DNA Repair and Recombination

Tong-Jin Zhao

School of Life Sciences, Xiamen University

Part I. DNA Repair



What We Have Learned Previously

TABLE 5–1 The Three Steps That Give Rise to High-Fidelity DNA Synthesis		
Replication step	Errors per nucleotide added	
5' \rightarrow 3' polymerization	1 in 10 ⁵	
$3' \rightarrow 5'$ exonucleolytic proofreading	1 in 10 ²	
Strand-directed mismatch repair	1 in 10 ³	
Combined	1 in 10 ¹⁰	

The third step, strand-directed mismatch repair, is described later in this chapter. For the polymerization step, "errors per nucleotide added" describes the probability that an incorrect nucleotide will be added to the growing chain. For the other two steps, "errors per nucleotide added" describes the probability that an error will not be corrected. Each step therefore reduces the chance of a final error by the factor shown.

Spontaneous DNA Damage

DNA lesion	Number repaired in 24 h	
Hydrolysis		
Depurination	18,000	
Depyrimidination	600	
Cytosine deamination	100	
5-Methylcytosine deamination	10	
Oxidation		
8-oxo G	1500	
Ring-saturated pyrimidines (thymine glycol, cytosine hydrates)	2000	
Lipid peroxidation products (M1G, etheno-A, etheno-C)	1000	
Nonenzymatic methylation by S-adenosylmethionine		
7-Methylguanine	6000	
3-Methyladenine	1200	
Nonenzymatic methylation by nitrosated polyamines and peptides		
O ⁶ -Methylguanine	20–100	

Depurination



Deamination



$G \equiv C \longrightarrow A = U$

Formation of Pyrimidine Dimers



C G

HI.

How Chemical Modifications of Nucleotides Produce Mutations





What Are the Consequences?

----- Various genetic diseases and cancer



Part I

Example 1

What Are the Consequences?

----- Various genetic diseases and cancer



Getting tanned



Skin cancer





Ames Test for Carcinogens, Based on Their Mutagenicity



How Does a Living Organism Respond to Mutations?

— DNA Repair

Types of DNA Repair System in E. Coli

- Mismatch repair
- Base-excision repair
- Nucleotide-excision repair
- Direct repair

Which Strand to Repair?





N⁶-Methyladenosine

In *E. Coli*, Dam methylase methylates DNA at the N⁶ position of all adenines within (5')GATC sequence.

Methylation and Mismatch Repair





Methyl-Directed Mismatch Repair



MutH has a site-specific endonuclease activity. It is inactive until the complex encounters a hemimethylated GATC sequence. MutH cuts at the 5'-side of the G in the GATC sequence in the unmethylated strand, which marks the strand for repair.

Methyl-Directed Mismatch Repair -continued



Proteins Required for Methyl-Directed Mismatch Repair

Dam methylase MutH, MutL, MutS proteins DNA helicase II SSB DNA polymerase III Exonuclease I Exonuclease VII RecJ nuclease Exonuclease X DNA ligase

Mismatches

Base-Excision Repair



Nucleotide-Excision Repair



Direct Repair

DNA photolyase

Two chromophores: MTHFpolyglu (folate) FADH⁻



Direct Repair of O⁶-methylguanidine



Direct Repair of Alkylated Bases by AlkB



Interaction of Replication Forks with DNA Damage Can Lead to Error-Prone Translesion DNA Synthesis



Translesion DNA Polymerases Can Use Damaged Templates



Translesion DNA Synthesis (TLS) is part of SOS Response.

Two Ways to Repair Double-Strand Breaks



Summary

Mismatch repair in E. Coli is directed by transient nonmethylation of the (5')GATC sequence on the newly synthesized strand.

Base-excision repair systems recognize and repair damage caused by environmental reagents and spontaneous reaction of nucleotides.

Nucleotide-excision repair systems recognize and remove a variety of bulky lesion and pyrimidine dimers.

Direct repair system works through direct reversal of the reaction causing the damage.

In bacteria, error-prone translesion DNA synthesis occurs in response to very heavy DNA damage.

Two ways to repair double-strand breaks: nonhomologous end joining and homologous recombination.

Part II. DNA Recombination



Genetic Recombination

Homologous Genetic Recombination

Site-Specific Recombination

DNA Transposition

Homologous Recombination Has Several Functions

In bacteria, it is mainly recombinational repair, which is directed at the reconstruction of replication forks stalled at the site of DNA damage.

In eukaryotes, homologous recombination has several roles in replication and cell division, including the repair of stalled replication forks. Recombination occurs with highest frequency during meiosis.

Repair of a Broken Replication Fork by Homologous Recombination



Meiosis in Animal Germ Cells



Crossing Over



Hot Spots

Homologous Recombination Has at Least Three Identifiable Functions

- It contributes to the repair of several types of DNA damage
- It provides, in eukaryotic cells, a transient physical link between chromatids that promotes the orderly segregation of chromosomes at the first meiotic cell division
- It enhances genetic diversity in population

Recombination During Meiosis



Holliday Intermediates Are a Feature of Homologous Recombination Pathways in All Organisms



Branch Migration



Recombination Requires a Host of Enzymes and Other Proteins

• In E. Coli, RecB, RecC and RecD form a heterotrimeric RecBCD, which has both helicase and nuclease activities.

Helicase and Nuclease Activities of the RecBCD Enzyme



RecB and RecD are helicase motors: RecB moves 3'-to-5' on one strand RecD moves 5'-to-3' on the other strand

Chi: (5')GCTGGTGG Chi binds tightly to a site on RecC subunit

Recombination Requires a Host of Enzymes and Other Proteins

- In E. Coli, RecD, RecC and RecD form a heterotrimeric RecBCD, which has both helicase and nuclease activities.
- The RecA promotes all central steps in homologous recombination process: the pairing of two DNAs, formation of Holliday intermediates, and branch migration.

The Active Form of RecA Protein Is an Ordered Helical Filament



Up to several thousand subunits



24-subunit filament

RecA Filaments Are Extended or Disassembled in the 5'-to-3' Direction



Filament Assembly Is Assisted by RecF, RecO and RecR, and Inhibited by RecX



RecA-Promoted DNA Strand Exchange in vitro



Model for RecA-Mediated DNA Strand Exchange



Recombination Requires a Host of Enzymes and Other Proteins

- In E. Coli, RecD, RecC and RecD form a heterotrimeric RecBCD, which has both helicase and nuclease activities.
- The RecA promotes all central steps in homologous recombination process: the pairing of two DNAs, formation of Holliday intermediates, and branch migration.
- The RuvA and RuvB (<u>r</u>epair of <u>UV</u> damage) form a complex that binds to Holliday complex, displaces RecA proteins, and promotes branch migration at higher rates than does RecA.
- Nucleases, often called resolvases, specifically cleave Holliday intermediates. RuvC is one of the at least two such nucleases in E. Coli.

All Aspects of DNA Metabolism Come Together to Repair Stalled Replication Forks



I. Recombinational DNA Repair of DNA Lesion in Stalled Replication Fork



II. Recombinational DNA Repair of DNA Nick in Stalled Replication Fork



Origin-Independent Replication Restart

Site-Specific Recombination

- Each site-specific recombination system consists of
 - An enzyme called recombinase
 - A short (20 to 200 bp), unique DNA sequence where the recombinase acts
 - One or more of the auxiliary proteins that regulate the timing or outcoming
- Two general classes of site-specific recombination systems
 - Rely on Tyr in the active site
 - Rely on Ser in the active site

A Site-Specific Recombination Reaction





Active site: Tyr

Site-Specific Recombination

In systems that employ an active-site Ser residue, both strands are cut concurrently and rejoined to the new partners without forming Holliday intermediate.

The exchange is always reciprocal and precise, regenerating the recombination sites when the reaction is complete.

The two recombination sites align in the same orientation during the recombination reaction.

Effects of Site-Specific Recombination



Outcomes Depends on the Location and Orientation of the Recombination Sites

• Two sites on the same DNA: inversion or deletion

- Two sites on different DNAs: Intermolecular
 - If one or both are circular : Insertion

Integration and Excision of Bacteriophage λ DNA at the Chromosomal Target Site



Complete Chromosome Replication Can Require Site-Specific Recombination





Transposition



Barbara McClintock 1902–1992



Jumping genes in maize

Transposable Genetic Elements Move from One Location to Another

- Transposition is a recombination that allows the movement of transposable elements, or transposons.
- DNA sequence homology is usually not required for this movement, called transpostition; the new location is determined more or less randomly.
- Bacteria have two classes of transposons.
 - Insertion sequences (simple transposons) contain only the sequences required for transposition and transposases.
 - Complex transposons contain one or more genes in addition to those needed for transposition.

Duplication of the DNA Sequence at a Target Site When a Transposon Is Inserted



Two General Pathways for Transposition: Direct and Replicative



Immunoglobulin Genes Assemble by Recombination



V (variable); J (joining); C (constant).

1.5x10⁷ possible IGs in human.

Mechanism of Immunoglobulin Gene Rearrangement



RSS: Recombinational signal sequence Just beyond each V segment and just before each J segment

RAG: Recombination Activating Gene

Summary

- DNA sequences are rearranged in recombinational reactions
- Homologous genetic recombination can take place between any DNA molecules that share sequence homology.
 - In meiosis, it helps to ensure accurate chromosomal segregation and create genetic diversity.
 - In both bacteria and eukaryotes, it serves in the repair of stalled replication forks.
 - A Holliday intermediate forms during homologous recombination.
- Site-specific recombination occurs at specific target sequences and this process can also involve Holliday intermediate.
- In virtually any cells, transposons use recombination to move within or between chromosomes.