

22. Black, sepia, cream, and albino are coat colors of guinea pigs. Individual animals (not necessarily from pure lines) showing these colors were intercrossed; the results are tabulated as follows, where the abbreviations A (albino), B (black), C (cream), and S (sepia) represent the following phenotypes:

Cross	Parental phenotypes	Phenotypes of progeny			
		B	S	C	A
1	B × B	22	0	0	7
2	B × A	10	9	0	0
3	C × C	0	0	34	11
4	S × C	0	24	11	12
5	B × A	13	0	12	0
6	B × C	19	20	0	0
7	B × S	18	20	0	0
8	B × S	14	8	6	0
9	S × S	0	26	9	0
10	C × A	0	0	15	17

- Deduce the inheritance of these coat colors and use gene symbols of your own choosing. Show all parent and progeny genotypes.
- If the black animals in crosses 7 and 8 are crossed, what progeny proportions can you predict by using your model?

		Phenotypes of progeny			
Cross	Parental phenotypes	B	S	C	A
1	B × B	22	0	0	7
2	B × A	10	9	0	0
3	C × C	0	0	34	11
4	S × C	0	24	11	12
5	B × A	13	0	12	0
6	B × C	19	20	0	0
7	B × S	18	20	0	0
8	B × S	14	8	6	0
9	S × S	0	26	9	0
10	C × A	0	0	15	17

60. Four homozygous recessive mutant lines of *Drosophila melanogaster* (labeled 1 through 4) showed abnormal leg coordination, which made their walking highly erratic. These lines were intercrossed; the phenotypes of the F1 flies are shown in the following grid, in which “+” represents wild-type walking and “-” represents abnormal walking:

Lines	1	2	3	4
1	-	+	+	+
2	+	-	-	+
3	+	-	-	+
4	+	+	+	-

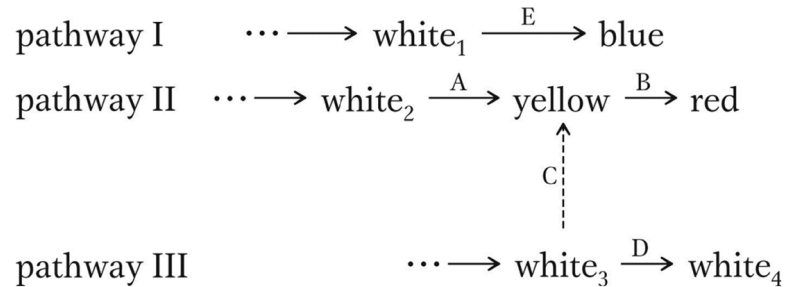
- What type of test does this analysis represent?
- How many different genes were mutated in creating these four lines?
- Invent wild-type and mutant symbols and write out full genotypes for all four lines and for the F1 flies.
- Do these data tell us which genes are linked? If not, how could linkage be tested?
- Do these data tell us the total number of genes taking part in leg coordination in this animal?

15. Several mutants are isolated, all of which require compound G for growth. The compounds (A to E) in the biosynthetic pathway to G are known, but their order in the pathway is not known. Each compound is tested for its ability to support the growth of each mutant (1 to 5). In the following table, a plus sign indicates growth and a minus sign indicates no growth.

Lines	A	B	C	D	E	G
1	-	-	-	+	-	-
2	-	+	-	+	-	+
3	-	-	-	-	-	+
4	-	+	+	+	-	+
5	+	+	+	+	-	+

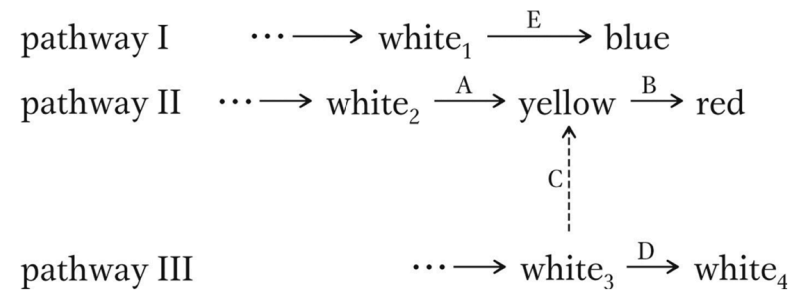
- What is the order of compounds A to E in the pathway?
- At which point in the pathway is each mutant blocked?
- Would a heterokaryon composed of double mutants 1, 3 and 2, 4 grow on a minimal medium? Would 1, 3 and 3, 4? Would 1, 2 and 2, 4 and 1, 4?

- 9 purple:3 green:4 blue
- 9 purple:3 red:3 blue:1 white
- 13 purple:3 blue
- 9 purple:3 red:3 green:1 yellow



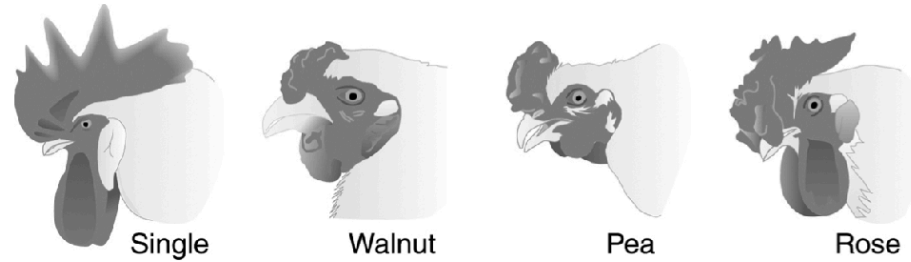
Assume that wild-type alleles are dominant and encode enzyme function and that recessive alleles result in a lack of enzyme function. Deduce which combinations of true-breeding parental genotypes could be crossed to produce F2 progeny in the following ratios:

- a. 9 purple:3 green:4 blue
- b. 9 purple:3 red:3 blue:1 white
- c. 13 purple:3 blue
- d. 9 purple:3 red:3 green:1 yellow



18. If a man of blood-group AB marries a woman of blood-group A whose father was of blood-group O, to what different blood groups can this man and woman expect their children to belong?

44. The genotype $r/r ; p/p$ gives fowl a single comb, $R/- ; P/-$ gives a walnut comb, $r/r ; P/-$ gives a pea comb, and $R/- ; p/p$ gives a rose comb (see the illustrations). Assume independent assortment.



- What comb types will appear in the F1 and in the F2 and in what proportions if single-combed birds are crossed with birds of a true-breeding walnut strain?
- What are the genotypes of the parents in a walnut x rose mating from which the progeny are 3/8 rose, 3/8 walnut, 1/8 pea, and 1/8 single?
- What are the genotypes of the parents in a walnut x rose mating from which all the progeny are walnut?
- How many genotypes produce a walnut phenotype? Write them out.

21. In the multiple-allele series that determines coat color in rabbits, c^+ encodes agouti, c^{ch} encodes chinchilla (a beige coat color), and c^h encodes Himalayan. Dominance is in the order $c^+ > c^{ch} > c^h$. In a cross of $c^+/c^{ch} \times c^{ch}/c^h$, what proportion of progeny will be chinchilla?

14. In *Drosophila*, the autosomal recessive *bw* causes a dark brown eye, and the unlinked autosomal recessive *st* causes a bright scarlet eye. A homozygote for both genes has a white eye. Thus, we have the following correspondences between genotypes and phenotypes:

$$\begin{aligned}st^+/st^+ ; bw^+/bw^+ &= \text{red eye (wild type)} \\st^+/st^+ ; bw / bw &= \text{brown eye} \\st / st ; bw^+/bw^+ &= \text{scarlet eye} \\st / st ; bw / bw &= \text{white eye}\end{aligned}$$

Construct a hypothetical biosynthetic pathway showing how the gene products interact and why the different mutant combinations have different phenotypes.

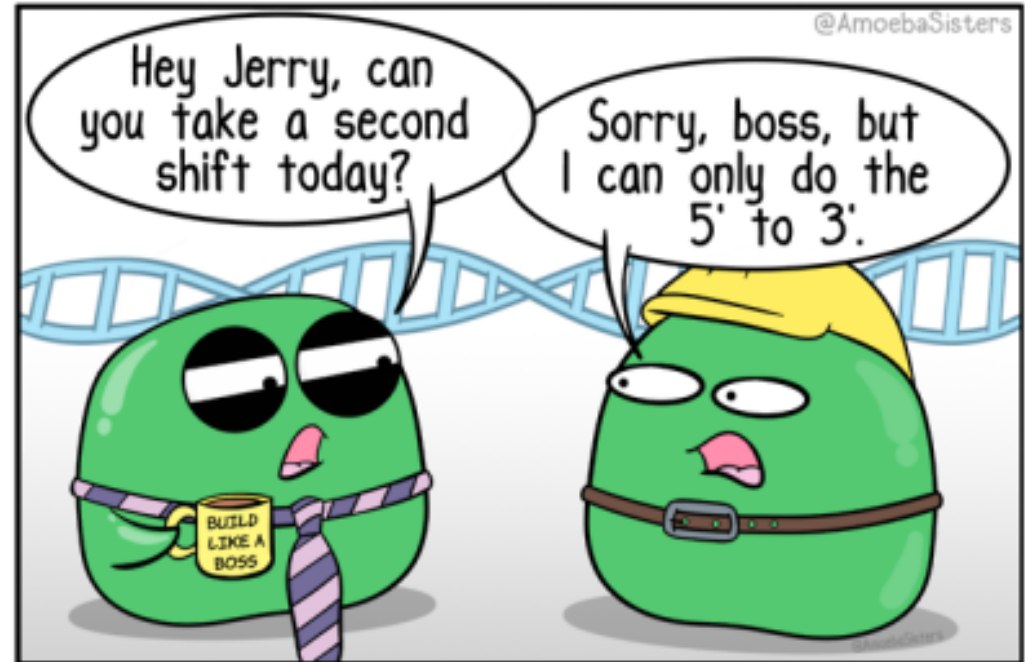
17. In sweet peas, the synthesis of purple anthocyanin pigment in the petals is controlled by two genes, B and D. The pathway is:

White intermediate —Gene B Enzyme—> blue intermediate —Gene D Enzyme—> anthocyanin (purple)

- a. What color petals would you expect in a pure-breeding plant unable to catalyze the first reaction?
- b. What color petals would you expect in a pure-breeding plant unable to catalyze the second reaction?
- c. If the plants in parts a and b are crossed, what color petals will the F1 plants have?
- d. What ratio of purple:blue:white plants would you expect in the F2?

Chapter 7: DNA Structure and Replication

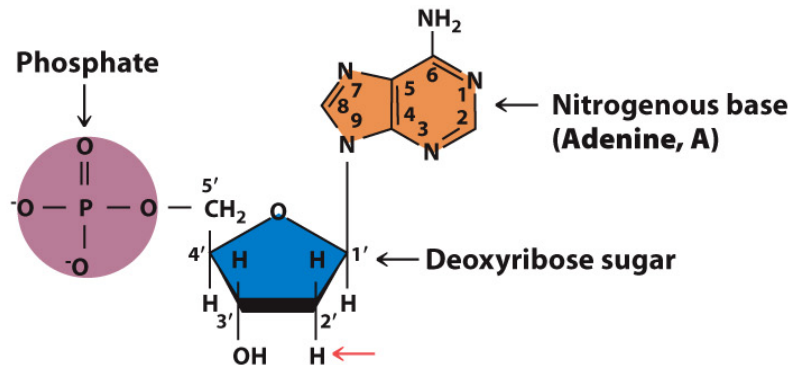
Paramecium Parlor



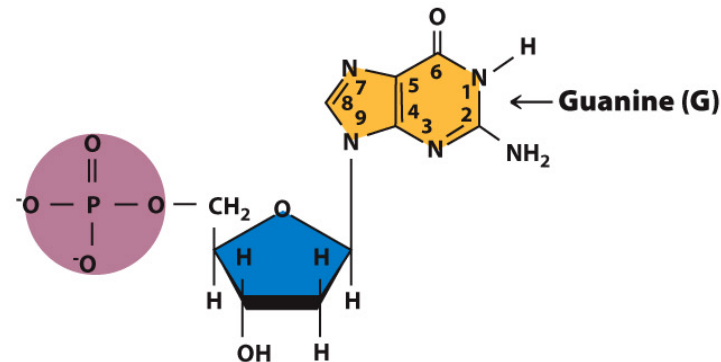
No one envied the job of the DNA polymerase shift manager.

Structure of the four DNA nucleotides

Purine nucleotides



Deoxyadenosine 5'-monophosphate (dAMP)



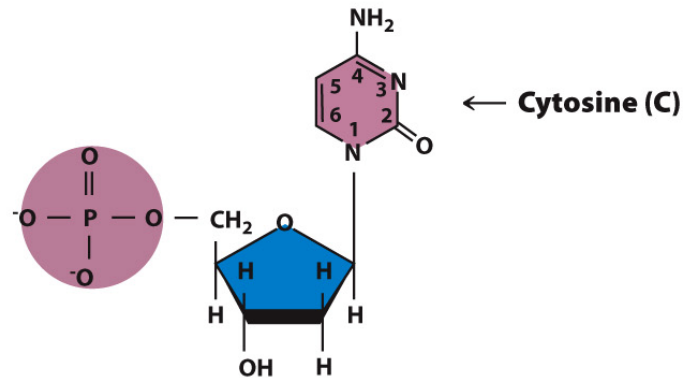
Deoxyguanosine 5'-monophosphate (dGMP)

Figure 7-5 part 1

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Structure of the four DNA nucleotides

Pyrimidine nucleotides

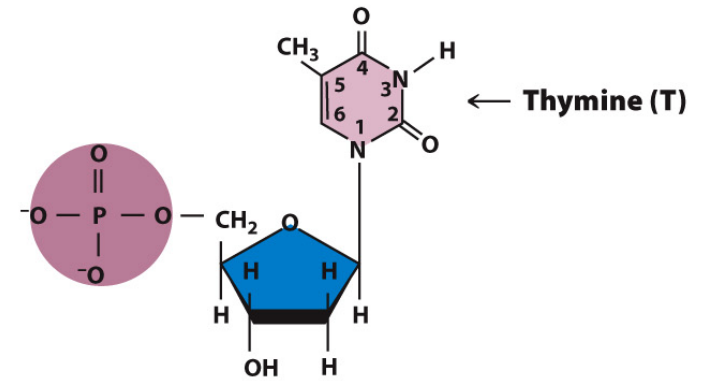


Deoxycytidine 5'-monophosphate (dCMP)

Figure 7-5 part 2

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Deoxythymidine 5'-monophosphate (dTMP)

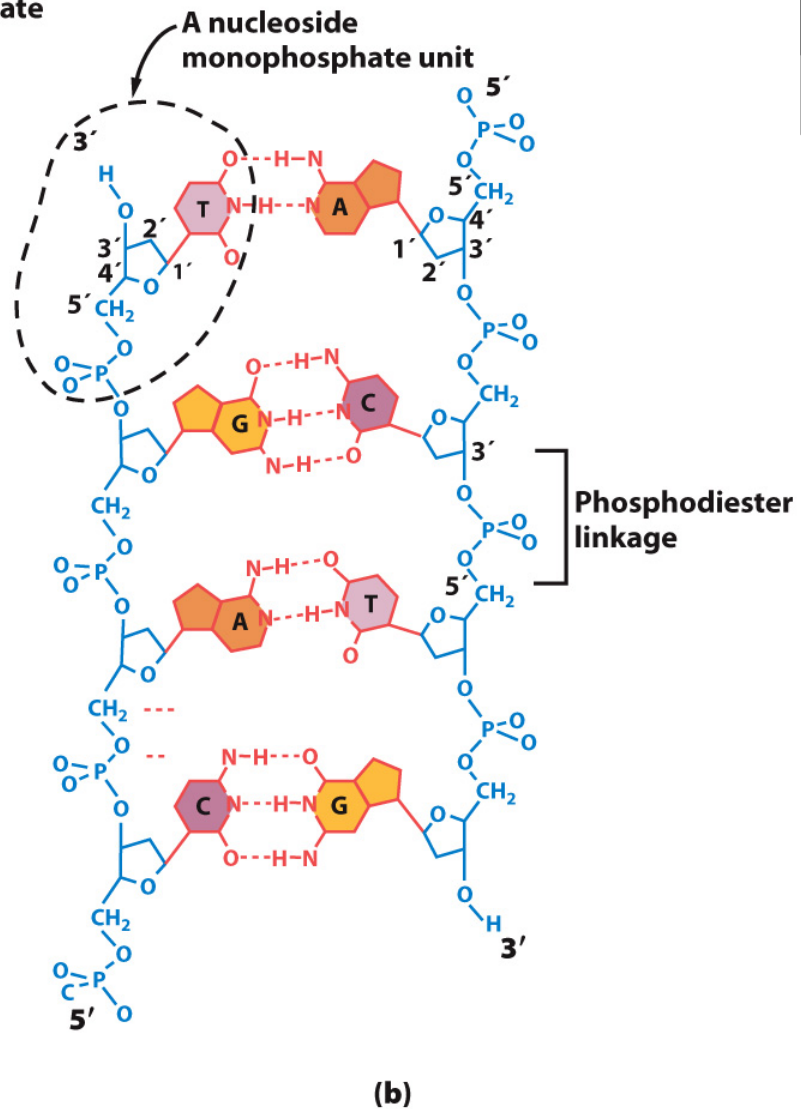
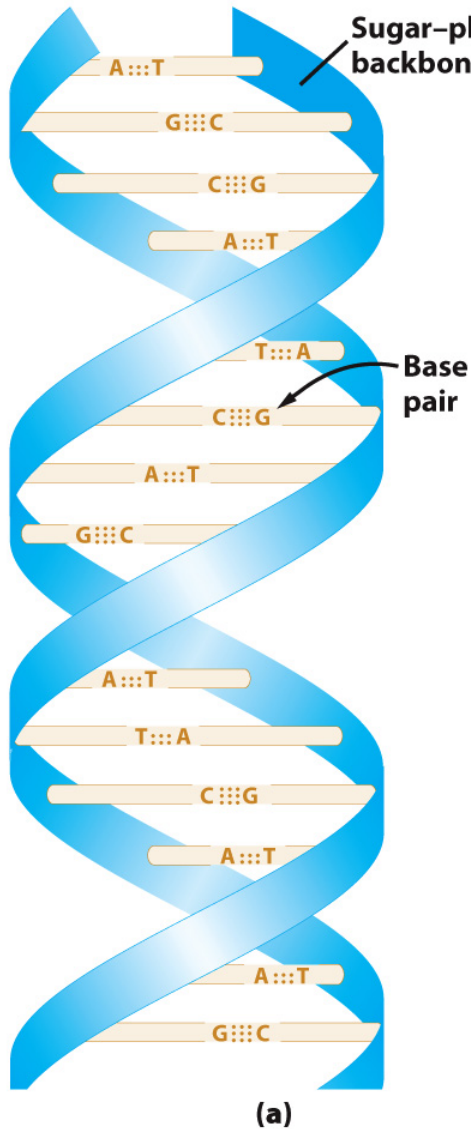
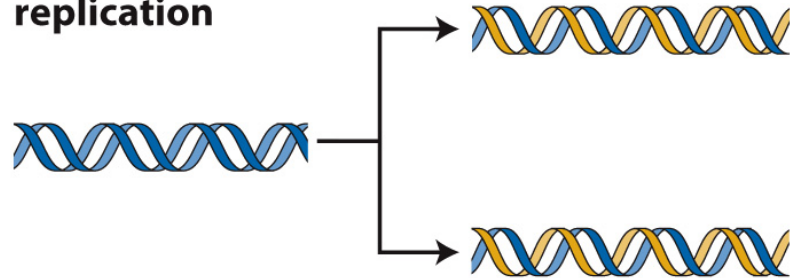


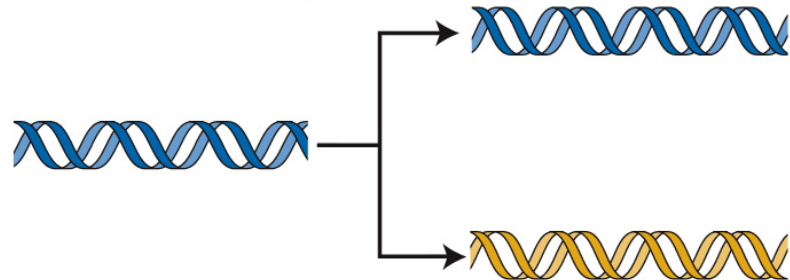
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Three alternative models for DNA replication

Semiconservative replication



Conservative replication



Dispersive replication

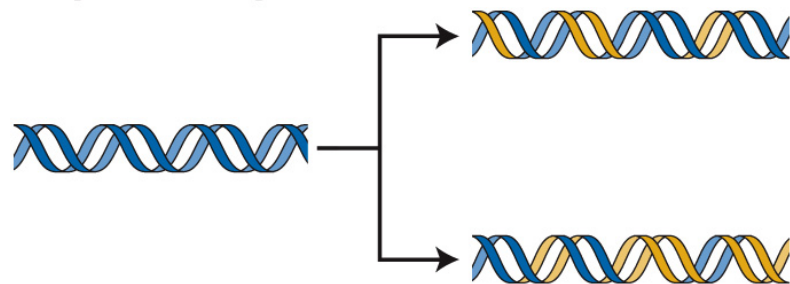


Figure 7-12

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DNA replication at the growing fork

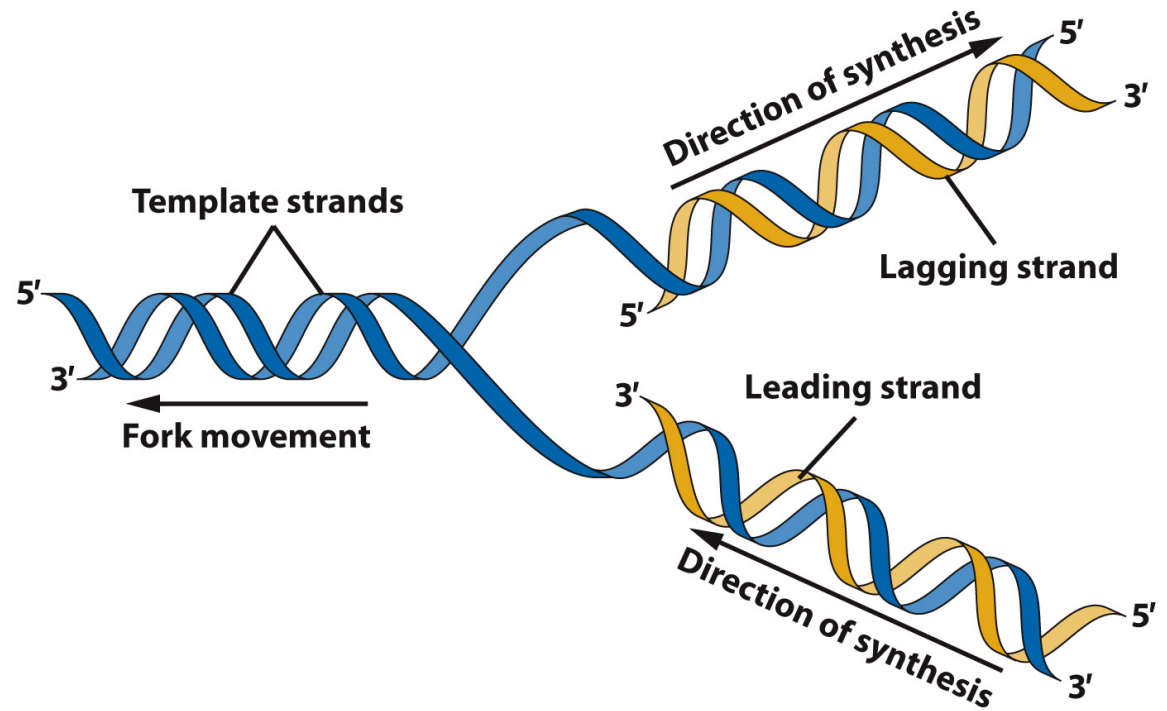


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Reaction catalyzed by DNA polymerase

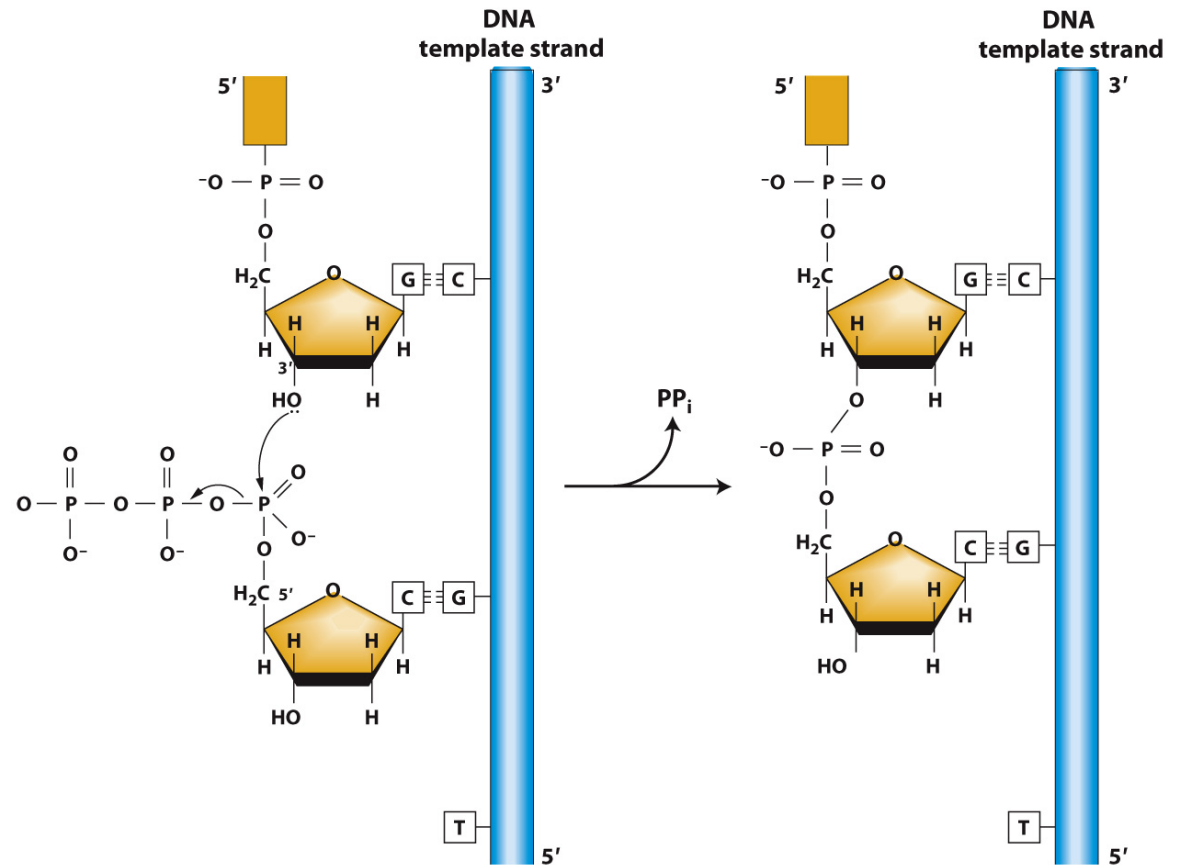


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Proteins at work at the replication fork

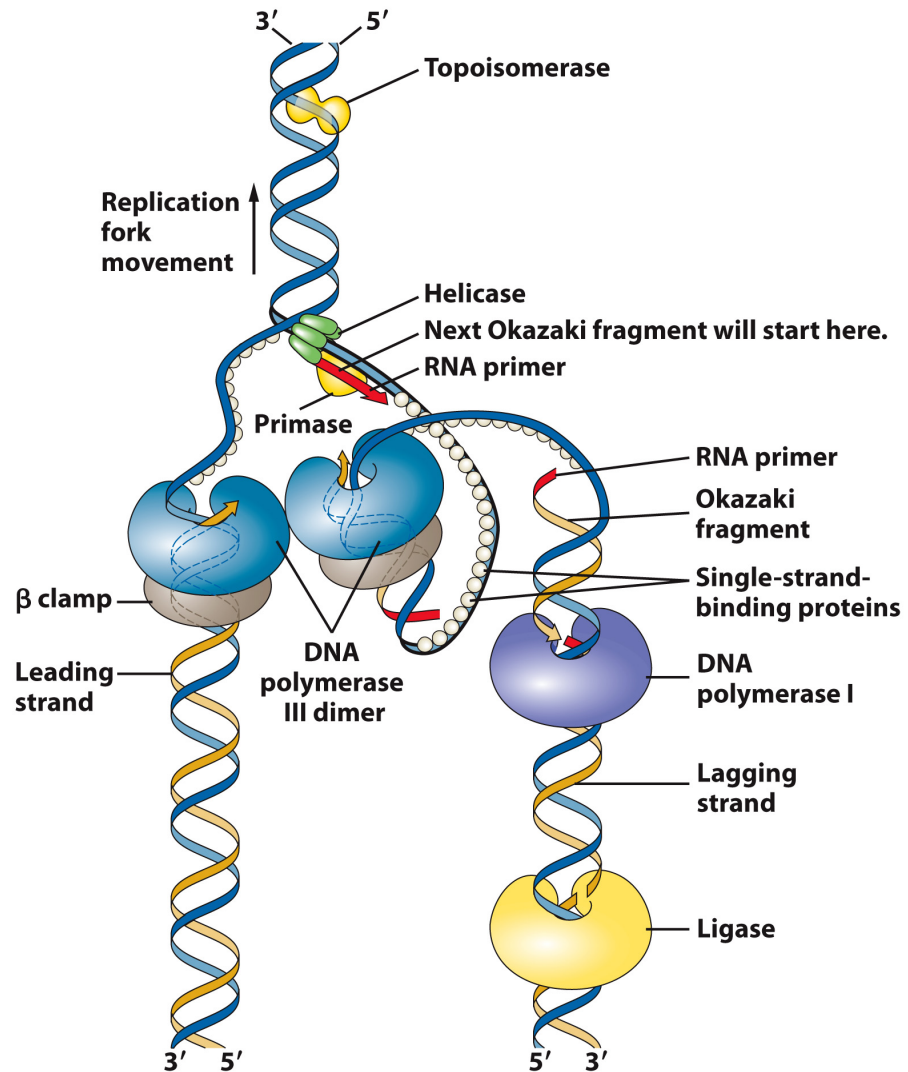


Figure 7-20

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DNA replication proceeds in two directions

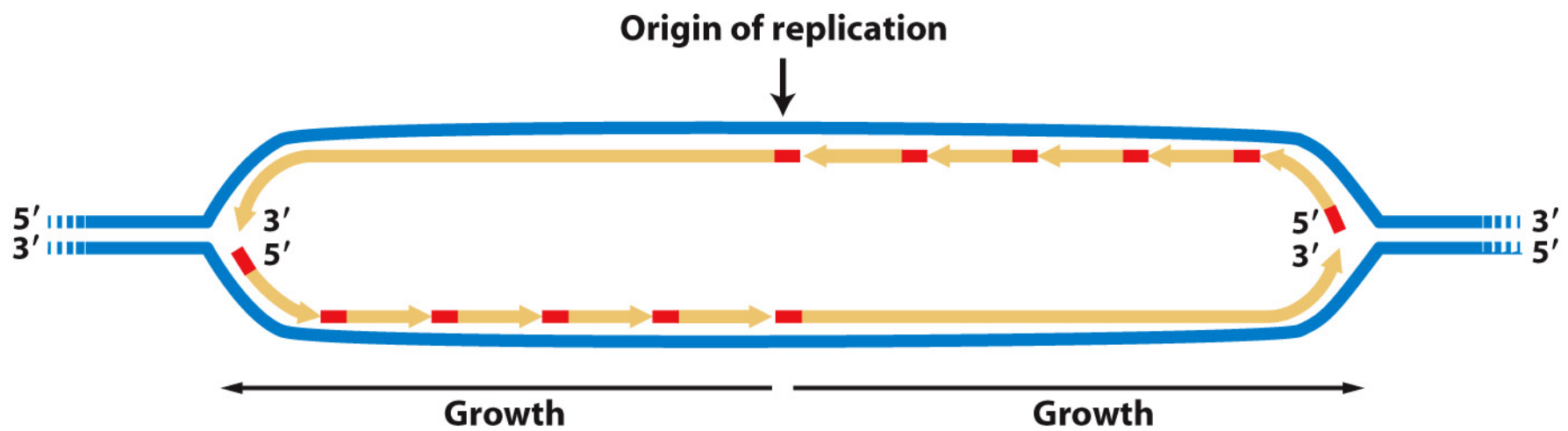


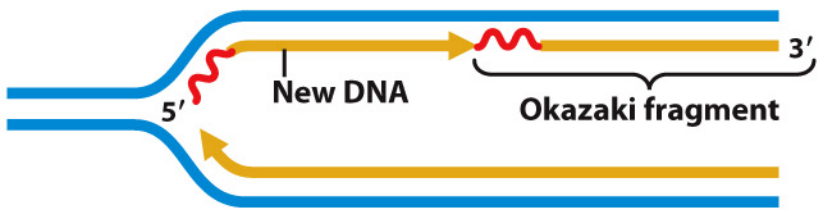
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Synthesizing the lagging strand

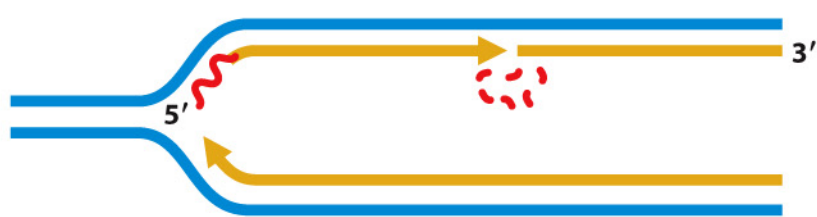
1. Primase synthesizes short RNA oligonucleotides (primers) copied from DNA.



2. DNA polymerase III elongates RNA primers with new DNA.



3. DNA polymerase I removes RNA at 5' end of neighboring fragment and fills gap.



4. DNA ligase connects adjacent fragments.



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The replication problem at chromosome ends

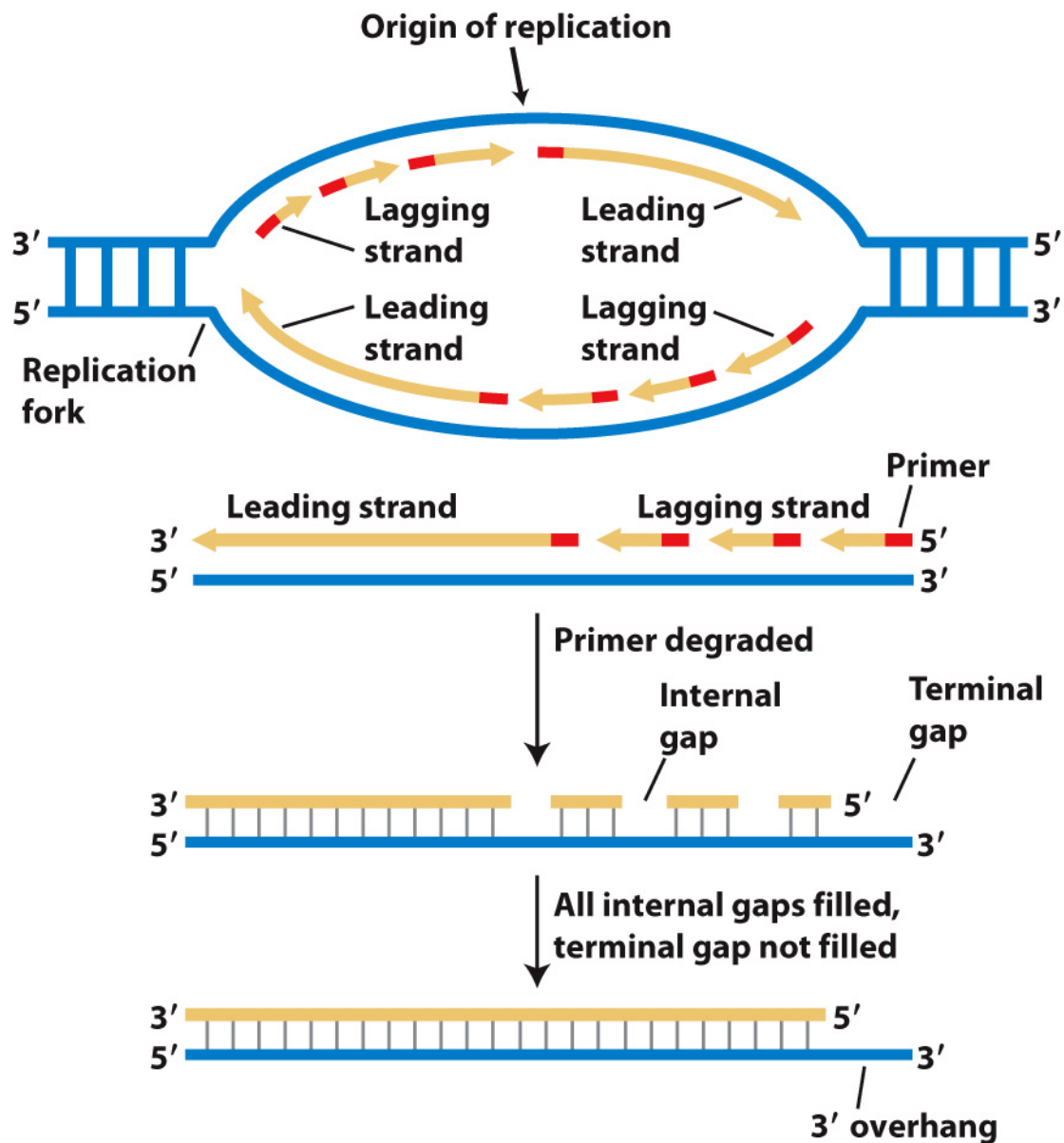


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Telomere lengthening

Lengthening of the 3' overhang

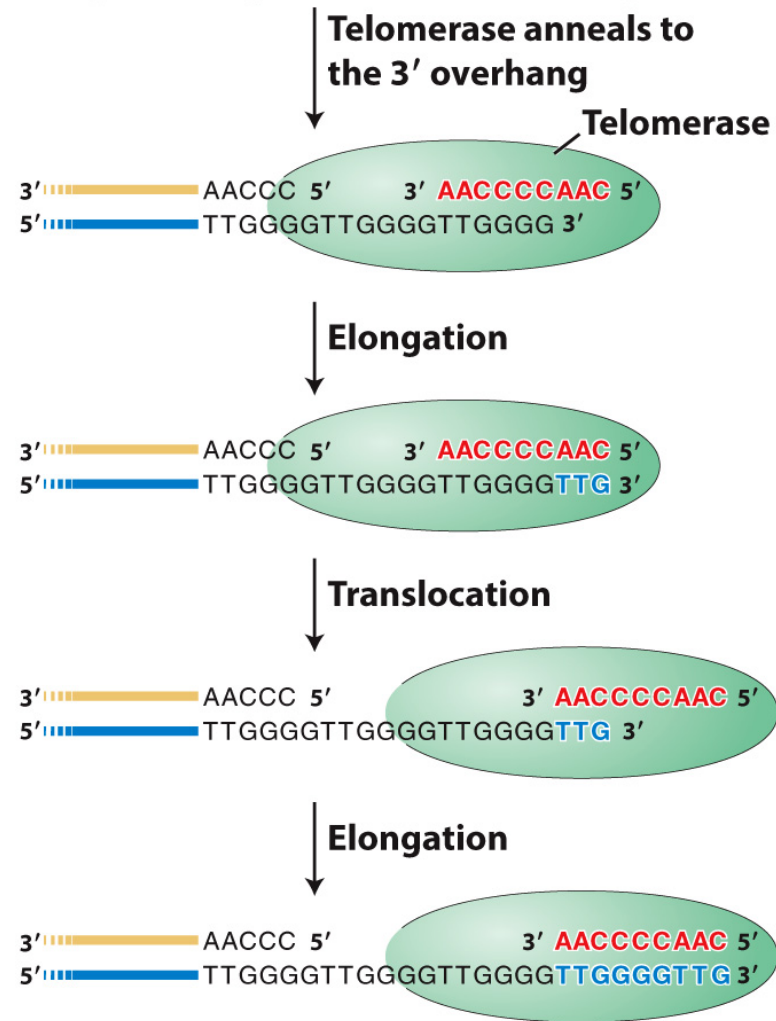


Figure 7-27a

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6. Describe the types of chemical bonds in the DNA double helix.

7. Explain what is meant by the terms *conservative* and *semiconservative* replication.

8. What is meant by a *primer*, and why are primers necessary for DNA replication?

9. What are helicases and topoisomerases?

10. Why is DNA synthesis continuous on one strand and discontinuous on the opposite strand?

12. If the helicases were missing during replication, what would happen to the replication process?

13. What would happen if, in the course of replication, the topoisomerases were unable to reattach the DNA fragments of each strand after unwinding (relaxing) the DNA molecule?

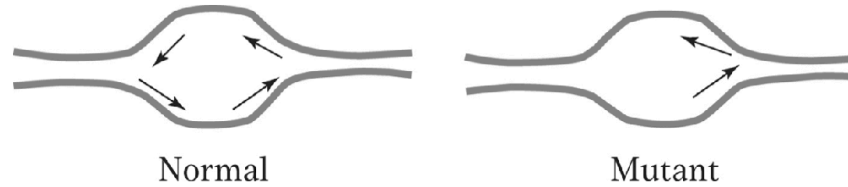
14. Which of the following is *not* a key property of hereditary material?
- a. It must be capable of being copied accurately.
 - b. It must encode the information necessary to form proteins and complex structures.
 - c. It must occasionally mutate.
 - d. It must be able to adapt itself to each of the body's tissues.

15. It is essential that RNA primers at the ends of Okazaki fragments be removed and replaced by DNA because otherwise which of the following events would result?
- The RNA would interfere with topoisomerase function.
 - The RNA would be more likely to contain errors because primase lacks a proofreading function.
 - The beta-clamp of the DNA pol II dimer would release the DNA and replication would stop.
 - The RNA primers would be likely to hydrogen bond to each other, forming complex structures that might interfere with the proper formation of the DNA helix.

16. Polymerases usually add only about 10 nucleotides to a DNA strand before dissociating. However, during replication, DNA pol III can add tens of thousands of nucleotides at a moving fork. How is this addition accomplished?

20. If the GC content of a DNA molecule is 48 percent, what are the percentages of the four bases (A, T, G, and C) in this molecule?

17. At each origin of replication, DNA synthesis proceeds bidirectionally from two replication forks. Which of the following would happen if a mutant arose having only one functional fork per replication bubble? (See diagram.)



- a. There is no change at all in replication
- b. Replication would take place only on one half of the chromosome.
- c. Replication would be complete only on the leading strand.
- d. Replication would take twice as long.

25. Consider the following segment of DNA, which is part of a much longer molecule constituting a chromosome:

5'....ATTCGTACGATCGACTGACTGACAGTC....3'
3'....TAAGCATGCTAGCTGACTGACTGTCAG....5'

If the DNA polymerase starts replicating this segment from the right,

- a. which will be the template for the leading strand?
- b. draw the molecule when the DNA polymerase is halfway along this segment.
- c. draw the two complete daughter molecules.
- d. is your diagram in part b compatible with bidirectional replication from a single origin, the usual mode of replication?