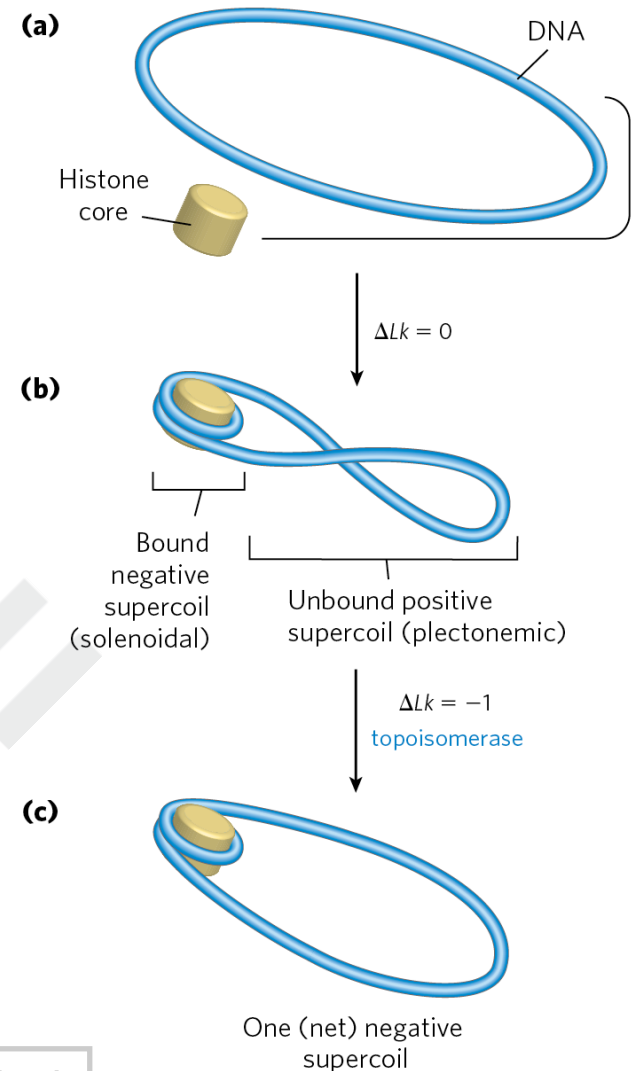
# Front and Side Views of Histone Amino-Terminal Tails

- the amino-terminal tails of the histones:
  - are intrinsically disordered
  - are where most of the histone modifications occur
  - play a key role in forming contacts between nucleosomes in the chromatin



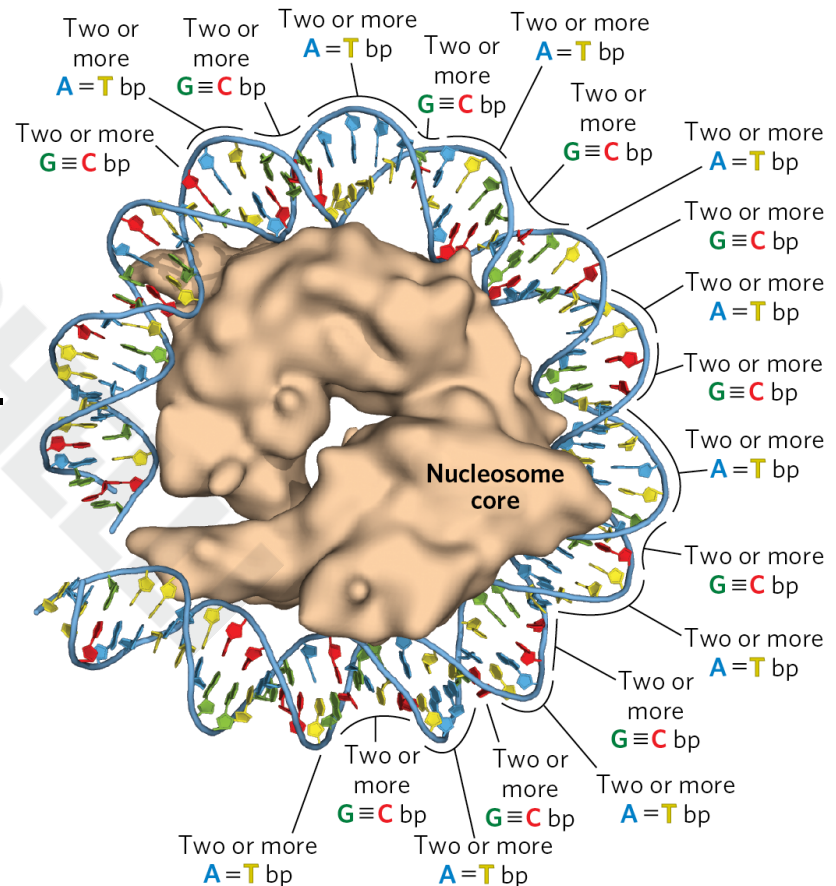
# Chromatin Assembly

- DNA wrapping around the histone core requires removal of  $\sim 1$  helical turn in the DNA
- binding of the histone core to a relaxed closed-circular DNA introduces a negative supercoil
  - accompanied by a positive supercoil in the unbound region
- when eukaryotic topoisomerases relax positive supercoils,  $Lk$  decreases



# The Effect of DNA Sequence on Nucleosome Binding

- histone binding is not random
- binding of the histone core seems to depend on a local abundance of A=T base pairs.
- staggering AA, AT, or TT dinucleotides at 10 bp intervals facilitates DNA binding around the histone core



# Depositing Nucleosome Cores on DNA

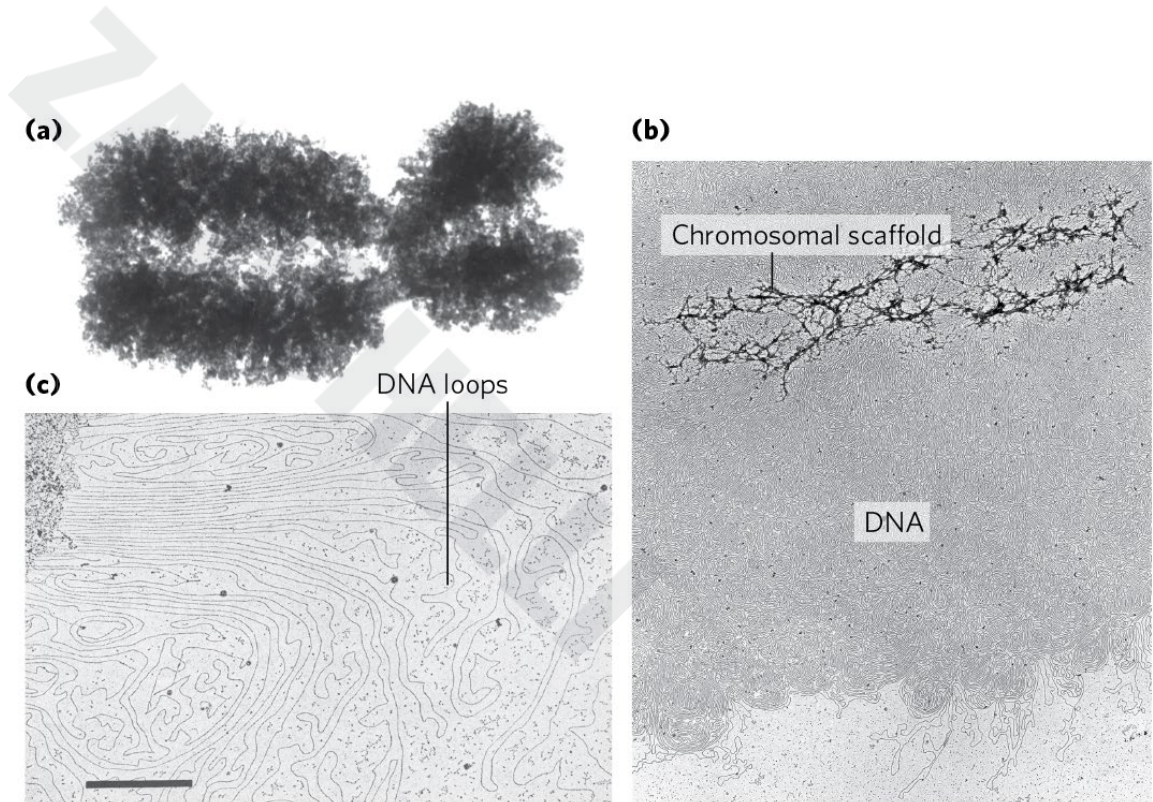
- deposited stepwise: a tetramer of two H3 and two H4 histones binds first, followed by two H2A–H2B dimers
- **histone chaperones** = mediate the incorporation of nucleosomes into chromosomes after chromosomal replication
  - conserved in all eukaryotes
- **histone exchange factors** = permit the substitution of histone variants for core histones in contexts other than postreplication

## **Principle 4** (4 of 7)

**Specialized proteins and RNAs maintain chromosome structure.** Topoisomerases control DNA underwinding. Histones, condensins, cohesins, and other DNA-binding proteins provide scaffolds to organize chromosome structure. Certain long, noncoding RNAs also play important roles in chromosome structure and function.

# Nucleosomes Are Packed into Highly Condensed Chromosome Structures

- certain regions of DNA seem to associate with a chromosomal scaffold



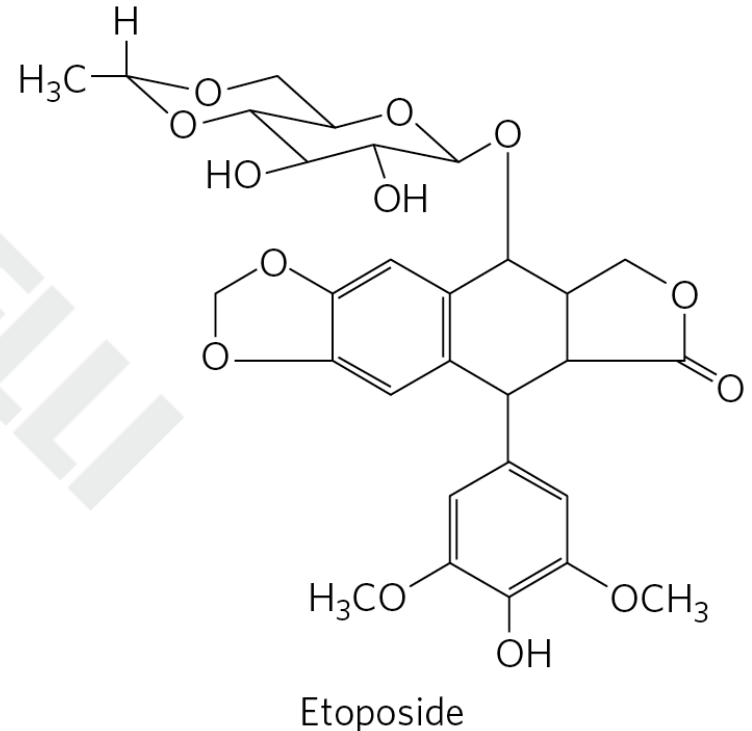
(a, b) Don W. Fawcett/Science Source. (c) U. K. Laemmli et al., "Metaphase chromosome structure: The role of nonhistone proteins," *Cold Spring Harb. Symp. Quant. Biol.* 42:351, 1978. © Cold Spring Harbor Laboratory Press.

## Principle 3 (6 of 7)

**Chromosomes in all cells are maintained in a state of torsional stress.** DNA is underwound relative to the stable B-form structure, facilitating both the packaging of DNA and access to the genetic information contained within it.

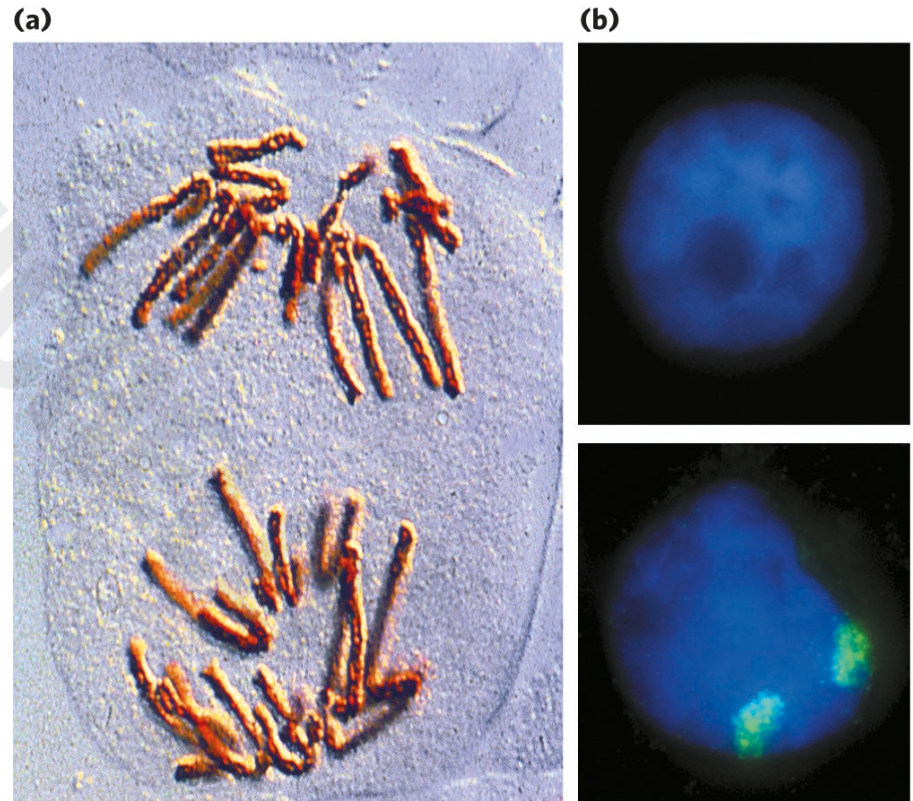
# Topoisomerase II Is One of the Most Abundant Proteins in the Chromosome

- topoisomerase II is important to the maintenance of chromatin structure
- inhibitors of topoisomerase II can kill rapidly dividing cells
  - several chemotherapy drugs are topoisomerase II inhibitors



# Chromosomal Organization in Eukaryotic Nucleus

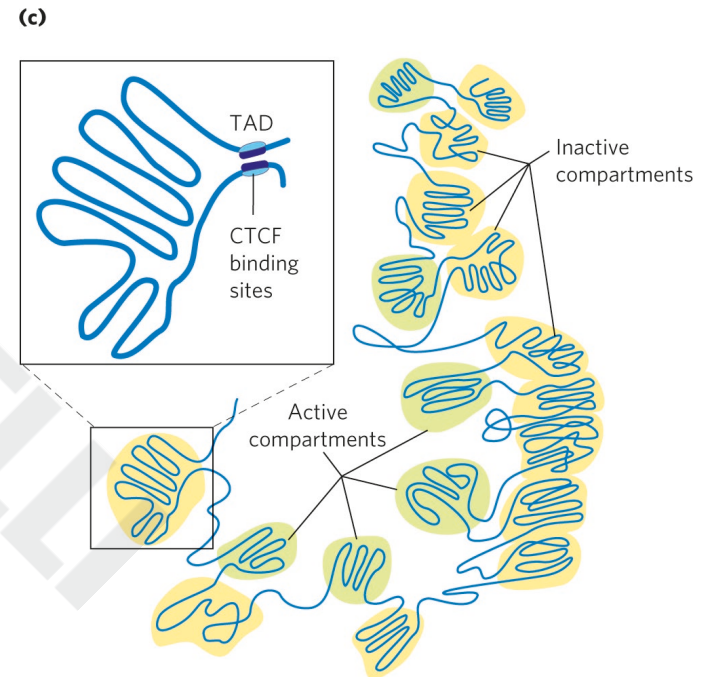
- chromosomes are highly condensed and organized just before cell division
- chromosomes appear dispersed during interphase
  - each chromosome is still organized, with two sets of compartments



(a) Pr. G. Giménez-Martín/Science Source. (b) Karen Meaburn and Tom Misteli/National Cancer Institute. Nelson & Cox, *Lehninger Principles of Biochemistry*, 8e, © 2021 W. H. Freeman and Company

# Active and Inactive Compartments

- active compartments have reduced chromatin condensation
- inactive compartments (**heterochromatin**) are highly condensed
- **topologically associating domains (TADs)** = large segments of DNA are organized in loops
  - binding of CTCF to bordering sites brings DNA together



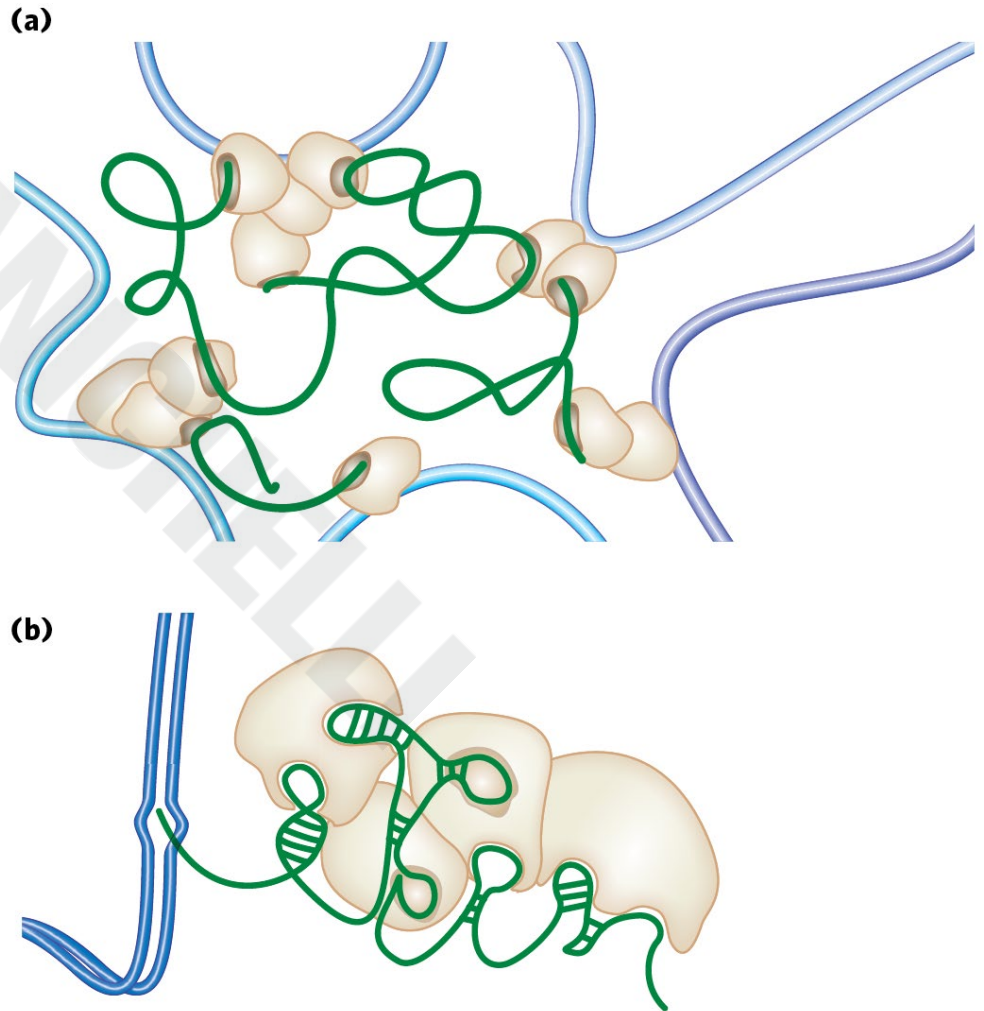
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## **Principle 4** (5 of 7)

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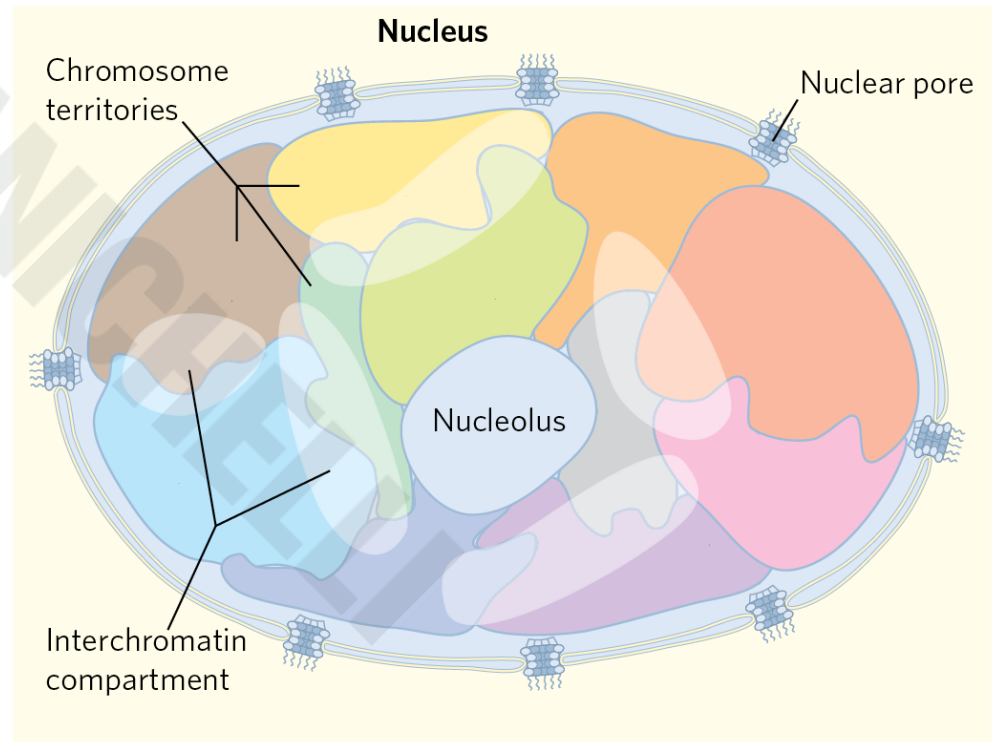
# Long Noncoding RNAs (lncRNAs)

- **long noncoding RNAs (lncRNAs)** = play a functional role in defining the chromosome structure
  - many provide a scaffold for proteins



# Chromosome Territory

- **chromosome territory** = subnuclear domain that constrains the entire structure of each chromosome
  - little or no intermingling of DNA in different territories

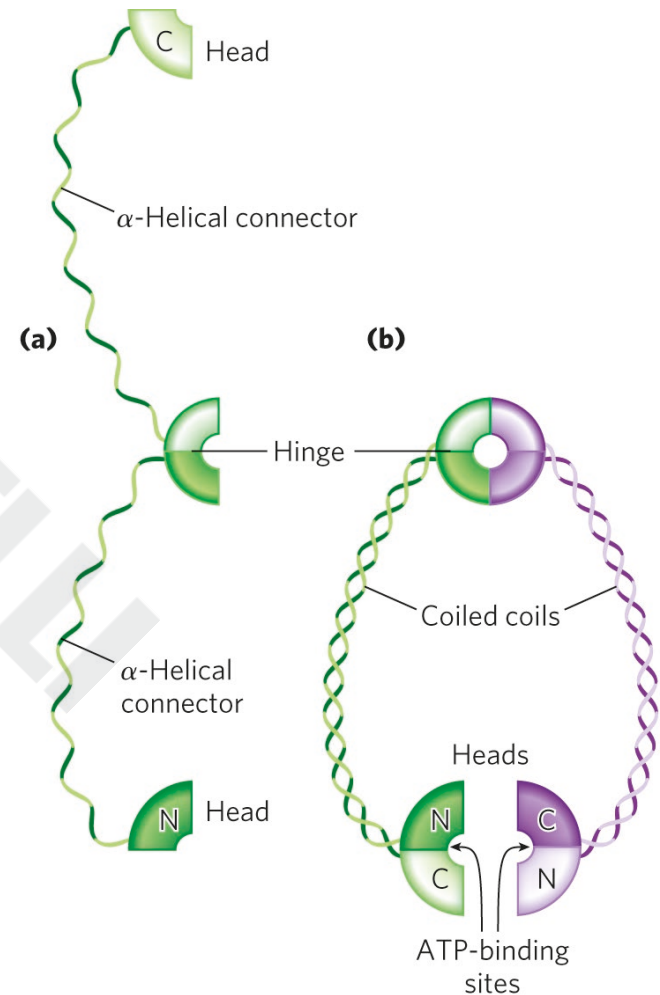


## **Principle 4** (6 of 7)

**Specialized proteins and RNAs maintain chromosome structure.** Topoisomerases control DNA underwinding. Histones, condensins, cohesins, and other DNA-binding proteins provide scaffolds to organize chromosome structure. Certain long, noncoding RNAs also play important roles in chromosome structure and function.

# Condensed Chromosome Structures Are Maintained by SMC Proteins

- **SMC proteins** (structural maintenance of chromosomes) = responsible for maintaining the structure and integrity of chromosomes following replication
  - consist of 5 distinct domains



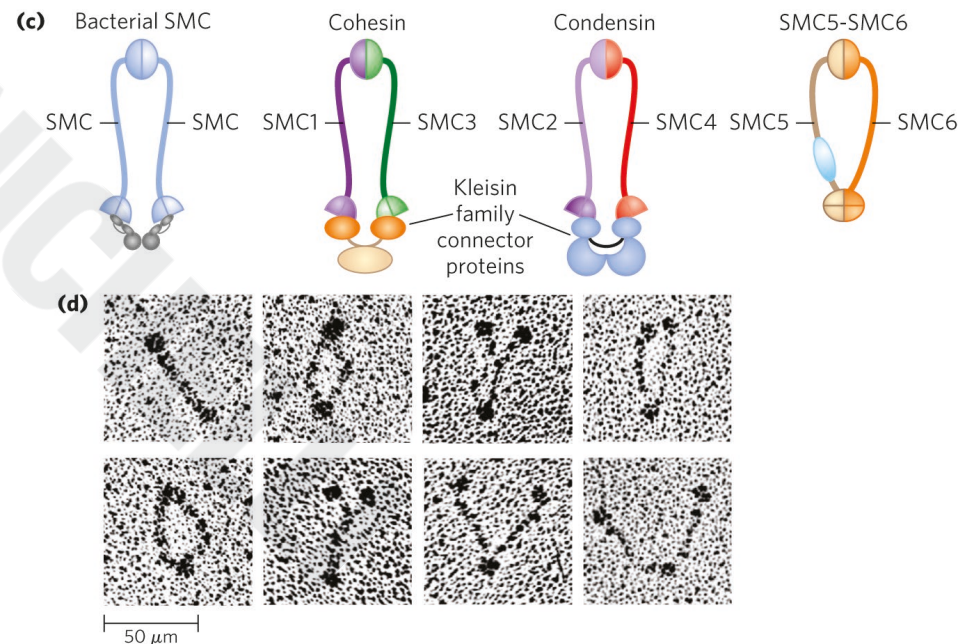
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# Eukaryotes Have Two Major Types of SMC Proteins

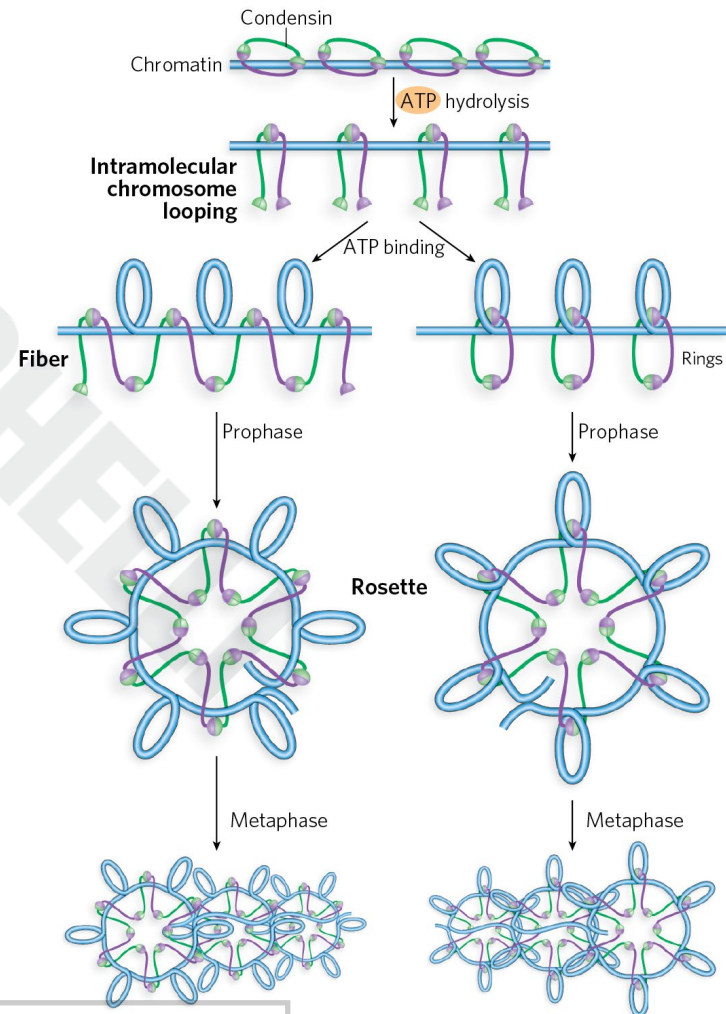
- **cohesins** = link sister chromatids together after replication and keep them together as the chromosomes condense to metaphase
- **condensins** = essential to chromosomal condensation as cells enter mitosis
  - create positive supercoils



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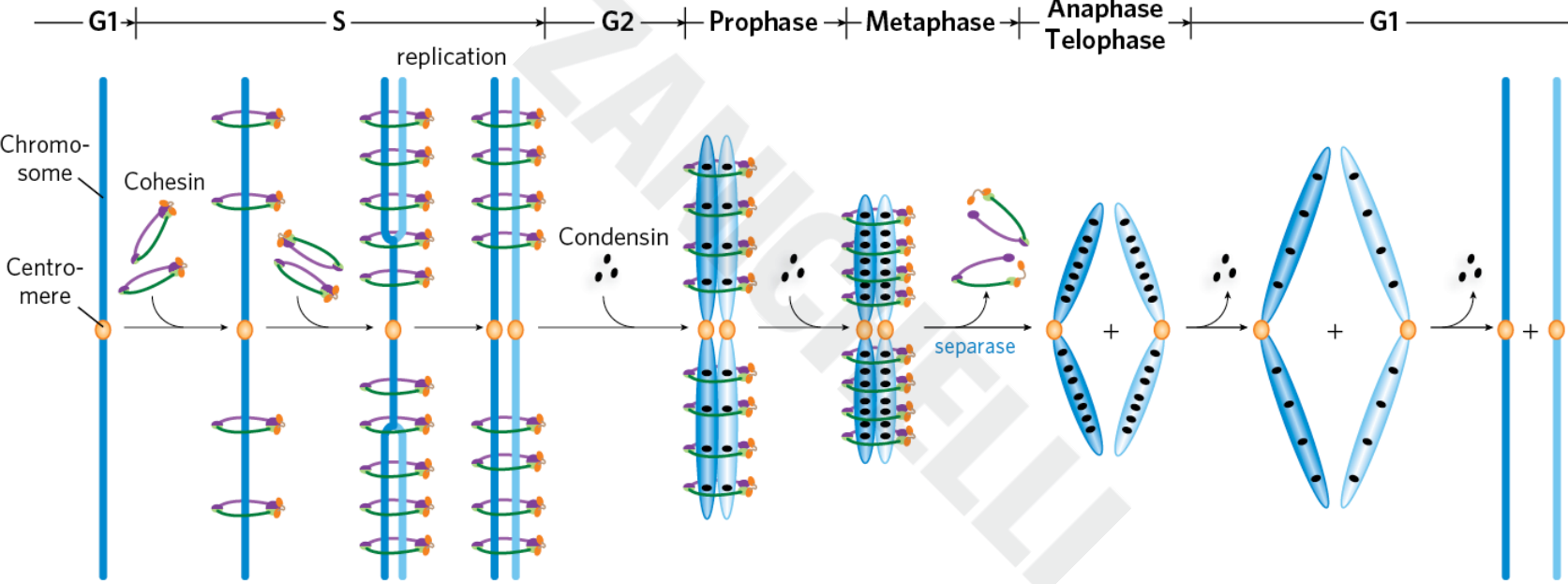
# Models of the Role of Condensins in Chromatin Condensation

- as DNA is compacted to form tighter and tighter loops, condensins stabilize the loops by binding at the base of each one



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# The Role of Cohesins and Condensins in the Cell Cycle



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## Principle 2 (6 of 6)

**Chromosomes are large.** To constrain them in a small space can require multiple layers and multiple modes of tertiary structure.

# Bacterial DNA Is Also Highly Organized

- bacterial DNA is compacted in a structure called the **nucleoid**
  - can occupy a significant fraction of the cell volume
  - appears attached at 1+ points in the inner surface of the plasma membrane



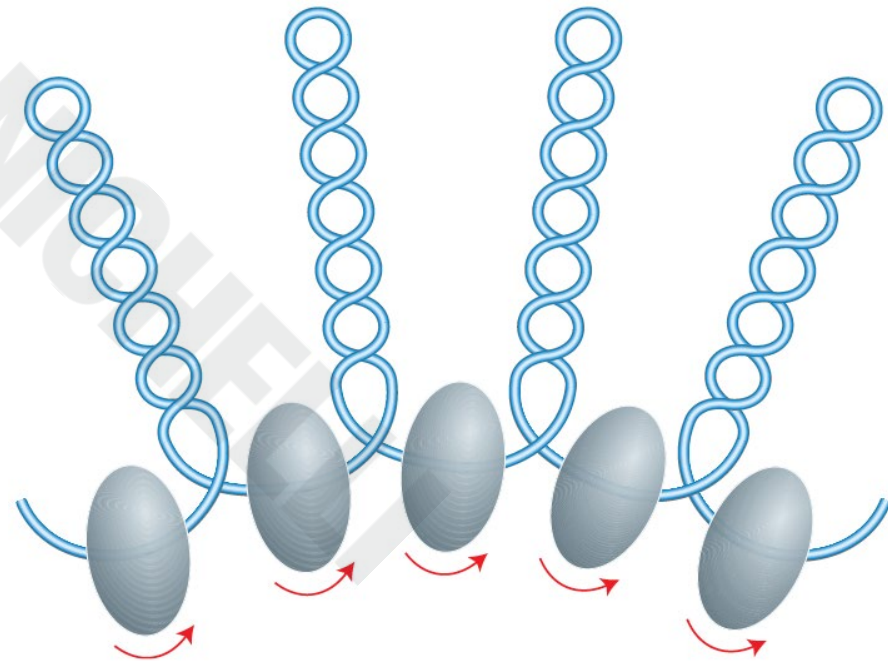
Lars Remmer.

## Principle 3 (7 of 7)

**Chromosomes in all cells are maintained in a state of torsional stress.** DNA is underwound relative to the stable B-form structure, facilitating both the packaging of DNA and access to the genetic information contained within it.

# Looped Domains of the *E. coli* Chromosome

- a scaffoldlike structure seems to organize the circular chromosome into a series of ~500 looped domains
  - domains are topologically constrained
  - domains do not have fixed end points



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## **Principle 4** (7 of 7)

**Specialized proteins and RNAs maintain chromosome structure.** Topoisomerases control DNA underwinding. Histones, condensins, cohesins, and other DNA-binding proteins provide scaffolds to organize chromosome structure. Certain long, noncoding RNAs also play important roles in chromosome structure and function.

# Histonelike Proteins Are Abundant in *E. coli*

- bacterial DNA does not seem to have comparable structure to eukaryotic nucleosomes
- histonelike proteins = proteins that bind and dissociate within minutes
  - no regular, stable DNA-histone structure has been found
- the dynamic and irregular structure of the bacterial chromosome reflects:
  - the shorter cell cycle
  - very active metabolism