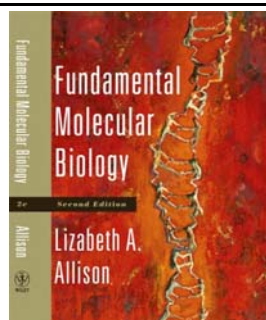


Terry Brown

Genomes
Third Edition

Chapter 17:
Recombination

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**Fundamental
Molecular Biology**
Second Edition

Lisabeth A. Allison

Chapter 7
DNA Repair Pathways

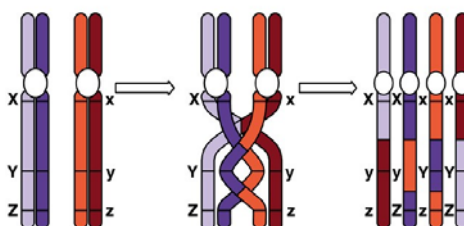
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Double Helix", oil and mixed media on canvas, © 2003

Recombination

- Outcome of crossing-over between pairs of homologous chromosomes during meiosis.
- Crossing-over results in daughter chromosomes that have different combinations of alleles compared with their parental chromosomes.

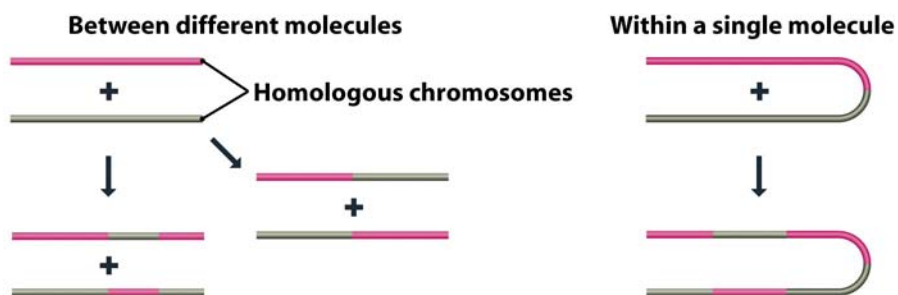
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Crossing over during meiosis



- A variety of processes that involve the breakage and reunion of polynucleotides.
 - 1) Homologous recombination
 - 2) Site-specific recombination
 - 3) Transposition.

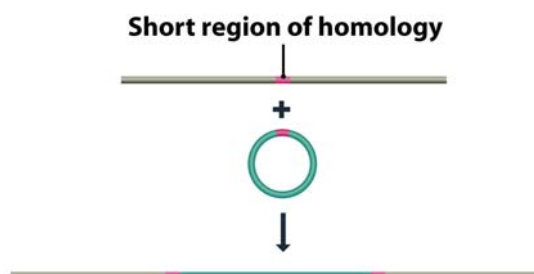
Homologous recombination



- Homologous recombination is exchange of DNA segments between two similar or identical molecules of DNA.
- It happens between different molecules or within a single molecule.
- It is responsible for crossing-over but its primary function is in DNA repair.

Figure 17.1a *Genomes 3* (© Garland Science 2007)

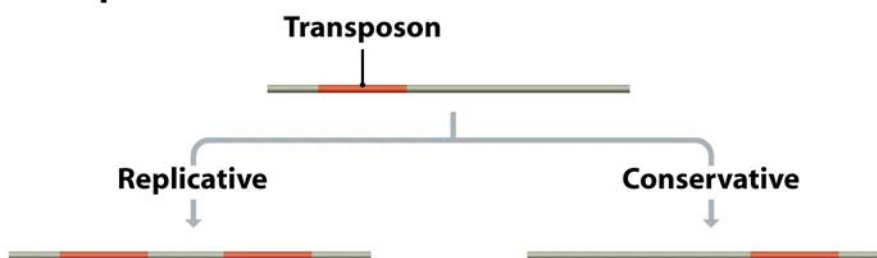
Site-specific recombination



- Site-specific recombination occurs between DNA segments possessing only a limited sequence homology, possibly just a few base pairs.
- The insertion of phage genome into bacterial genome is mediated by site-specific recombination.

Figure 17.1b *Genomes 3* (© Garland Science 2007)

Transposition

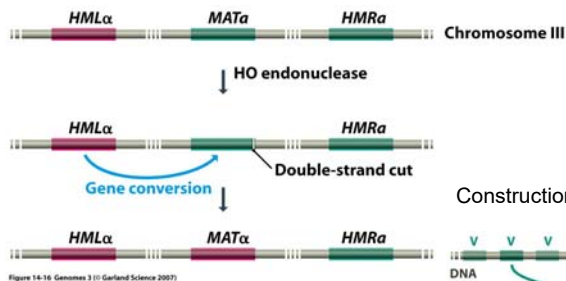


- Transposition is the transfer of a segment of DNA from one position in the genome to another position.

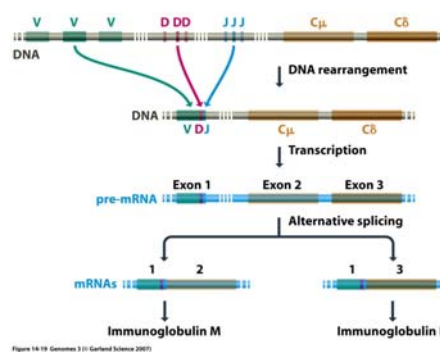
Figure 17.1c *Genomes 3* (© Garland Science 2007)

Other recombination events

Mating type switching in yeast



Construction of immunoglobulin genes



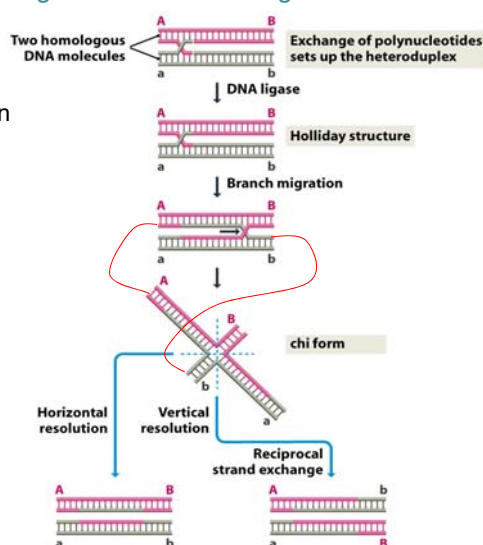
17.1 Homologous recombination

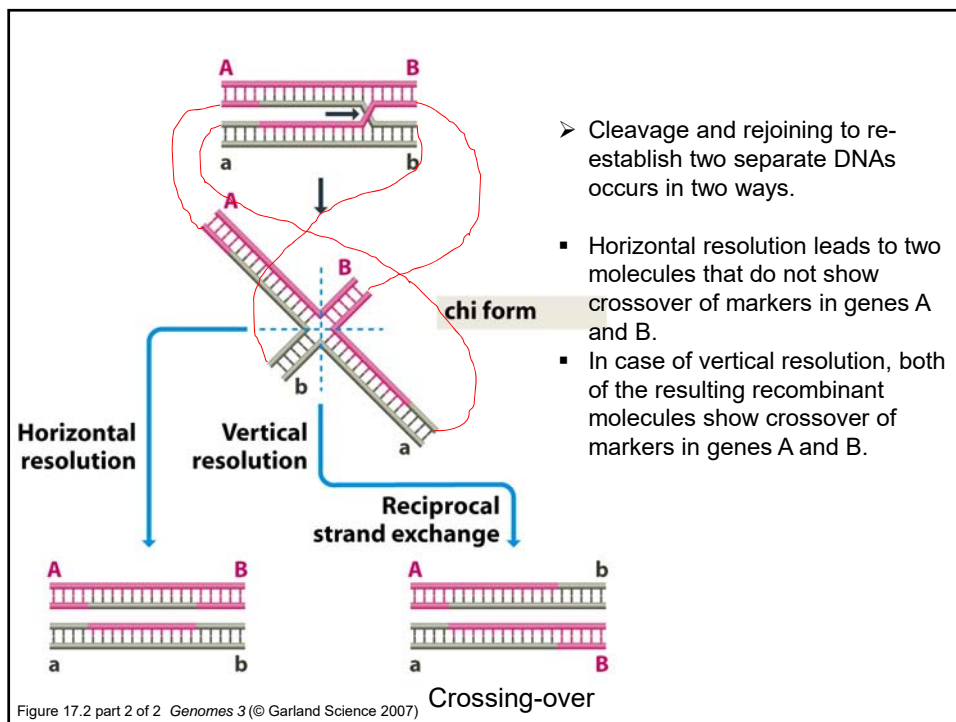
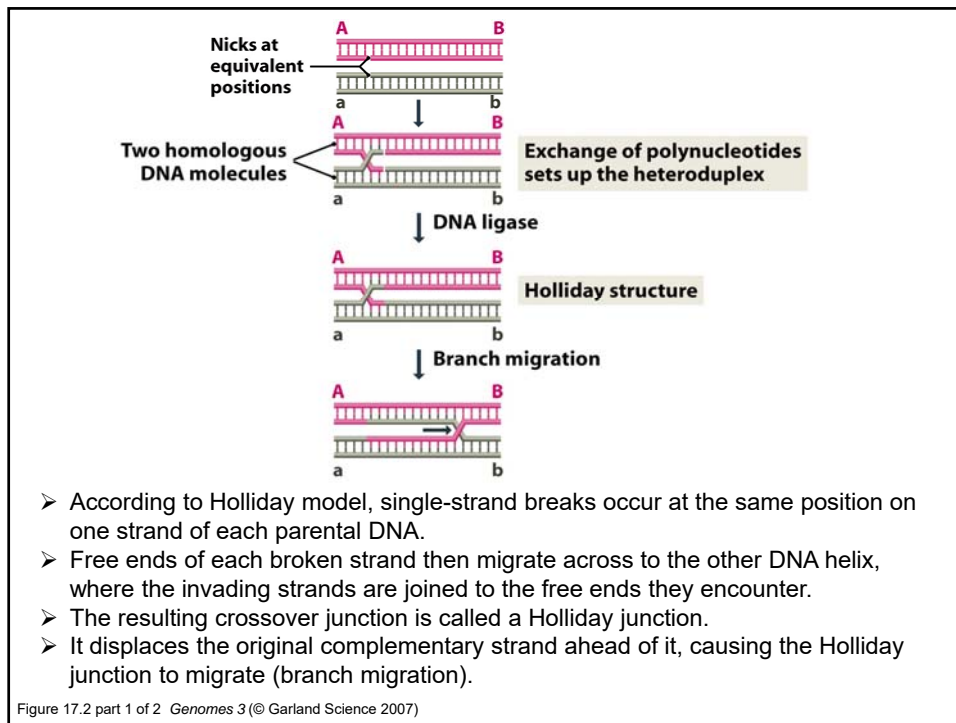
17.1.1 Models for homologous recombination

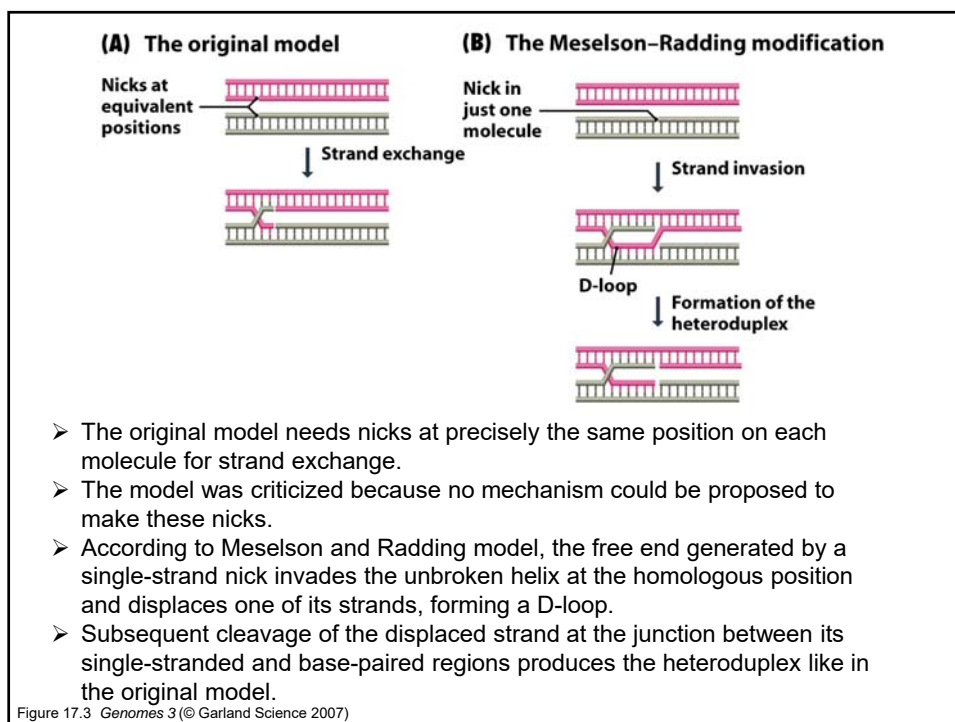
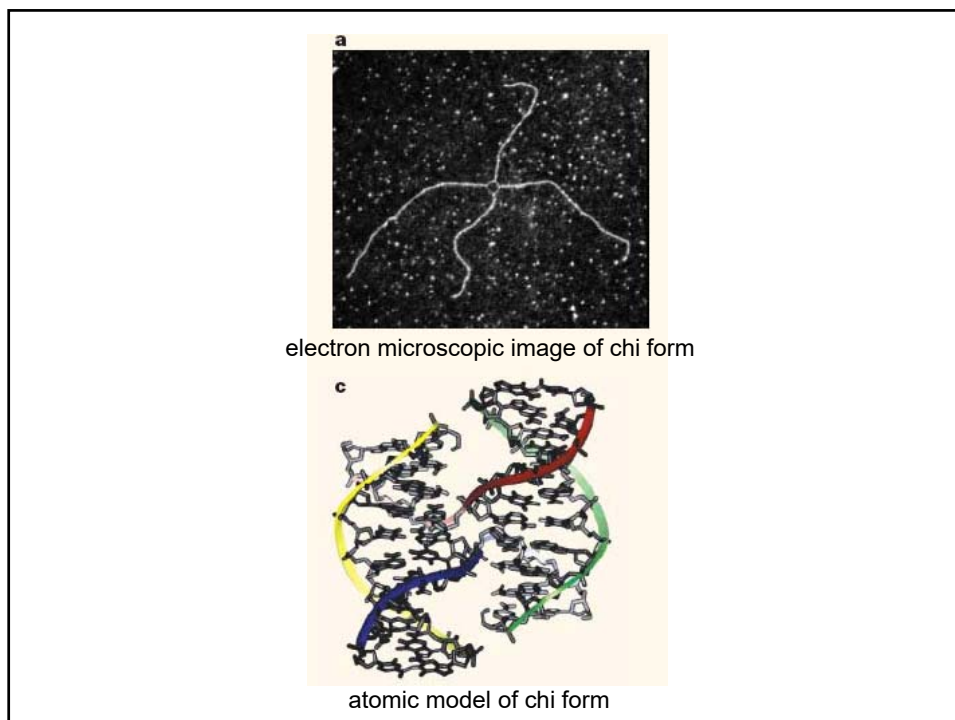
1. The Holliday and Meselson-Radding models for homologous recombination

➤ Robin Holliday and Matthew Meselson

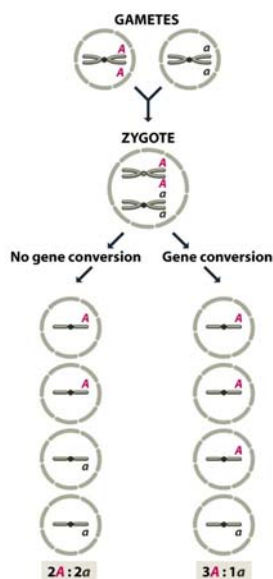
➤ The key feature of Holliday model is formation of a heteroduplex resulting from the exchange of polynucleotide segments between the two homologous molecules, called chi form.







2. The double-strand break model for homologous recombination



- Gene conversion is a process that results in the four haploid products of meiosis displaying an unusual segregation pattern.
- Holliday's original and Meselson's modified model cannot explain gene conversion.
- Without gene conversion, two gametes with allele 'A' and allele 'a' form a zygote, and produce four haploid spores 2A;2a alleles.
- If gene conversion occurs, the ratio will be changed like 3A;1a.

Figure 17.4 *Genomes 3* (© Garland Science 2007)

Double-strand break model

- 1) 100-1,000 times during meiosis
- 2) homologous recombination is involved in DNA repair (DSB repairing)

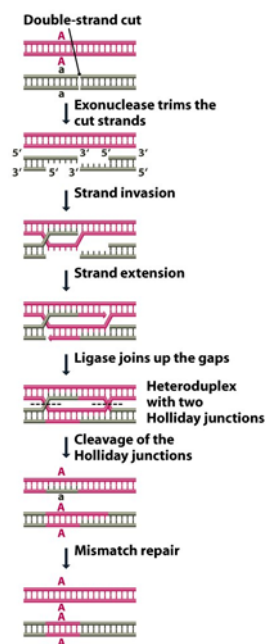
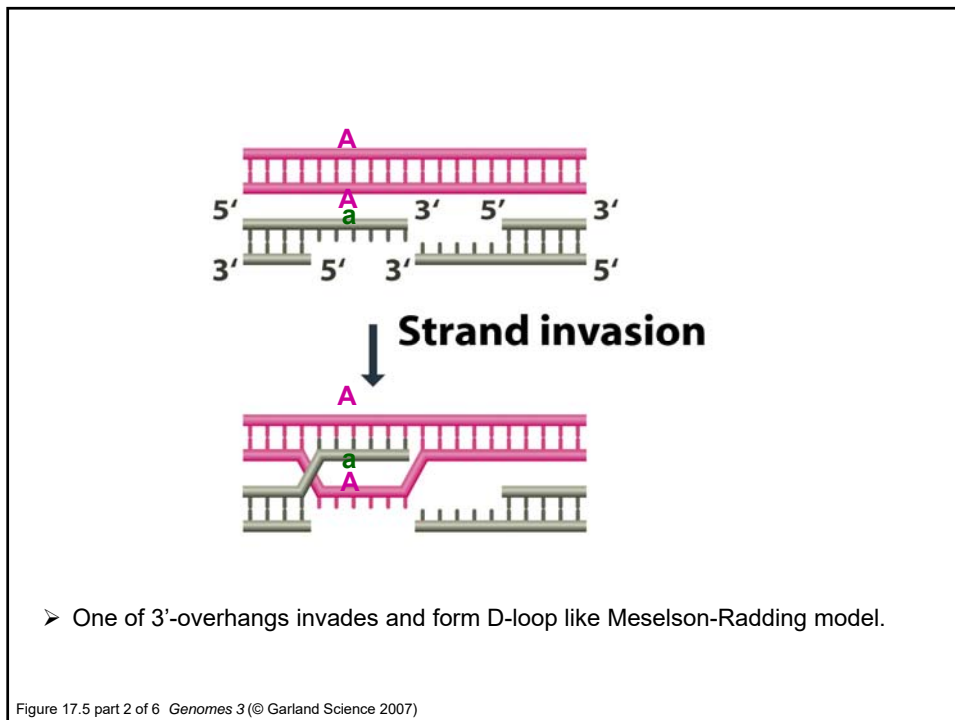
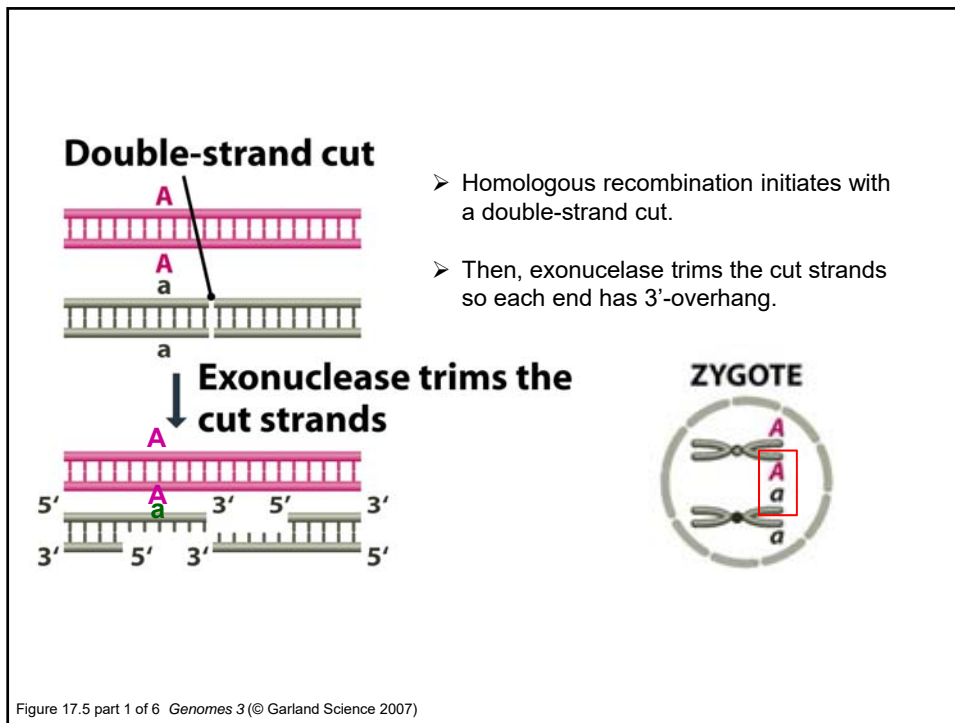
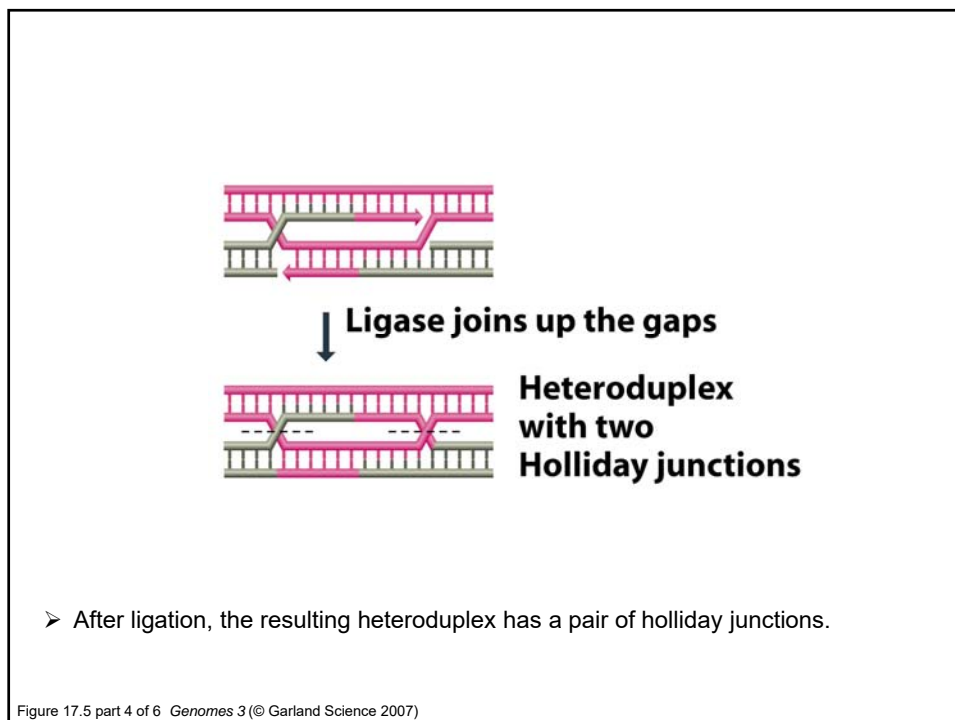
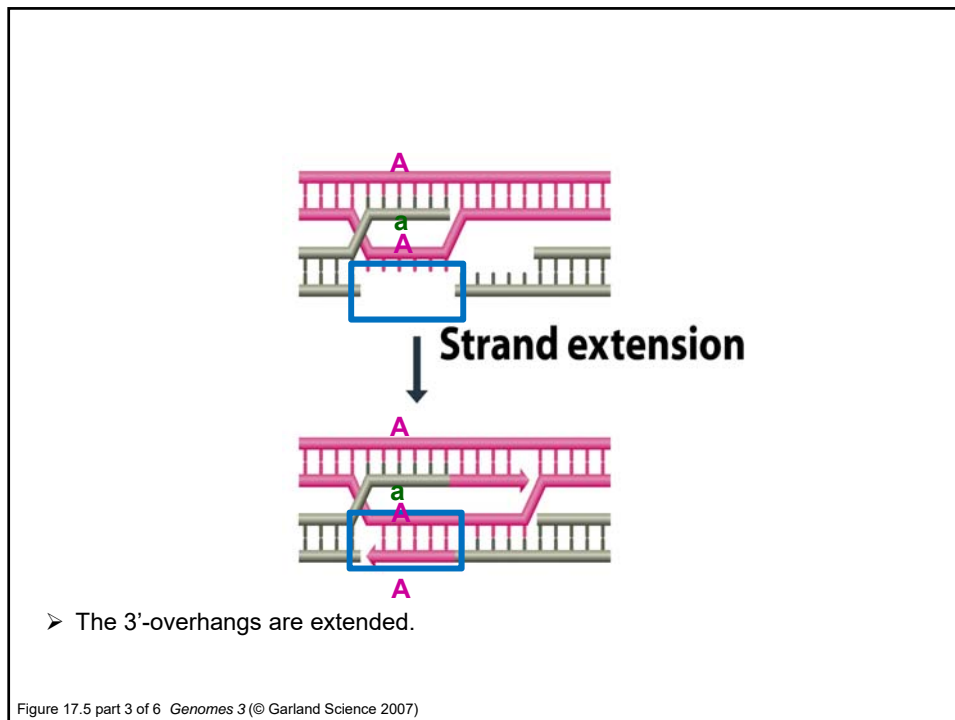
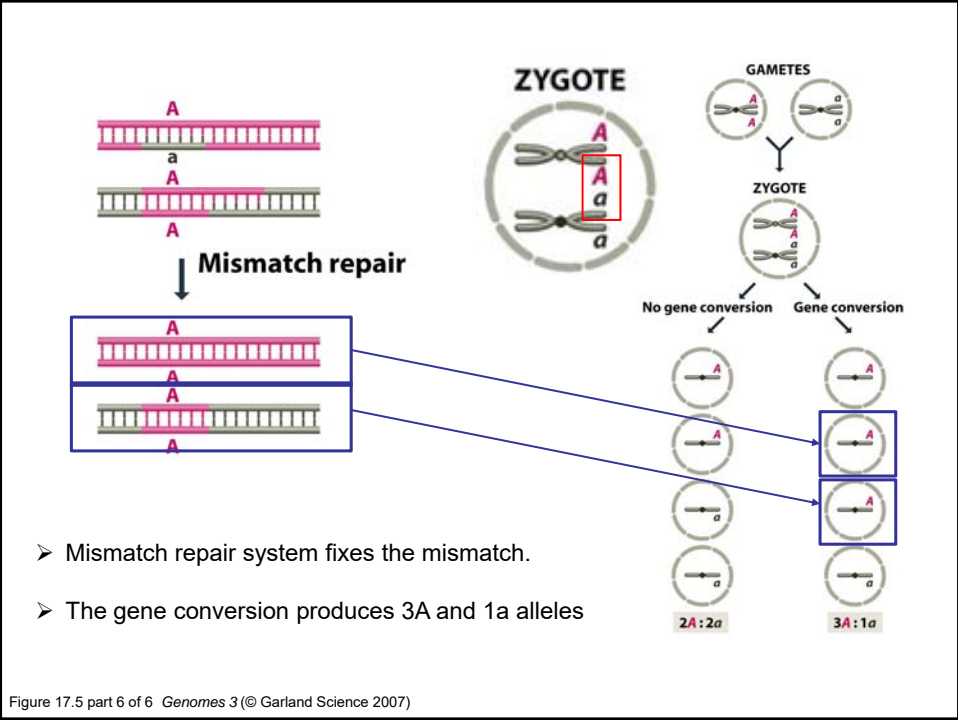
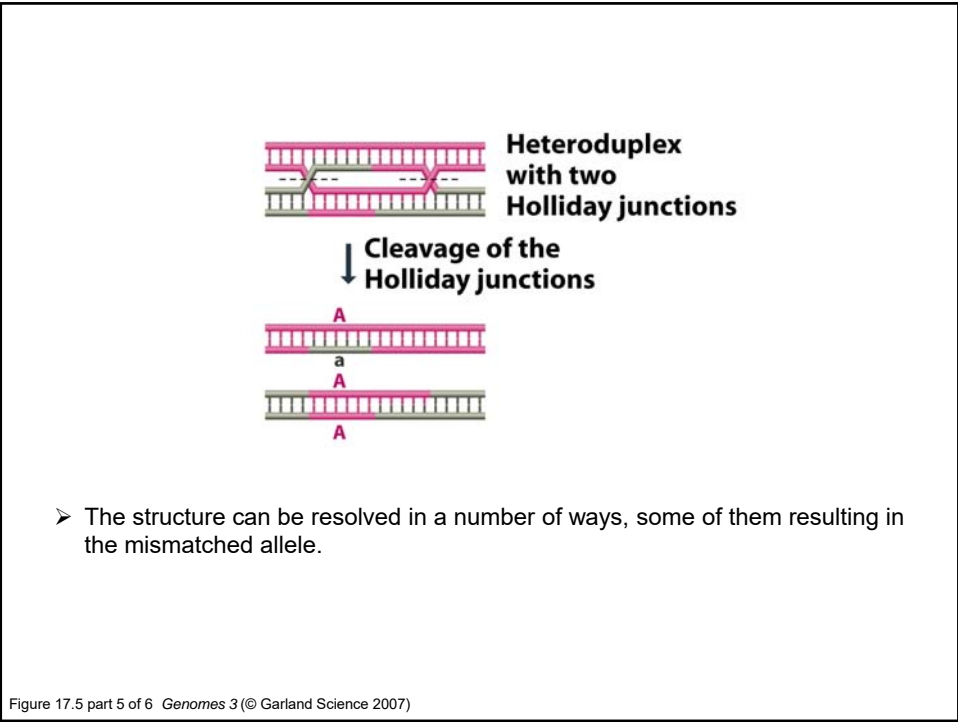


Figure 17.5 *Genomes 3* (© Garland Science 2007)







17.1.2 The biochemistry of homologous recombination

1. The RecBCD pathway of Escherichia coli

Progression of RecBCD along a DNA molecule

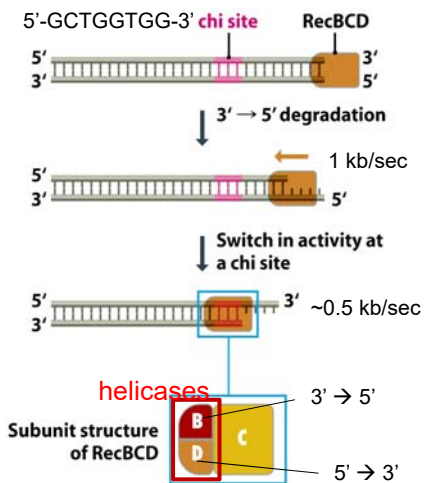


Figure 17.6 Genomes 3 (© Garland Science 2007)

- RecBCD enzymes mediate E. coli homologous recombination.
- To initiate the recombination, RecBCD complex attaches to the free ends of a chromosome at a double-strand break.
- RecD has a 5' to 3' helicase activity and RecB has a 3' to 5' helicase activity and 3' to 5' exonuclease activity.
- The enzyme complex moves and degrades the strand.
- The enzyme complex progresses along the DNA at a rate of ~1kb/sec until it reaches the chi site.
- The chi site occurs on average once every 6 kb.
- At the chi site the RecBCD complex undergoes conformation changes.
- The conformation changes slow down the migration of the RecBCD and abolish the 3' to 5' exonuclease activity of RecB, and make a single endonucleolytic cut in the other strand.

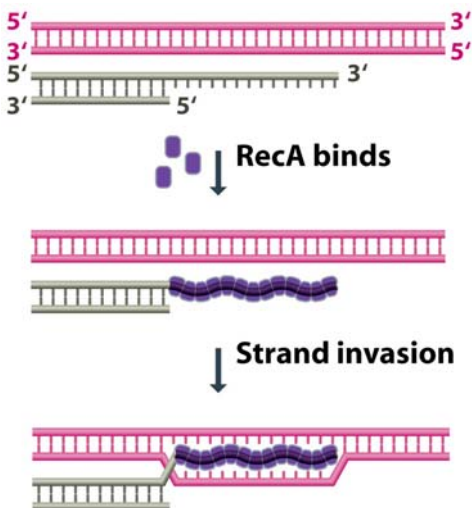
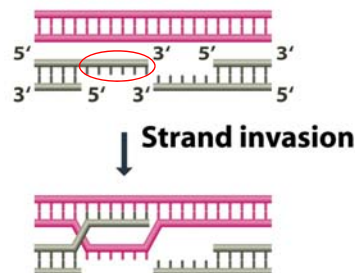
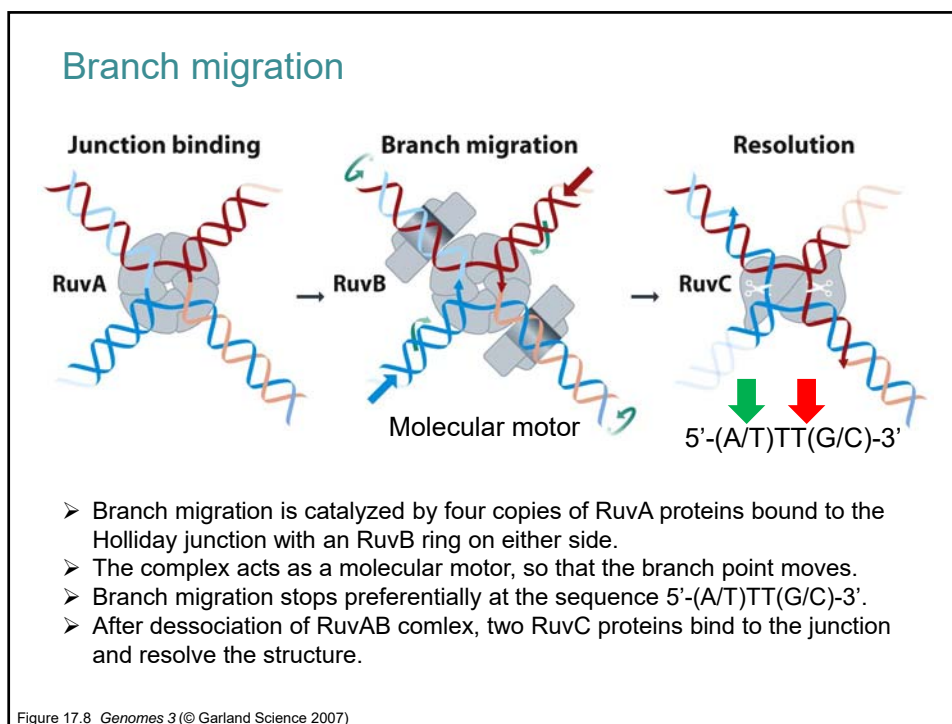


Figure 17.7 Genomes 3 (© Garland Science 2007)

- The result is a double-strand molecule with a 3'-overhang.
- RecA proteins form a protein-coated DNA filament.
- The filament invades the intact double helix and set up the D-loop.





2. Other homologous recombination pathways in *E. coli*

- *E. coli* has at least two alternative homologous recombination pathways, RecE and RecF although their recombination efficiency is low.

RecF pathway

- 1) RecT, RecA, RecF, RecQ, RecJ, RecO, RecR, RuvA, and UvrD
- 2) RecJ: exonuclease (remove 5'-end)
- 3) RecQ: helicase
- 4) RecO: promote annealing of complementary single-stranded DNA ends and strand exchange in RecA-mediated homologous recombination
- 5) RecR: interact with RecO, enhances RecO's ability to stimulate displacement of SSB protein from ssDNA by RecA
- 6) UvrD: helicase

RecE pathway

- 1) RecA, RecJ, RecO, RecR, RecQ, RecF

3. Model for mammalian homologous recombination to repair DSBs

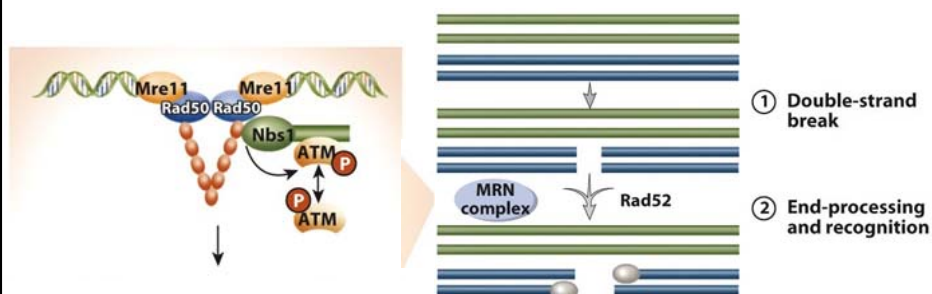
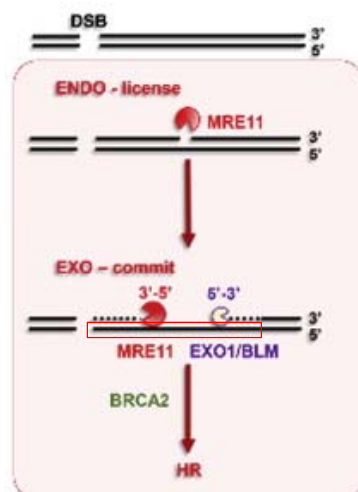


Figure 7.13
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- 1) Double-strand break (DSB) is induced by ionizing radiation.
- 2) End-processing and recognition:
 - a. The MRN (Mre11-Rad50-Nbs1) complex is recruited to the DSB and initiates repair.
 - b. The 3'→ 5' exonuclease activity of Mre11 generates single-strand DNA tails.
 - c. The MRN complex forms a bridge between free DNA ends via the coiled coil arms of the Rad50 dimers.
 - d. The ssDNA tails are then recognized by Rad52.



Molecular Cell Volume 53, Issue 1, p7-18

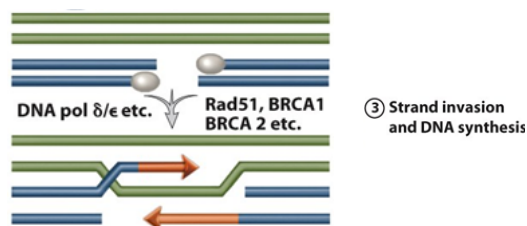


Figure 7.13
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- 3) Strand invasion and DNA synthesis:
- a. Rad51 initiates strand invasion of the 3' tails with homologous intact sequences.
 - b. Rad54, Rad55, Rad57, BRACA1, and BRACA2 are involved in homologous recombination.
 - BRACA2: interacts with RAD 51
 - c. Strand exchange generates a hybrid molecule.
 - d. Missing sequence information at the DSB is restored by DNA synthesis.

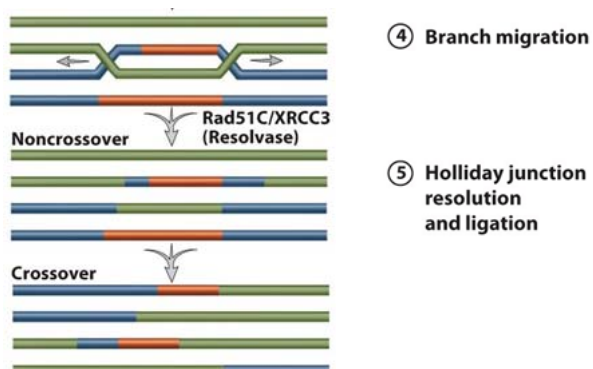


Figure 7.13
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- 4) Branch migration
- a. Rad54 is a motor protein that is important for branch migration.
 - b. Rad51C/XRCC3 complex is required for Holliday junction processing.
- 5) Holliday junction resolution and ligation

➤ ATM (ataxia telangiectasia mutated) and DSBs

- 1) ATM is a serine-threonine kinase, which is a key signal transducer.
- 2) MRN complexes form a bridge between free DNA ends via Rad50 dimers.
- 3) Inactive ATM dimers are recruited to the DSBs.
- 4) Conformation change of Nbs1 may activates ATM.
- 5) Activated ATM monomers either phosphorylate proteins (BRCA1, Smc1, Chk2) involved in DNA repair or phosphorylate proteins (p53, Creb) involved in cell cycle control.
- 6) Ataxia telangiectasia
 - Extreme sensitivity to radiation
 - Increased susceptibility to developing cancer
 - Immunodeficiency
 - Premature aging
 - Neurodegenerative disorder

17.1.3 Homologous recombination and DNA repair

1) Single-strand gap repair

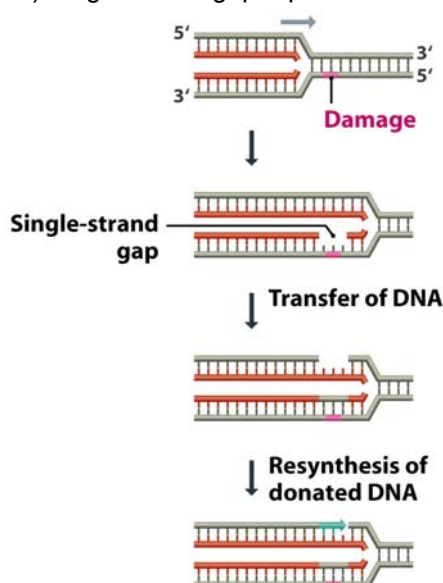
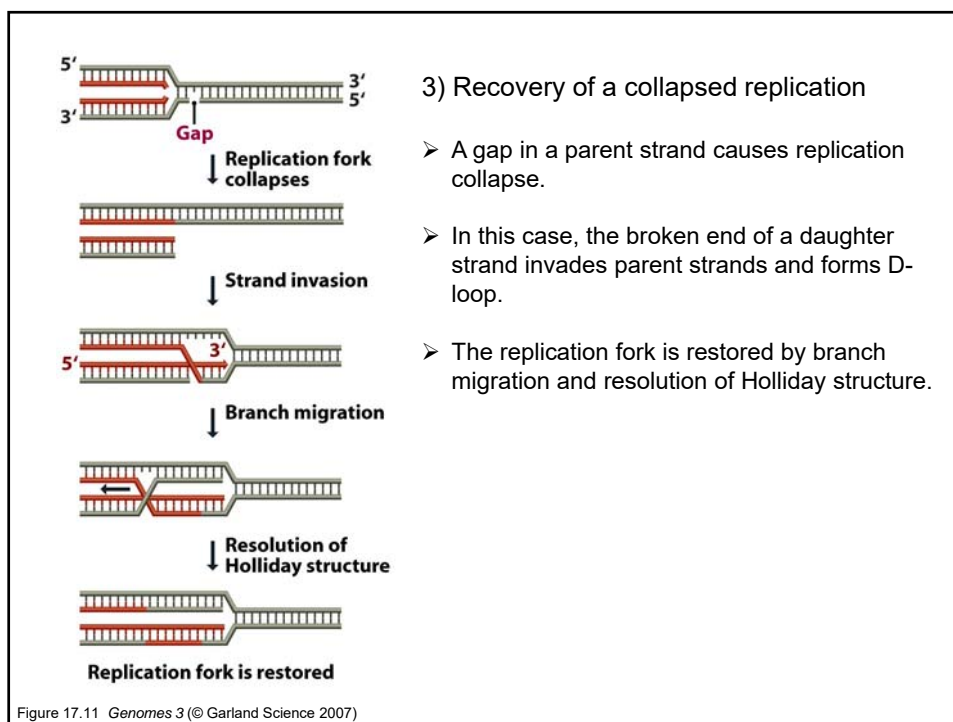
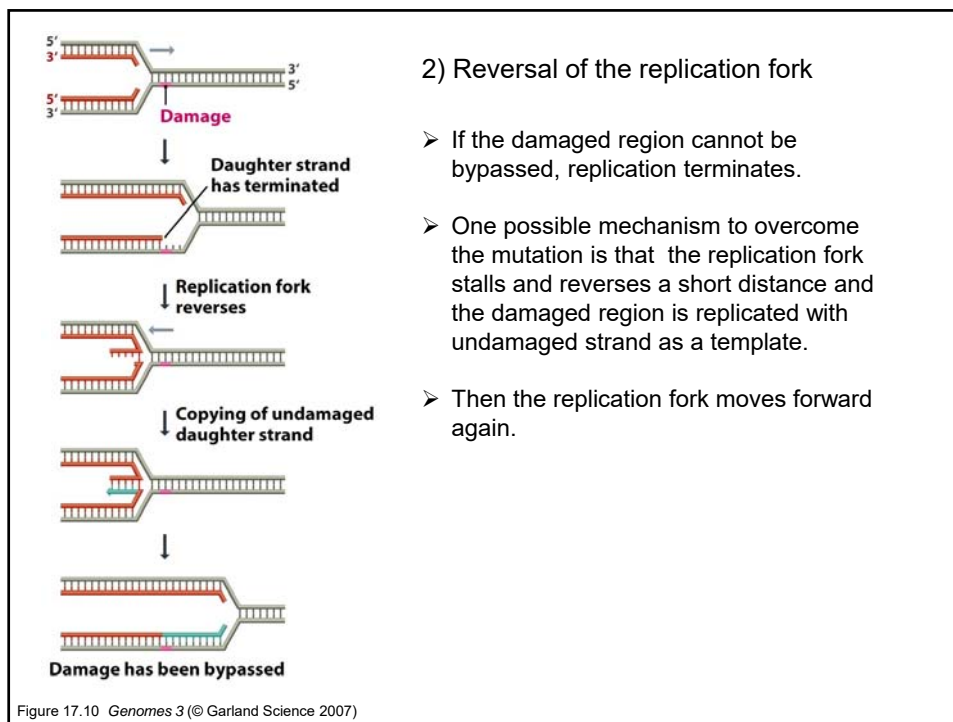


Figure 17-9 Genomes 3 (© Garland Science 2007)

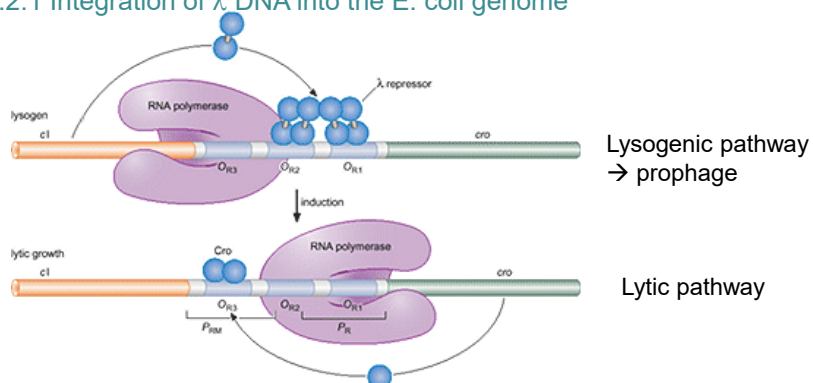
- The primary function of homologous recombination is in postreplicative DNA repair.
- The DNA polymerase encounters heavily damaged region like cyclobutyl dimers, the polymerase jumps ahead and restarts replication.
- The gap is repair by E. coli RecF pathway.





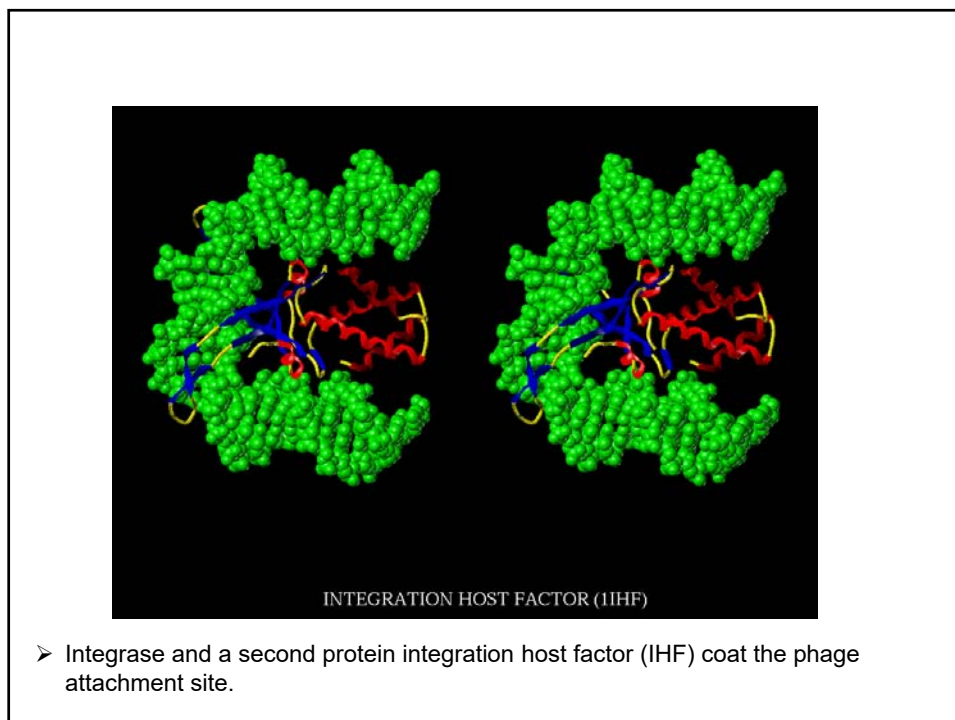
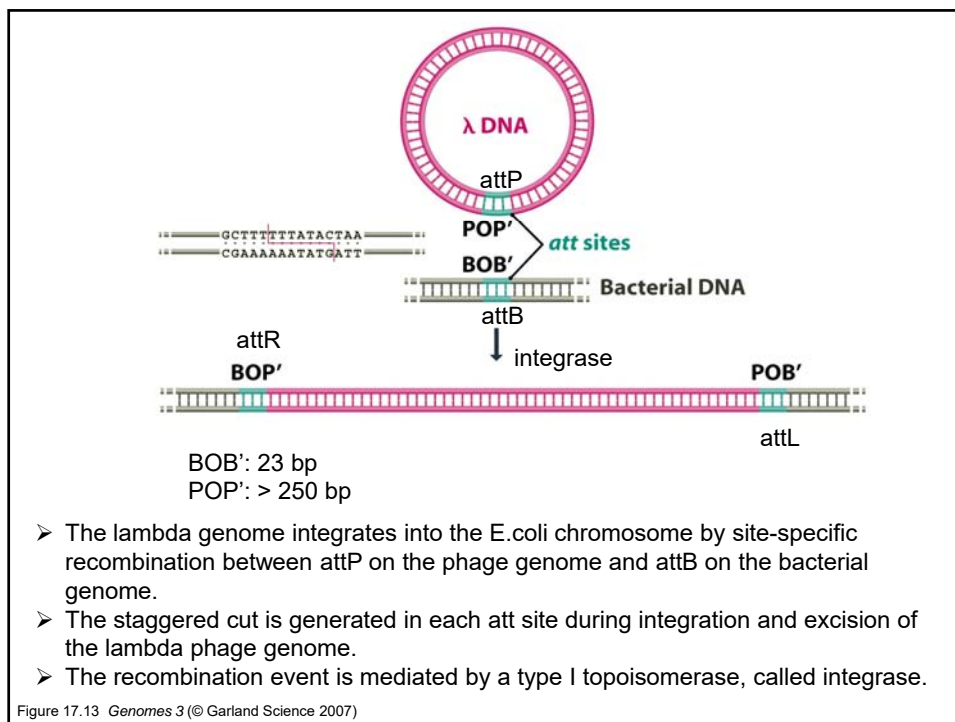
17.2 Site-Specific Recombination

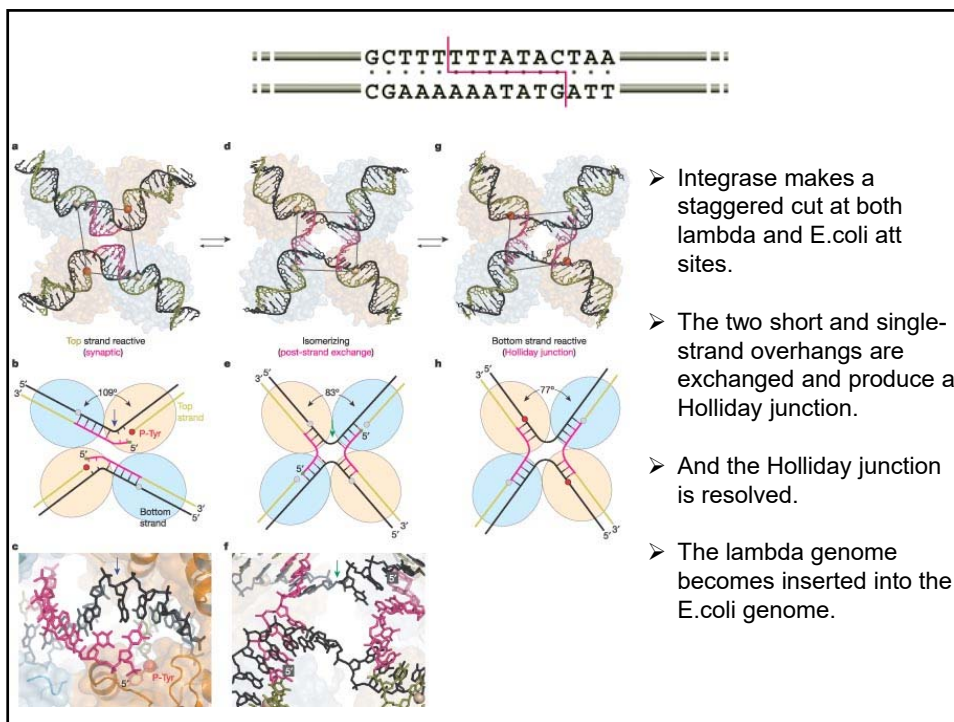
17.2.1 Integration of λ DNA into the E. coli genome



- After injecting its DNA into an E.coli cell, bacteriophage λ can follow either of two infection pathways.
- In the lytic pathway, new phages are rapidly generated in host E.coli and released within about 45 mins after infection.
- In the lysogenic pathway, lambda genome is integrated into bacterial genome and stays resident within the host's genome without apparent harm to the host.

Figure 17.12 Genomes 3 (© Garland Science 2007)

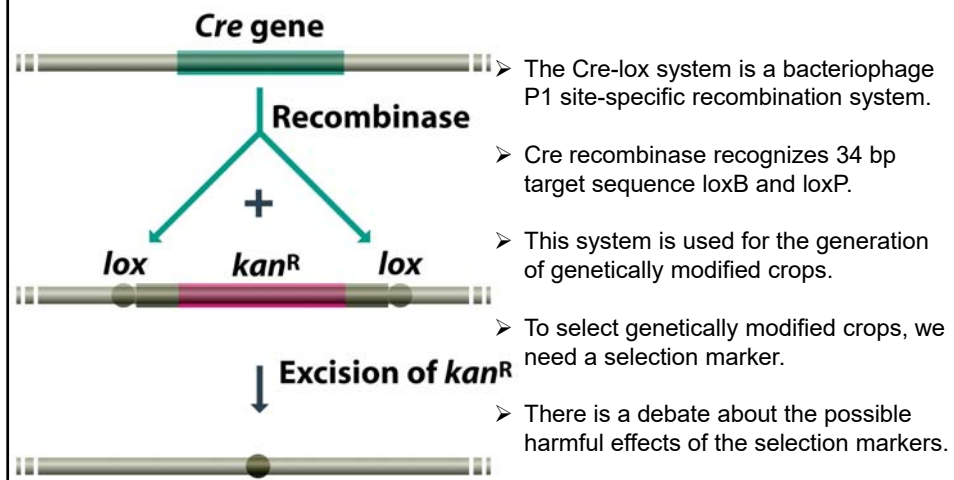




17.2.2 Site-specific recombination is an aid in genetic engineering

Bacteriophage P1

13bp 8bp 13bp
 ATAACCTCGTATA - GCATACAT -TATACGAAGTTAT



- The Cre-lox system is a bacteriophage P1 site-specific recombination system.
- Cre recombinase recognizes 34 bp target sequence loxB and loxP.
- This system is used for the generation of genetically modified crops.
- To select genetically modified crops, we need a selection marker.
- There is a debate about the possible harmful effects of the selection markers.

Figure 17.14 Genomes 3 (© Garland Science 2007)

Cre-loxP targeting

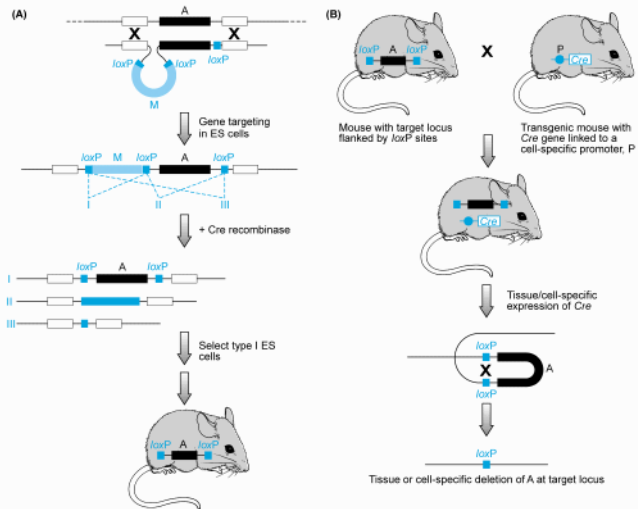
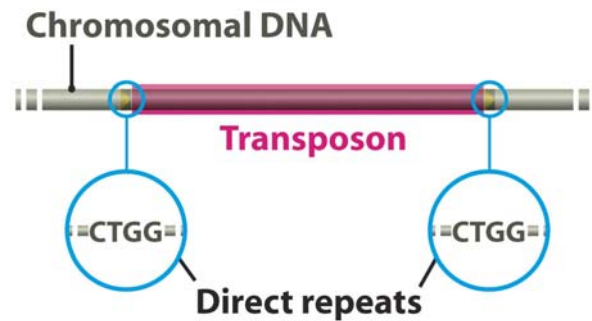


Fig. 21.8

17.3 Transposition



- Transposition is not a type of recombination.
- It often utilizes recombination.
- The end result is the transfer of a segment of DNA from one position in the genome to another position.
- A characteristic feature of transposition is a pair of short direct repeats flanking the transferred segment.
- Barbara McClintock discovered mobile genetic elements in maize (1983 Nobel prize)

Figure 17.15 *Genomes 3* (© Garland Science 2007)

- Abundant in the genomes of bacteria, plants, and animals.
 - 1) Mammals: nearly half the genome
 - 2) Some higher plants: ~90%
- Transposition (movement) may disrupt genetic function and result in phenotypic variation.
- In vertebrates and higher plants, only a low percentage of spontaneous mutations are caused by transposable elements.
- The primary function of eukaryotic DNA methylation may be defense of the genome from transposition of transposable elements.

Table 12.3 Classes of transposable elements.

Class	Transposition intermediate	Examples
Class I		
LTR retrotransposons	RNA	Yeast: Ty elements Human: Human endogenous retroviruses (HERV) Mouse: Intracisternal A particles (IAPs)
Non-LTR retrotransposons	RNA	Human: L1 elements Alu elements
Class II		
DNA transposons	DNA	Bacteria: Insertion sequence Bacteriophage Mu Transposons (e.g., Tn7) <i>Drosophila</i> : P elements Maize: Ac and Ds elements Invertebrates and vertebrates: Tc1/mariner superfamily

Table 12.3
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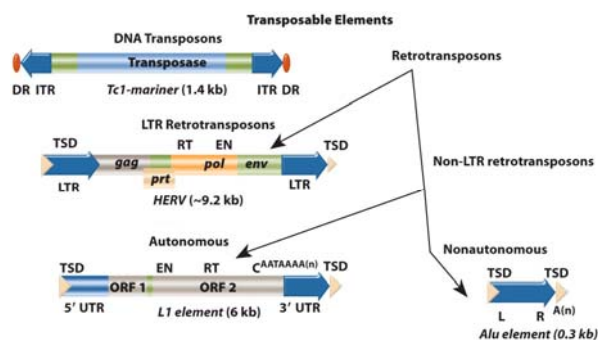


Figure 12.10
Adapted with permission from Ottertag & Kazanin, 2001. Annual Review of Genetics 35:501-538. Copyright © 2001 by Annual Reviews.

➤ Two main classes of transposable elements

1) DNA transposons: DNA intermediate during transposition.

- Inverted terminal repeats (ITRs), transposase, short direct repeats (DRs)

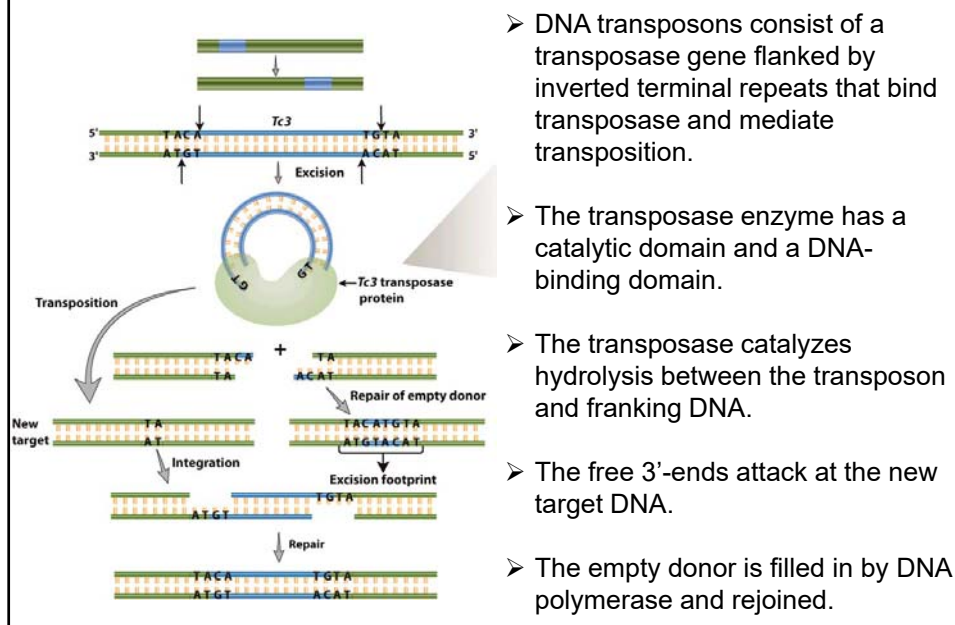
2) Retrotransposons: RNA intermediate during transposition.

a. LTR retrotransposons: long terminal repeats (LTRs), RT and endonuclease domains (EN), group-specific antigen (gag), protease (prt), polymerase (pol), envelope (env).

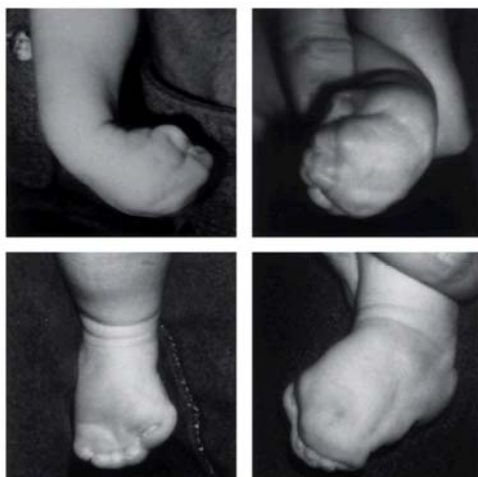
b. Autonomous: 5'UTR with promoter, RT, EN, ORF1, ORF2, 3' UTR.

c. Nonautonomous: L and R sequences, poly (A).

17.3.1 DNA transposons move by a “cut-and paste” mechanism



- Transposable elements provide material for DNA mispairing and unequal crossing-over.
- They are potential causal agents of human disease through insertion mutagenesis.

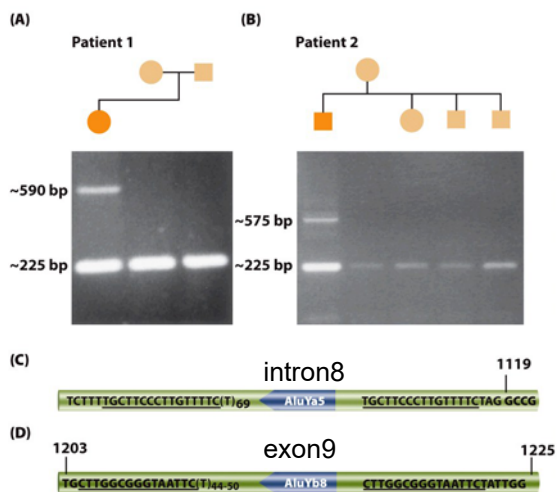
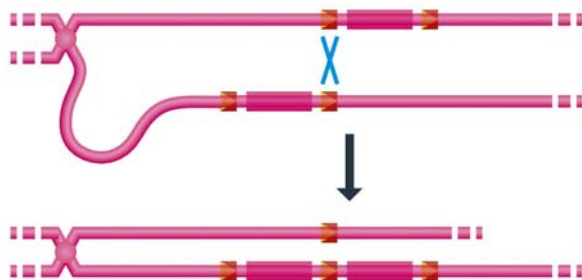


➤ Apert syndrome

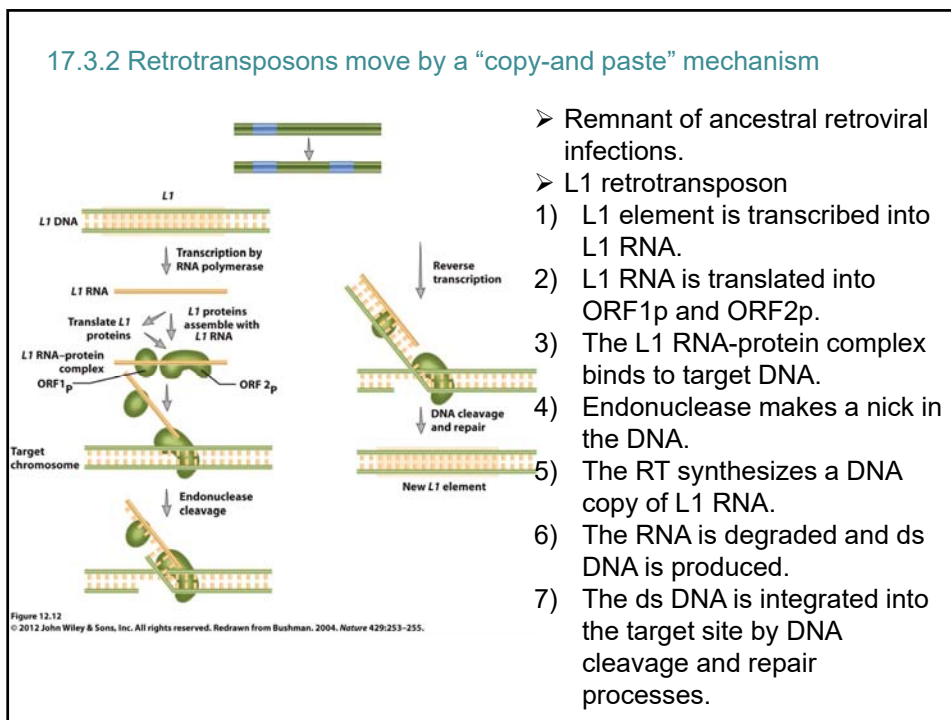
- 1) Craniofacial abnormalities and syndactyly (condition wherein two or more digits are fused together) of the hands and feet.
- 2) Usually results from a missense mutation in exon 7 of the fibroblast growth factor receptor II gene.
- 3) In two cases, insertion of an Alu element in or near exon 9 causes the disease.

Disease Box 12.4 Figure 1e
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Unequal sister chromatid exchange



17.3.2 Retrotransposons move by a “copy-and paste” mechanism



- Remnant of ancestral retroviral infections.
- L1 retrotransposon
 - 1) L1 element is transcribed into L1 RNA.
 - 2) L1 RNA is translated into ORF1p and ORF2p.
 - 3) The L1 RNA-protein complex binds to target DNA.
 - 4) Endonuclease makes a nick in the DNA.
 - 5) The RT synthesizes a DNA copy of L1 RNA.
 - 6) The RNA is degraded and dsDNA is produced.
 - 7) The dsDNA is integrated into the target site by DNA cleavage and repair processes.