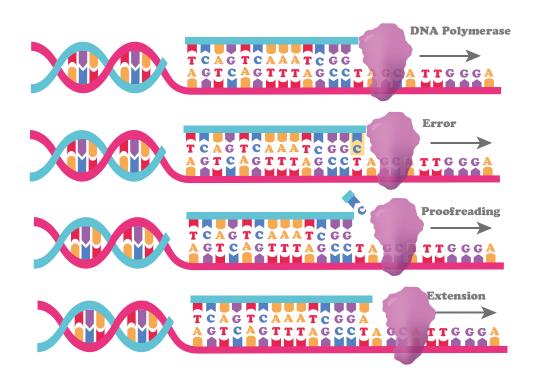
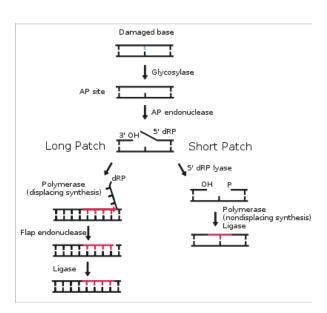
## **CONCEPT: DNA REPAIR**

- - $\hfill\Box$  DNA polymerase proofreading quickly fixes a mispaired base
  - $\hfill \Box$  Enzymes can reverse damaged DNA
    - Many different enzymes exist to counteract damaged DNA
      - Ex: CPD photolyase is an enzyme that repairs dimers caused from UV light



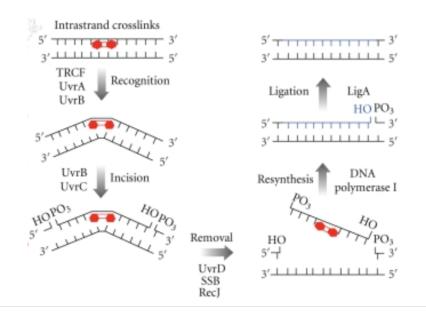
## Repair Pathways

- □ Base-excision repair (BER) removes and replace damaged nucleotides
  - DNA glycoslyases are enzyme that remove the damaged base from the sugar
  - A deoxyribophosphodiesterase enzyme removes a stretch of neighboring DNA
  - DNA polymerase fills the gap with complementary nucleotides
  - DNA ligase seals the new nucleotides into the backbone

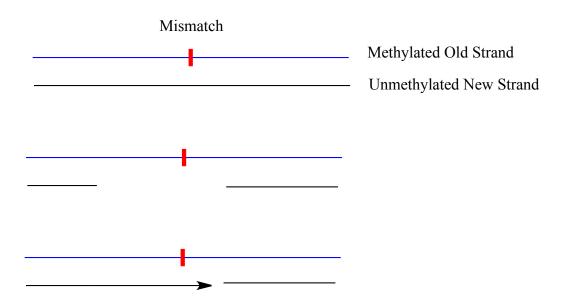


- □ Nucleotide-excision repair (NER) repairs damage that \_\_\_\_\_\_ the DNA helix
  - 1. Global genomic NER repairs damage anywhere in the genome
  - 2. Transcription-coupled NER repairs only transcribed regions of DNA
  - Damaged base is recognized by proteins, which recruit more proteins
  - A 30 nucleotide segment of DNA is removed from the damaged area
  - DNA polymerase fills the gap, and DNA ligase seals the new nucleotides to the backbone
  - Distortions in NER cause diseases like Xoderma pigmentosum and cockayne syndrome
    - Causes light sensitivities

## **EXAMPLE:**



- □ **Mismatch repair** repairs damaged from insertions or deletions that occur immediately following replication
  - First, the mismatched base is recognized, and proteins are recruited to the area
  - The methylation status of the DNA strands is used to determine which is the mismatched base
    - Methylated strand = old strand, unmethylated strand = new strand
  - DNA polymerase fills the gap, and DNA ligase seals the new nucleotides to the backbone



## **Translesion Synthesis**

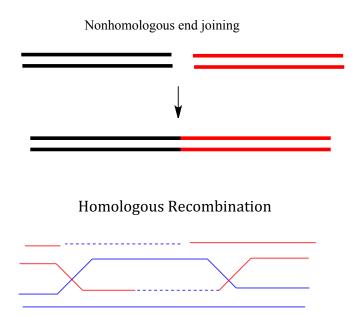
- □ **Translesoin DNA synthesis** is a last resort mechanism the cell uses to prevent mutations from causing death
  - DNA damage causes DNA polymerase to \_\_\_\_\_\_- while replicating
  - Translesion (bypass) polymerases are recruited to the area
    - These polymerases can overcome helical distortions and continue replicating
    - But they have problems including: high error rate, no proofreading, and only add few nt at a time
  - So translesion polymerases unblock stalled replication forks to allow for replication to continue
  - They don't fix the damage that caused the stall and they can miss mutations without proofreading

#### **EXAMPLE:**

# 

## **Double Strand Breaks**

- Double strand breaks can be repaired in \_\_\_\_\_ ways
  - □ Nonhomologous end joining connects two broken ends together
    - Proteins recognize DNA damage
    - Other proteins are recruited which trim nucleotides off the damaged ends
    - DNA ligase connects the two broken ends
    - This type of repair occurs outside of S phase
  - □ **Homologous recombination** repairs double strand breaks directly after replication
    - Synthesis-dependent strand annealing uses sister chromatids as a template to repair the broken strands
    - Process is similar to crossing over, but uses sister chromatids instead of nonsister chromatids



## PRACTICE:

- 1. Which of the following repair pathways repairs damage that causes distortions in the double helix?
  - a. Base Excision Repair
  - b. Nucleotide Excision Repair
  - c. Mismatch Repair
  - d. Homologous Recombination

- 2. Which of the following repair pathways uses a methylated strand of DNA to correct DNA damage?
  - a. Base Excision Repair
  - b. Nucleotide Excision Repair
  - c. Mismatch Repair
  - d. Homologous Recombination

3.	True or False: Translesion DNA synthesis is the first mechanism the cell uses to repair DNA damage?  a. True  b. False
4.	Which of the following pathways is an error-free way to repair double-stranded breaks?  a. Homologous recombination  b. Non-homologous end joining  c. Translesion synthesis  d. Mismatch repair