

# MOD1 – DNA ENGINEERING

Engelward, Spring 2008

**Day 3**

## **About the experiments in Mod1**

- how is recombination used to fix double strand breaks
- how your two-plasmid assay works

## **Agarose Gels – How do we ‘look’ at DNA?**

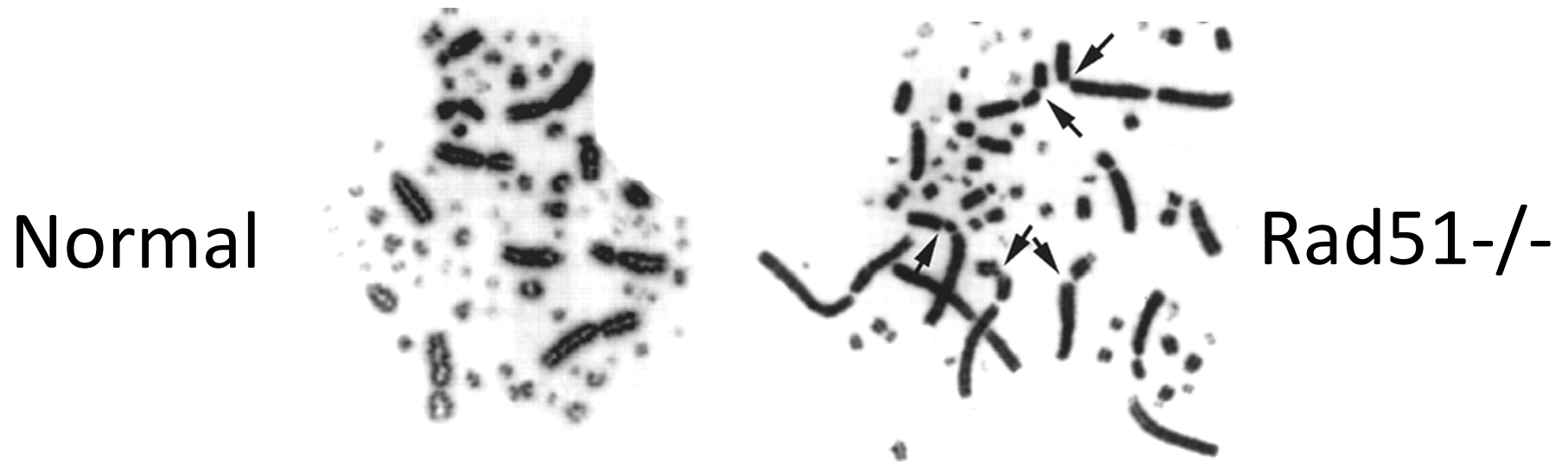
## **Anticipating Potential Problems & Pitfalls**

# Background & Significance:

“Homology-Directed Repair” for double strand breaks

You will need to understand this material in order to write your final report.

# Why you owe Your Life to Homologous Recombination...

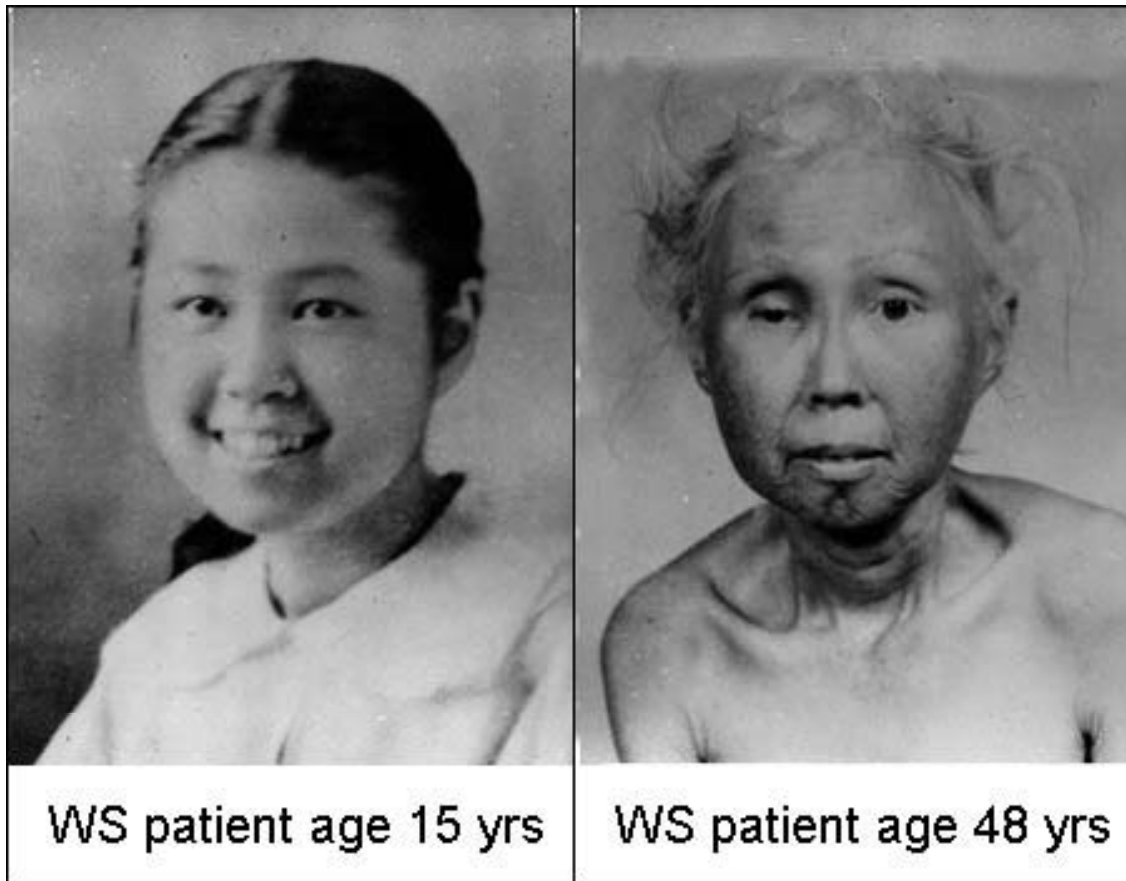


Turn Off Homologous Recombination  
→ Chromosomes Fall Apart

Sonada *et al.*, *EMBO J.* **17**, 598–608 (1998).

# *Why you owe Your Youthfulness to Homologous Recombination...*

Loss of Helicase → Faulty Recomb.



Werner's  
Syndrome

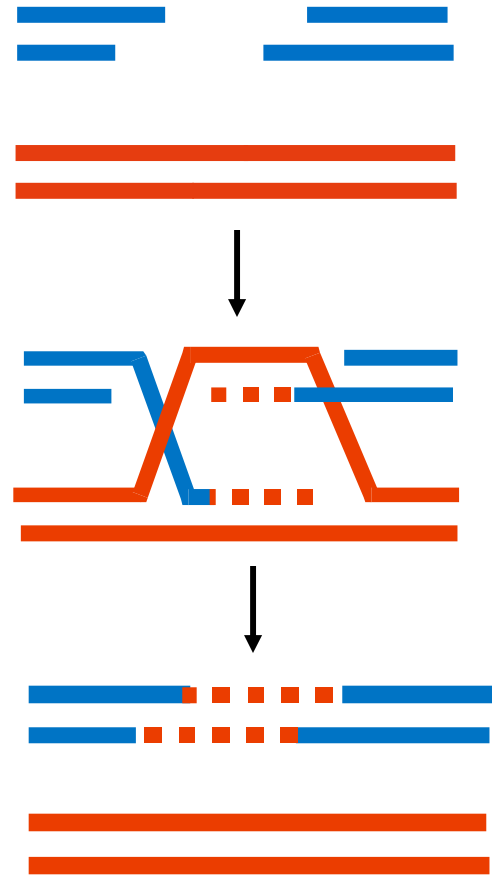
# Double Strand Breaks

NHEJ



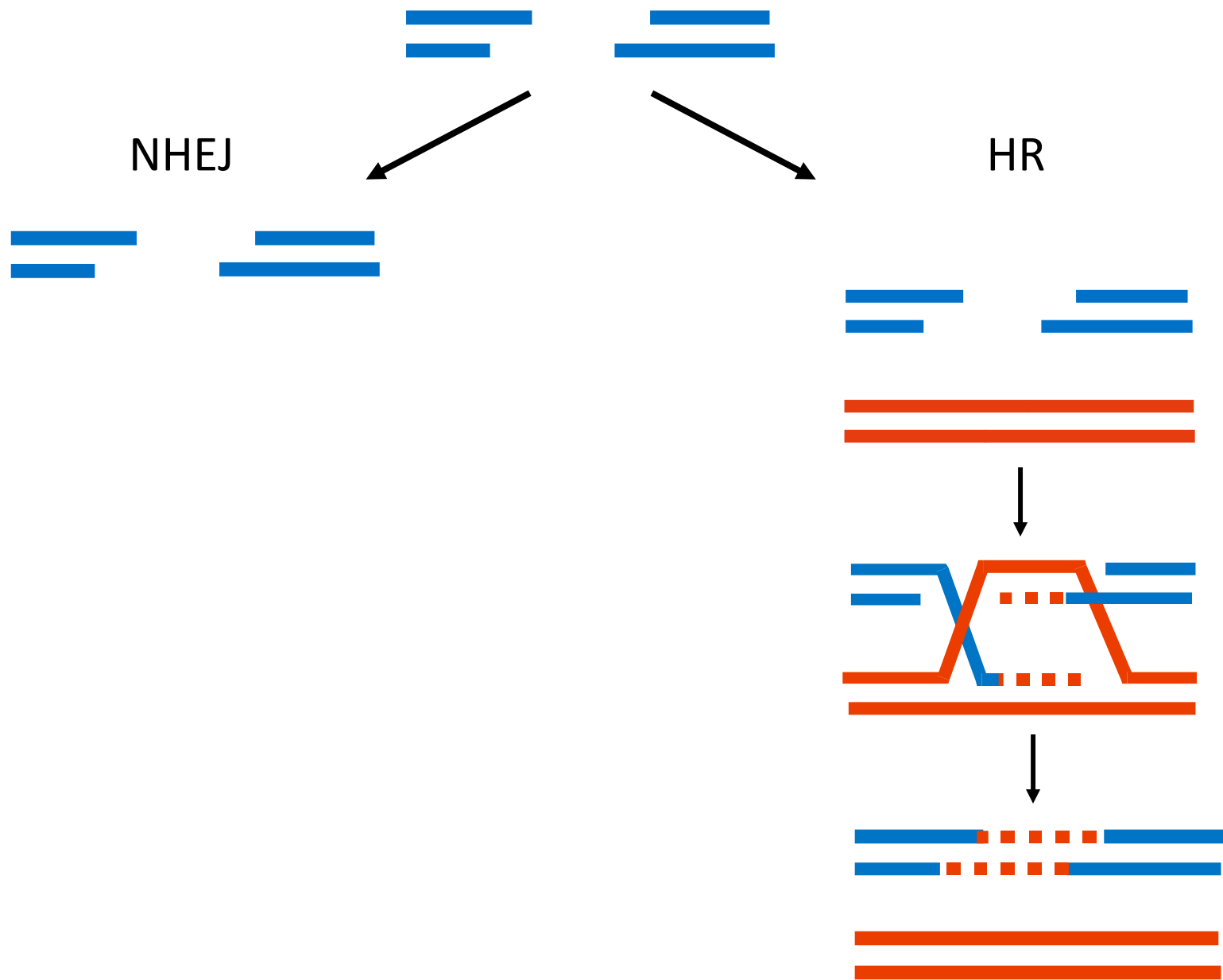
KU heterodimer  
(Ku70&80)  
DNA-PKcs  
DNA ligase IV  
XRCC4

HR

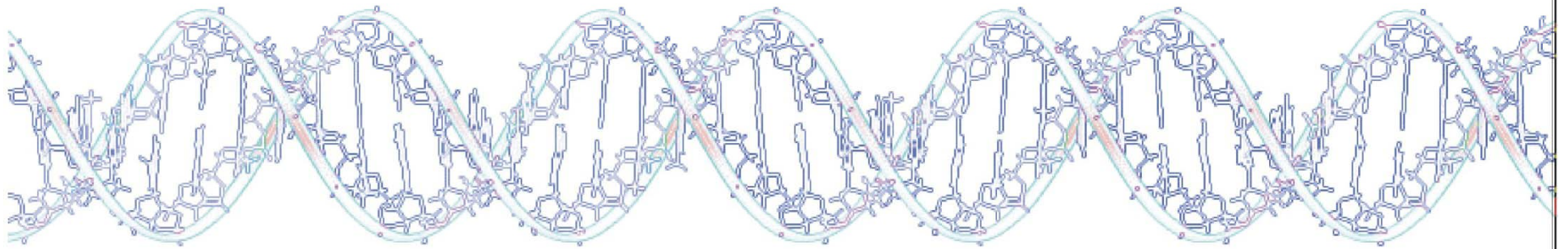


**See NHEJ Animation by Justin Lo**

# Double Strand Breaks

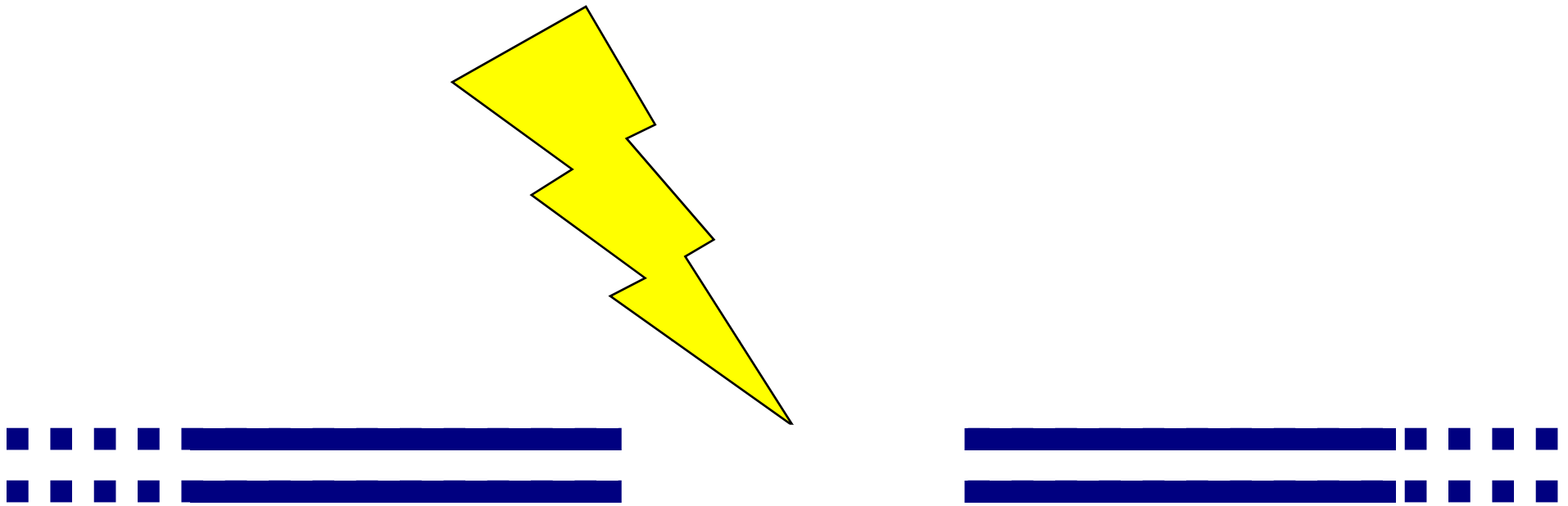






=





Imagine HR is initiated by the fragment on the left....

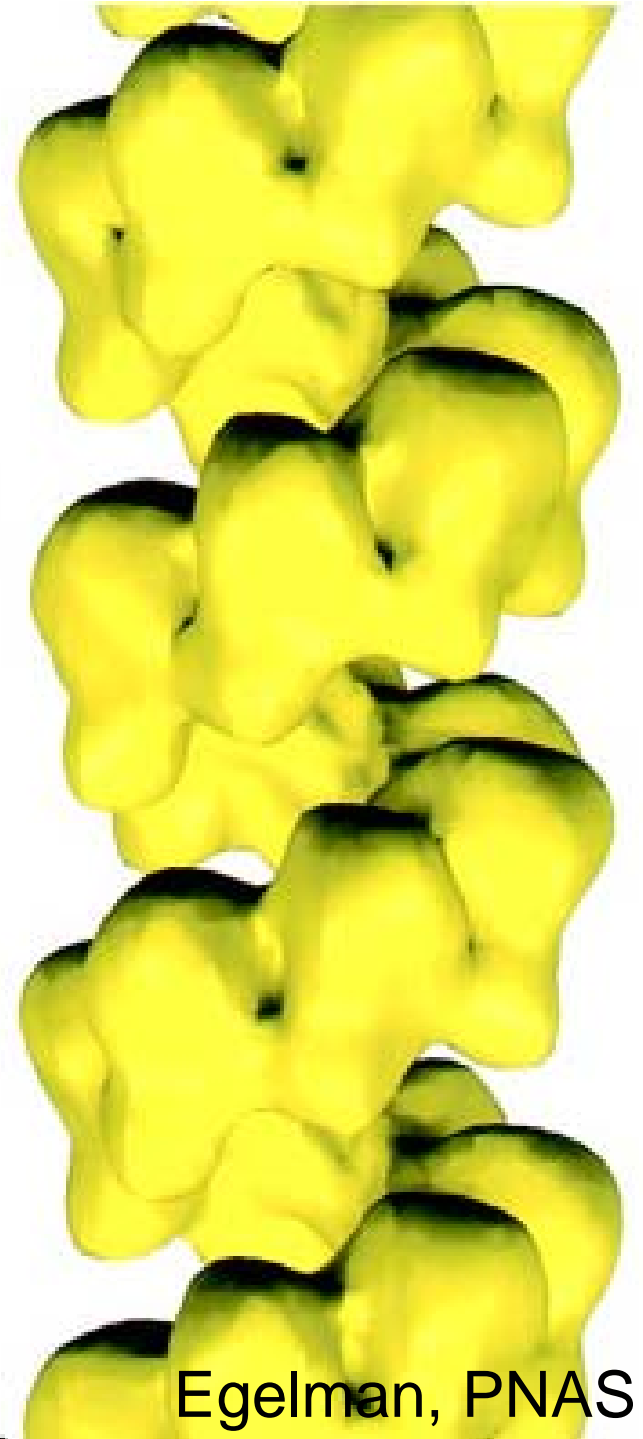
Step 1: A double stranded end has been created



Step 2: Resect the end to create a 3' overhang



Step 3: Create a nucleoprotein filament capable of homology searching



Egelman, PNAS

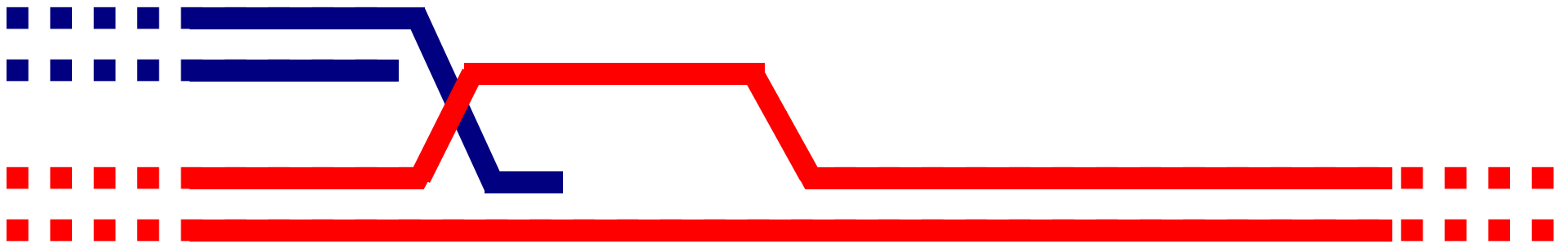


Step 4: Search and Invade



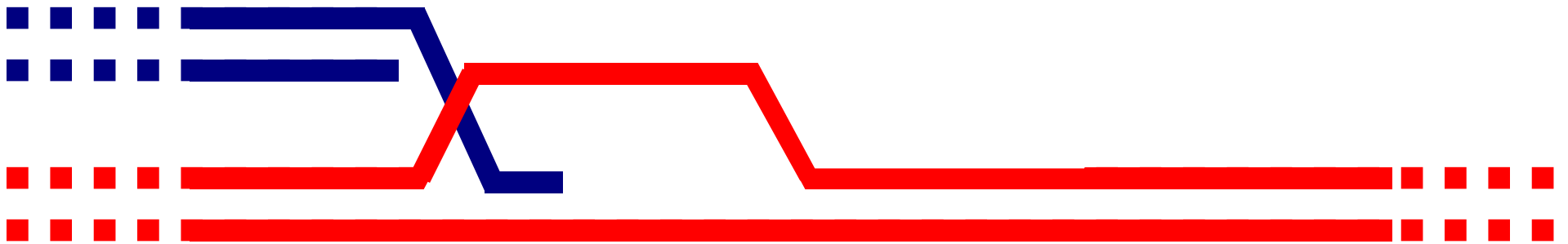


Step 4: Search and Invade

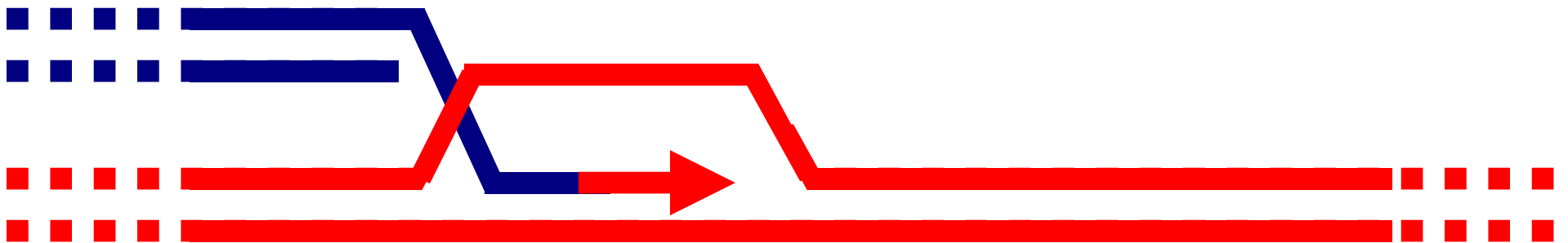


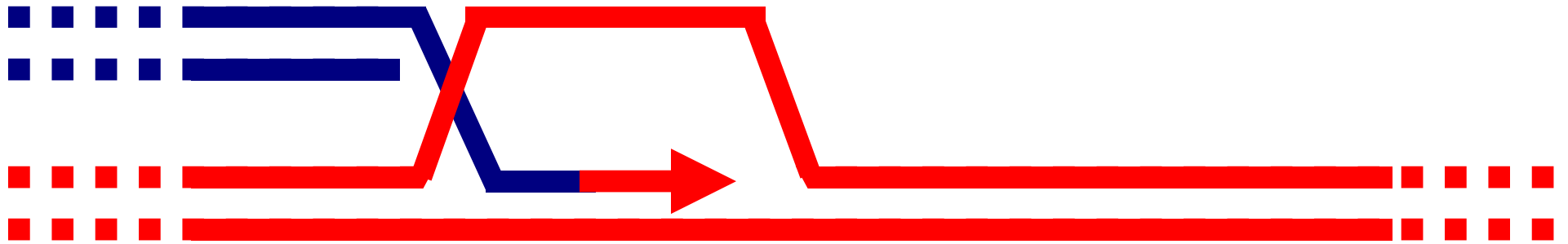


Step 4: Search and Invade

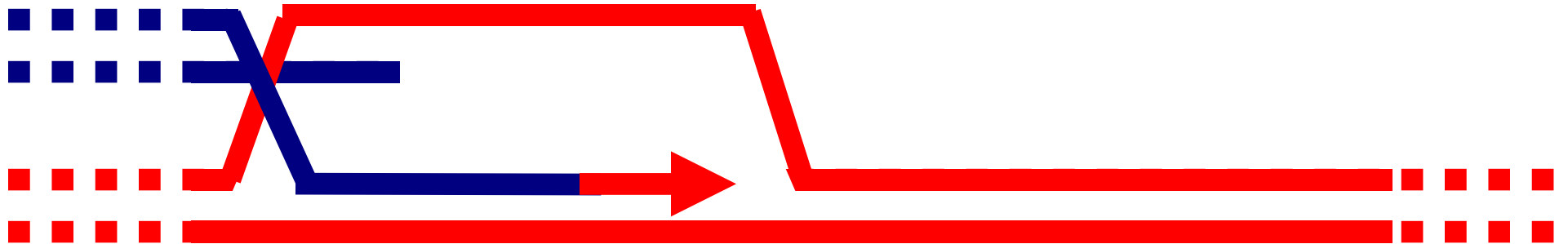


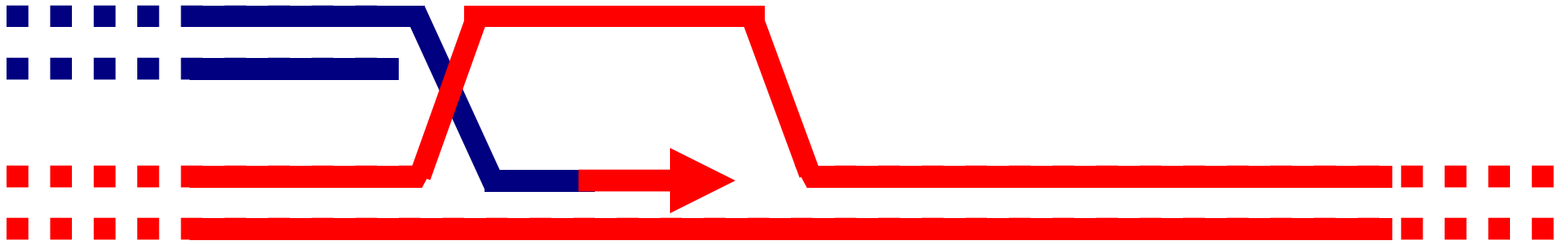
Step 5: Polymerize DNA using invading strand with 3'OH as a primer and the homologous donor DNA as a template



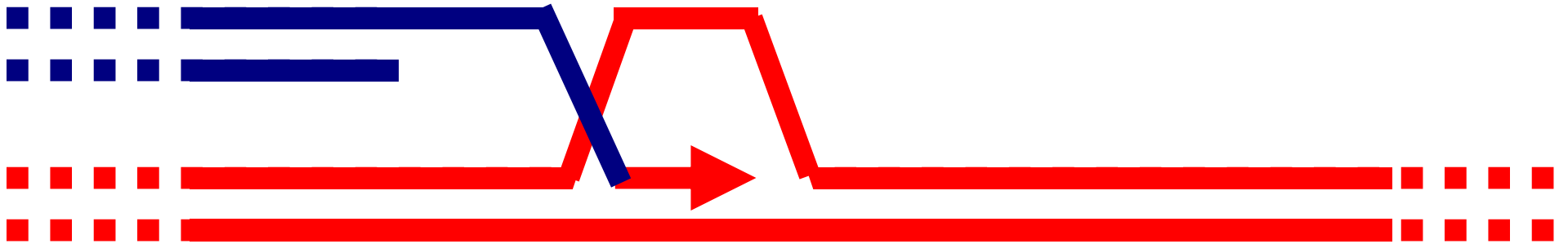


Step 6: Branch Migration (Backwards)

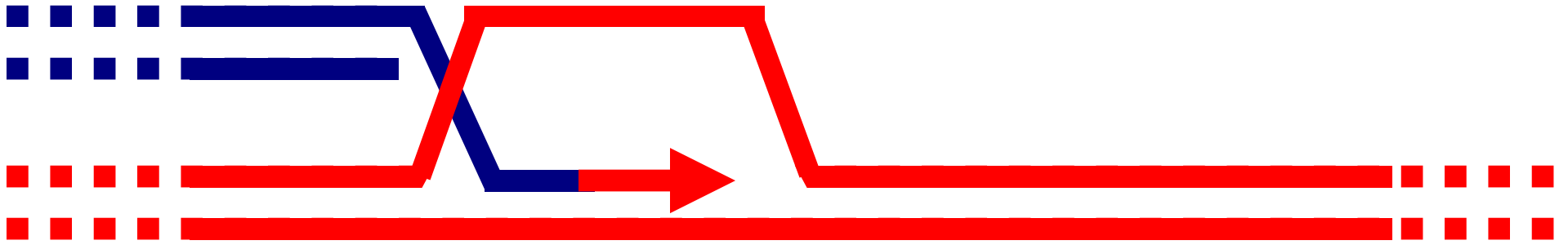




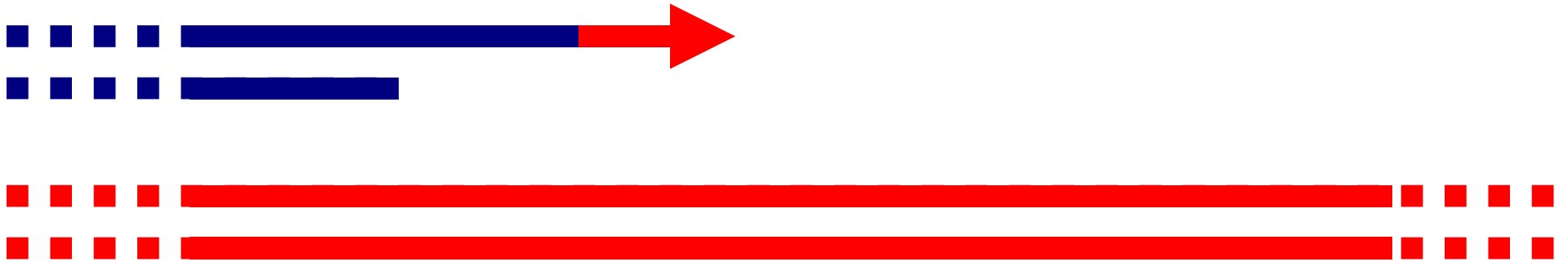
Step 6: Branch Migration (Forwards)

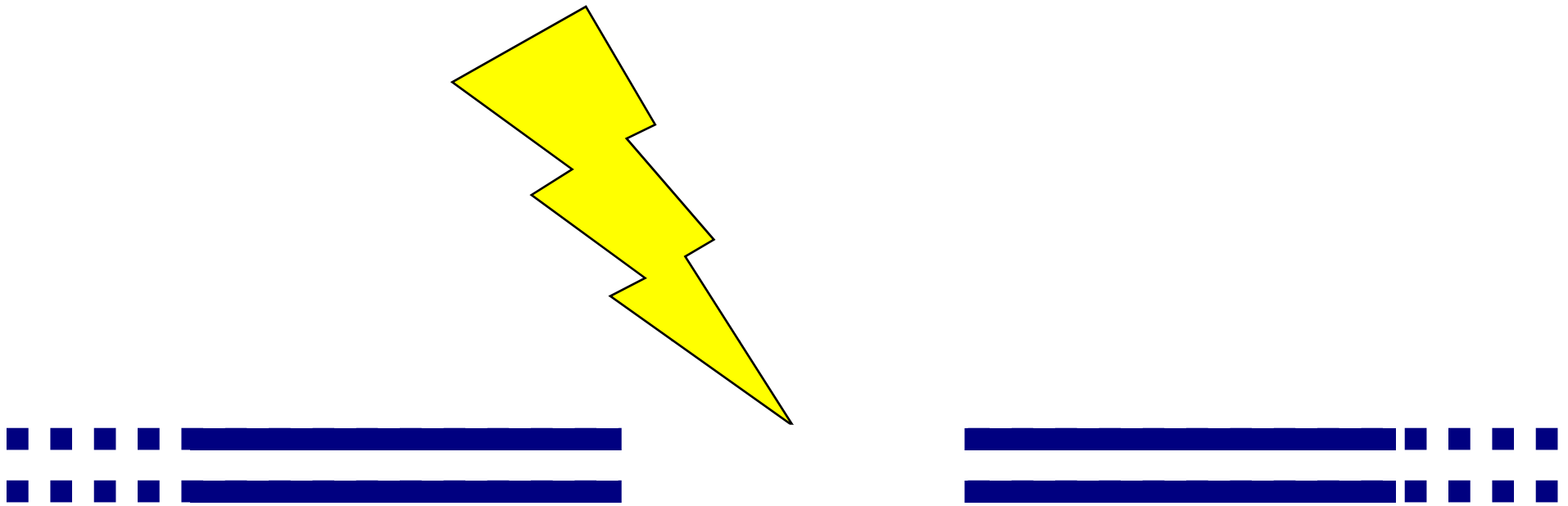






Step 6: Possible Release





This process started with a two-ended DSB...



↓ Now let's imagine the same thing happened at the other end...



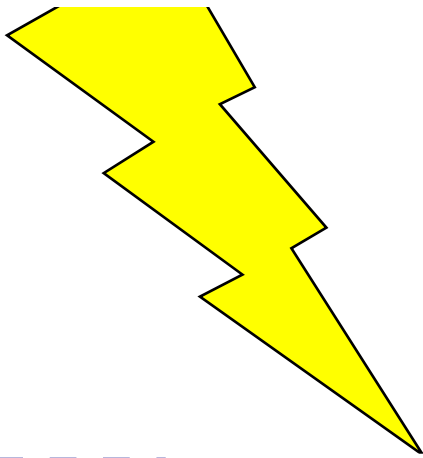
↓ Annealing





Final Steps: Filling, Trimming, Ligating

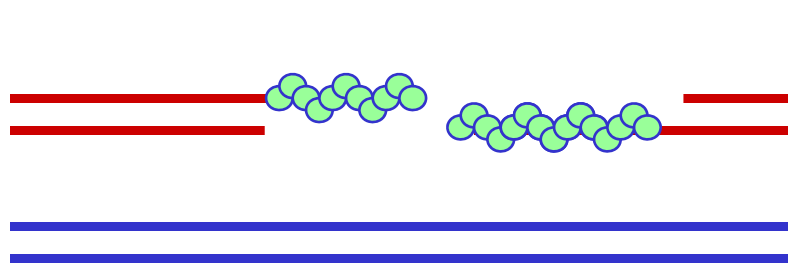




+



**See SDSA Animation  
by Justin Lo**

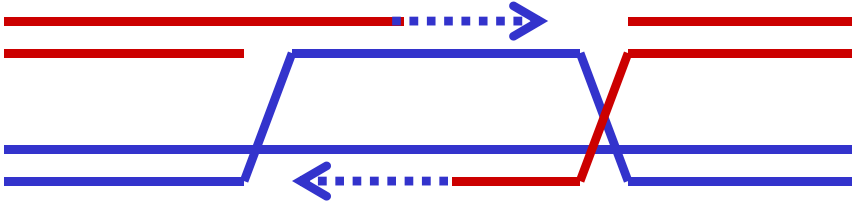
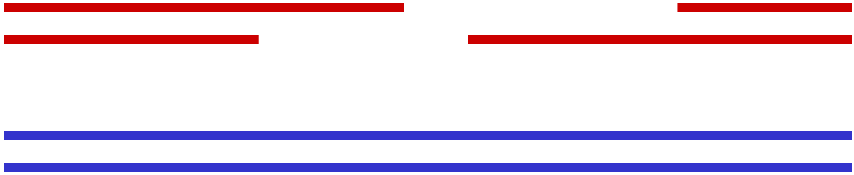


*E. coli*: RecA\*

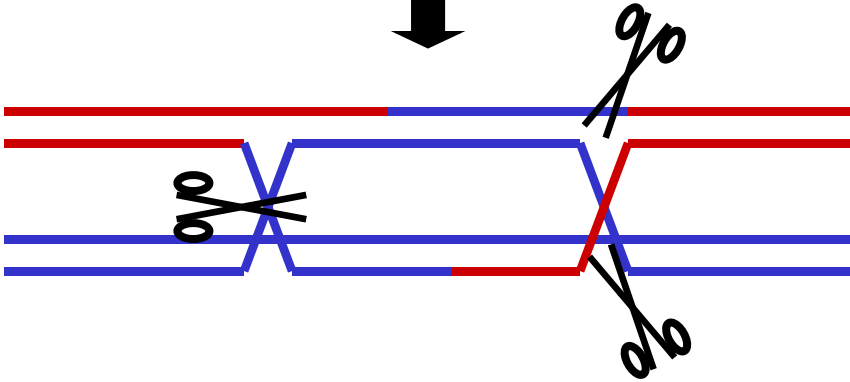
Yeast: Rad51

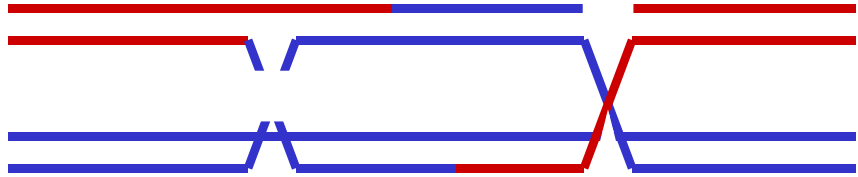
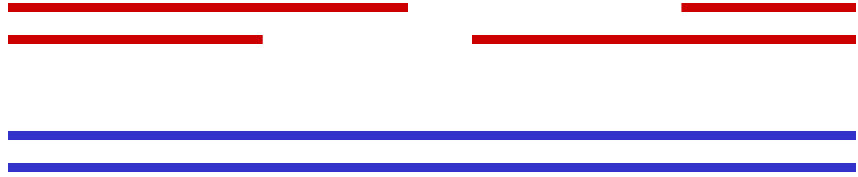
Mammals: RAD51

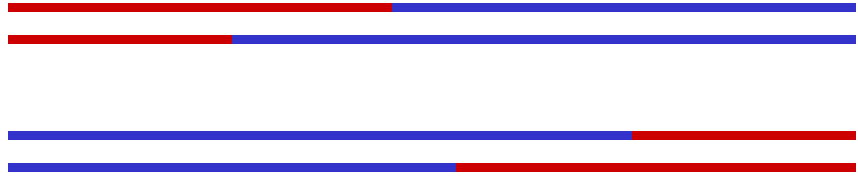
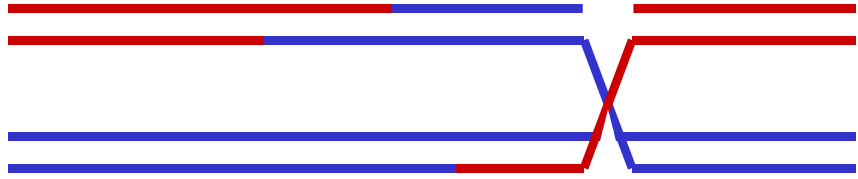


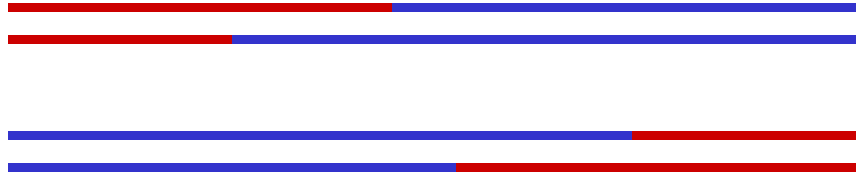
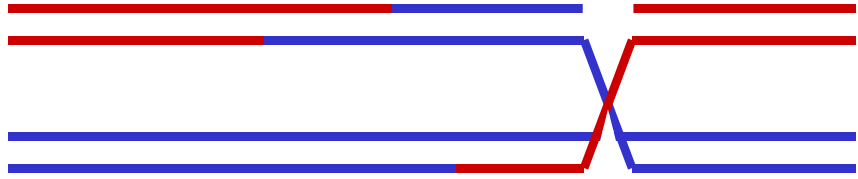






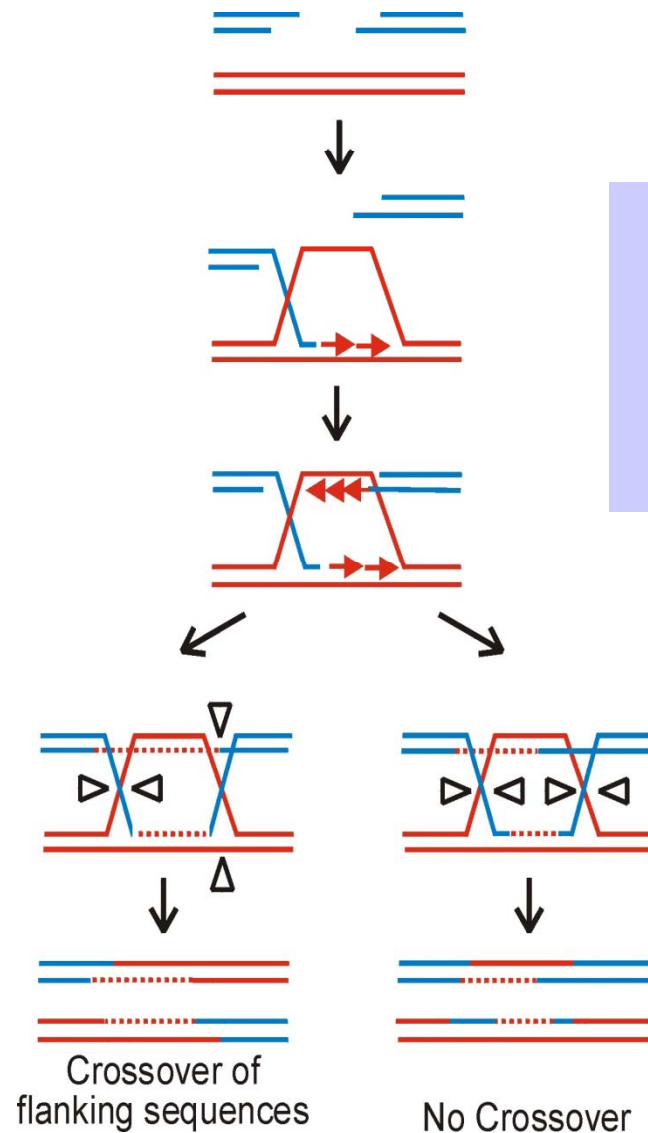






**See Prototypic Model Animation  
by Justin Lo**

# DNA Damage can be repaired by **Homology Directed Repair (HDR)**



This is the 'prototypic' model of repair of how homologous recombination can repair a double strand break

**NOTE: BREAKPOINT  
TURNS FROM BLUE TO  
RED**

Decision to initiate HDR,  
resection of DNA ends

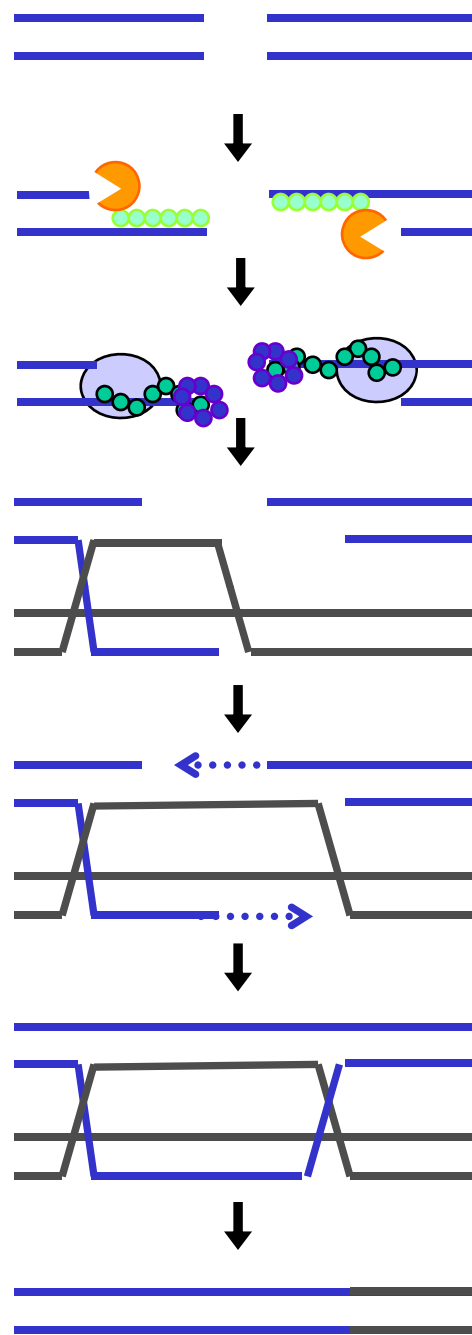
Displacement of RPA &  
Loading Rad51

Homology searching,  
histone remodeling,  
& invasion

Holliday junction migration,  
inhibition by mismatches

Repair synthesis,  
Holliday junction migration,  
Possible resolution  
without junction cleavage

Junction resolution  
Repair of mismatches



ATM, ATR, cAbl, Chk1, p53,  
BRCA1, Fanconi genes, Mre11,  
Rad50, Nbs1, Exonucleases

RPA, Rad52, BRCA2,  
Rad51, Rad51B, Rad51C,  
Rad51D, XRCC2, XRCC3

Rad54, Rad54B, Rad52

MMR, WRN, BLM, Rad54, p53

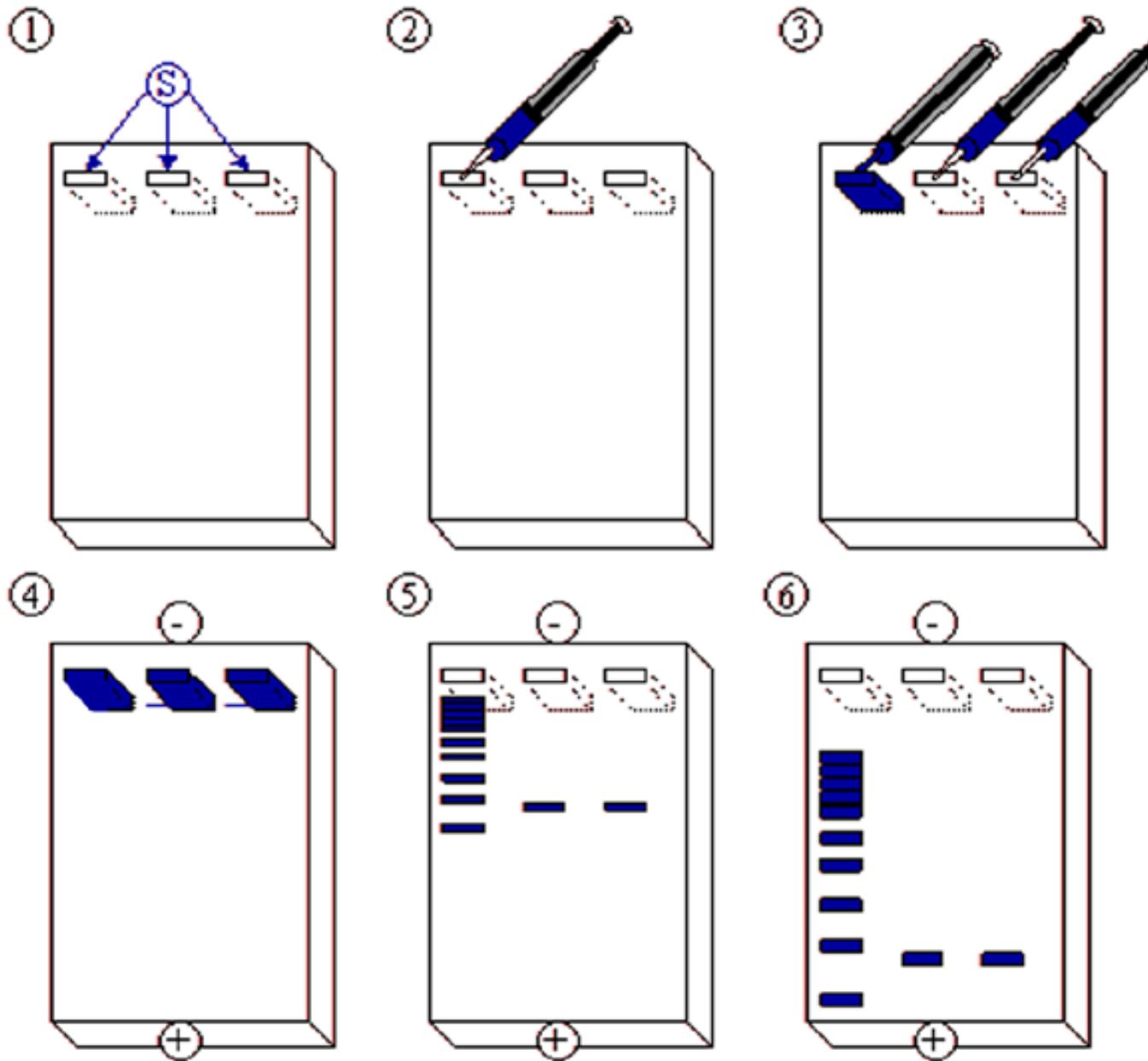
Polymerase(s), topoisomerase(s)  
WRN, BLM, Rad54

Rad51C & possible additional  
proteins; possible resolution by  
topoisomerases; MMR

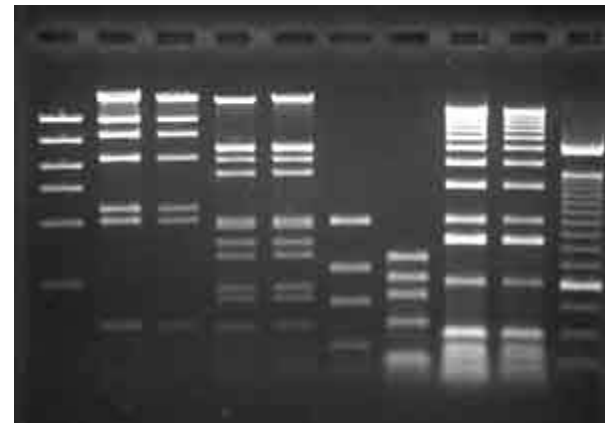
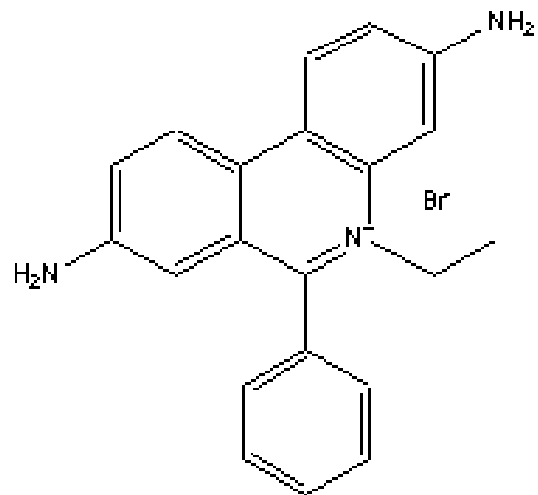
# Agarose Gels & Gel Purification

- How do we 'look' at DNA?
- How do we get our DNA out of a gel?

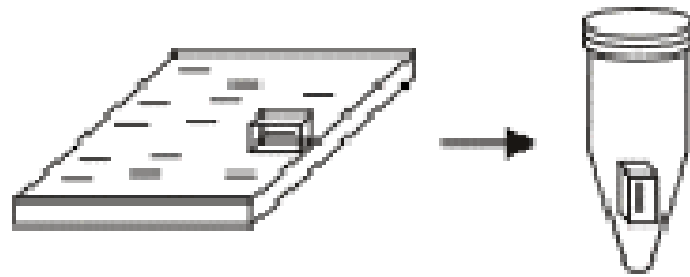




How can you see your DNA?

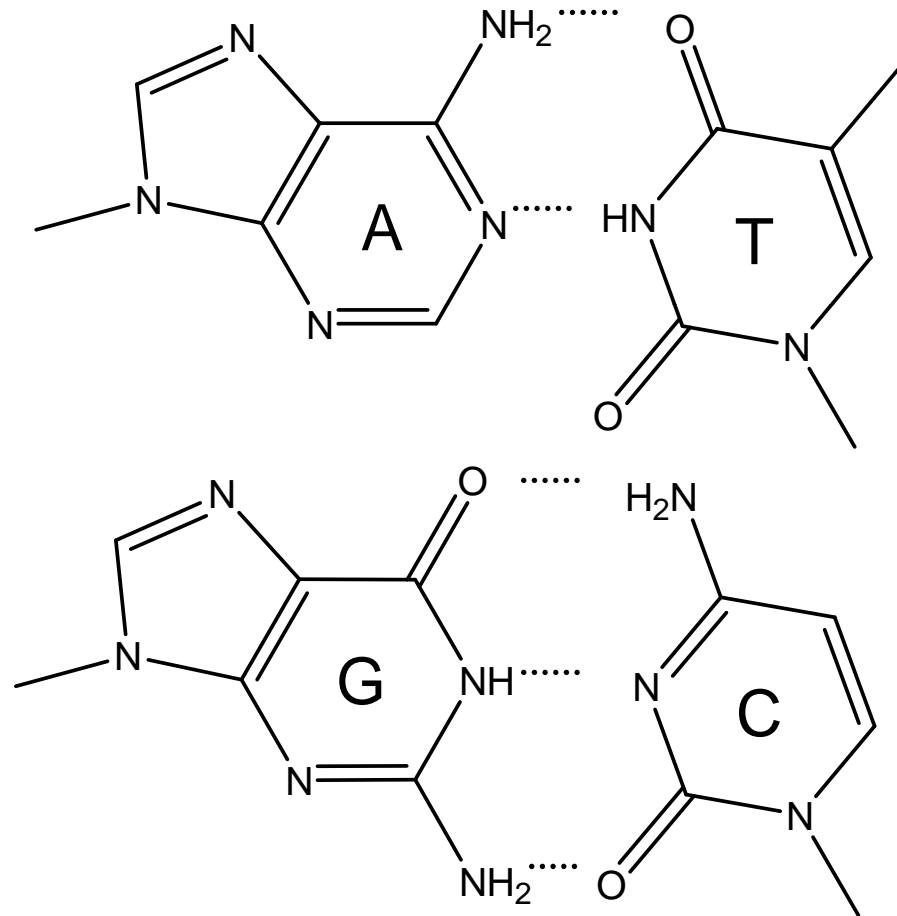


# Gel Purification



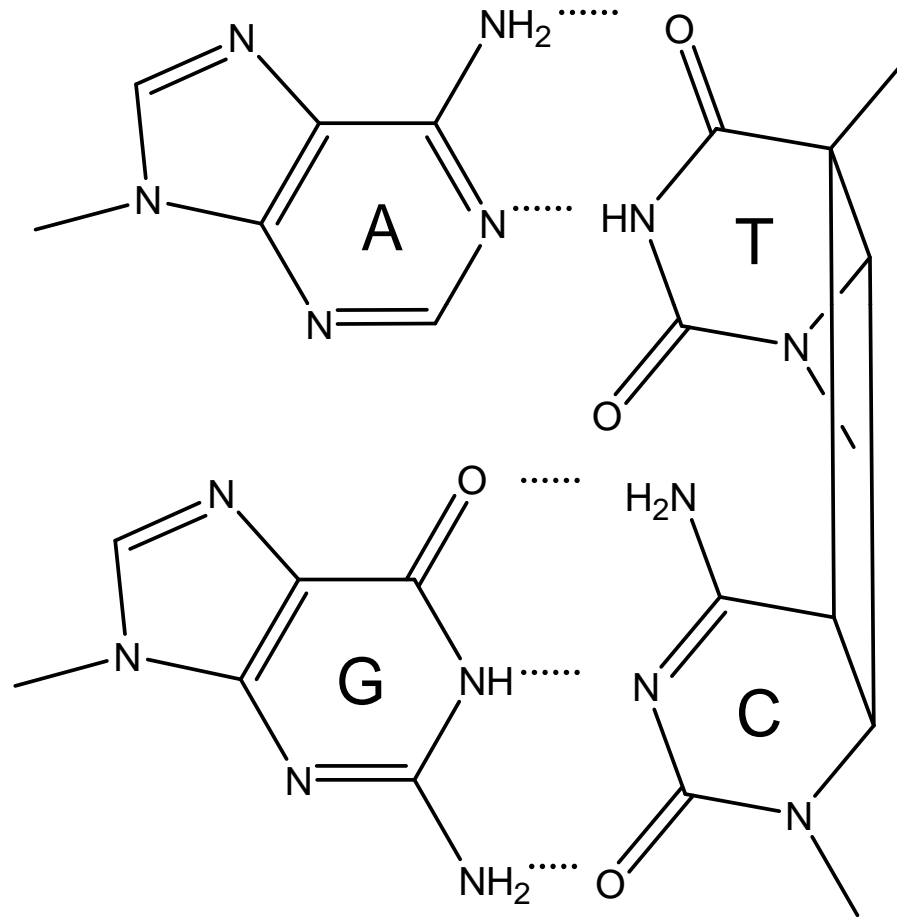
Why do you need to cut out your band fairly quickly?

# Sunlight Damages DNA



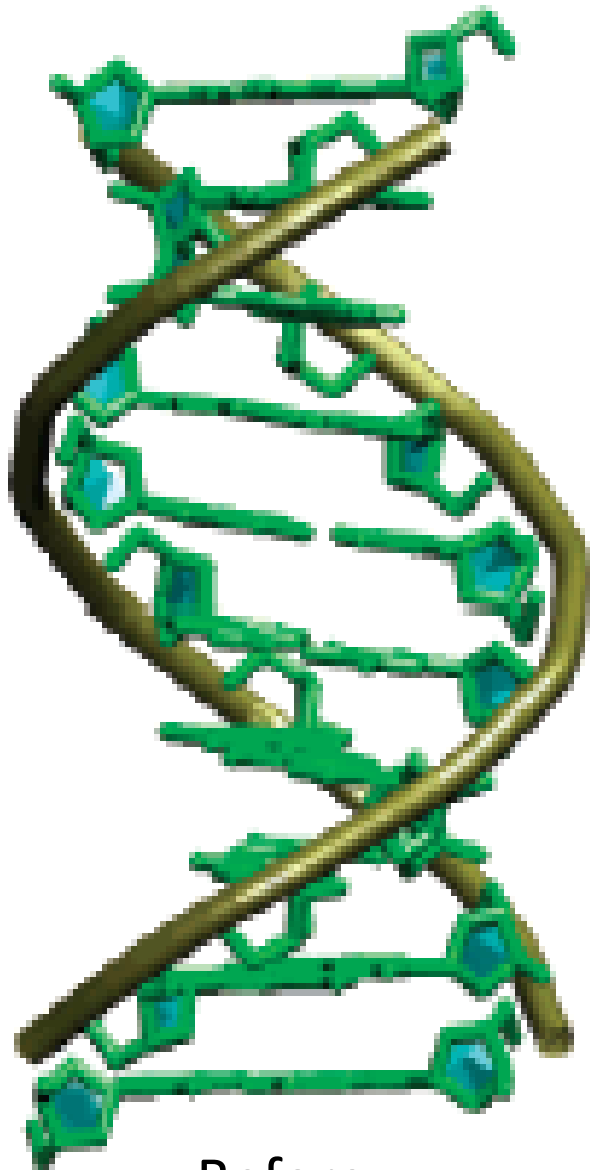
Normal Base Pairs

# Sunlight Damages DNA

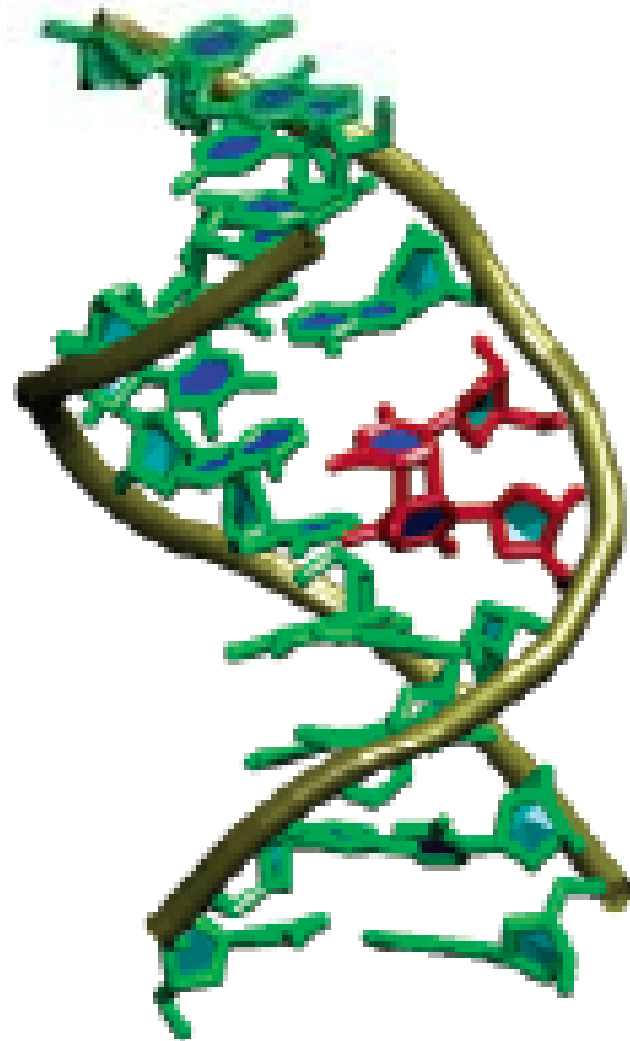


Sunlight-Induces Crosslinks Between Bases

# Sunlight Damages DNA

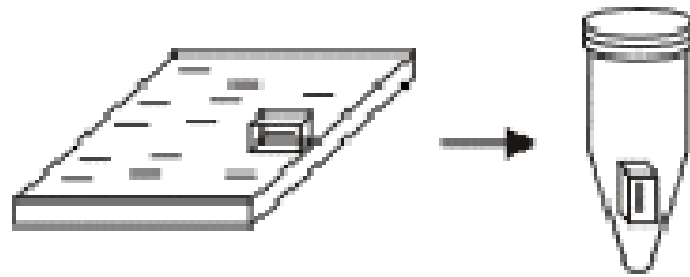


Before



After

# Gel Purification



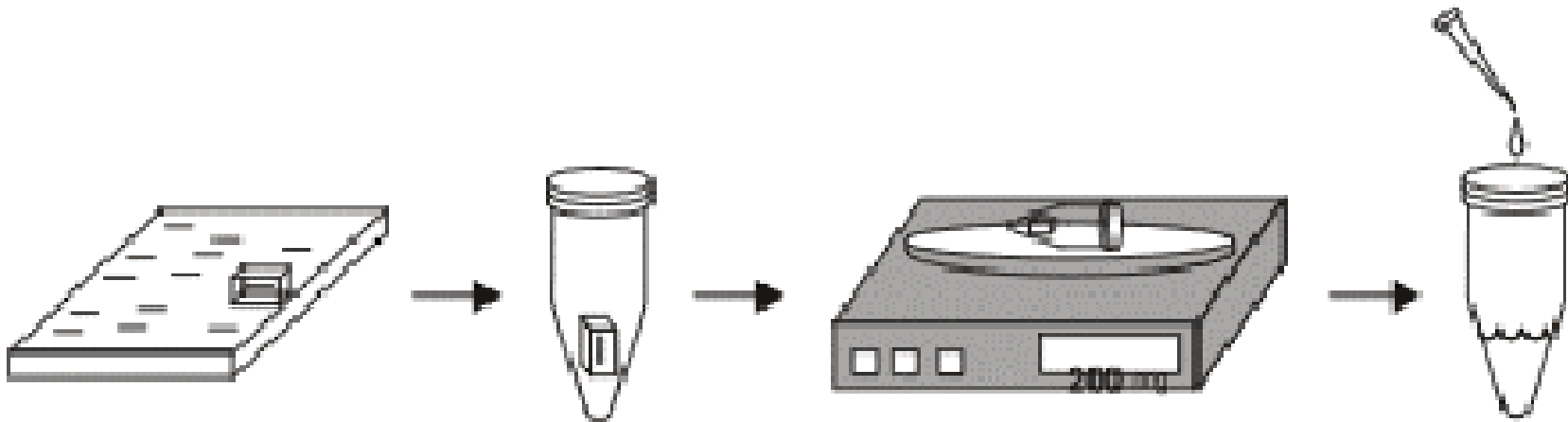
Why do you need to cut out your band fairly quickly?

You will need to dissolve the gel to get the DNA out.. You do this by adding 3 volumes of a gel-dissolving solution.

What does it mean to 'add 3 volumes'?

How can you estimate the volume of your gel slice?





# Agarose Gels – How do we ‘look’ at DNA?

- Loading

- Standards

- Parameters that affect migration

  - gel concentration

  - length of DNA

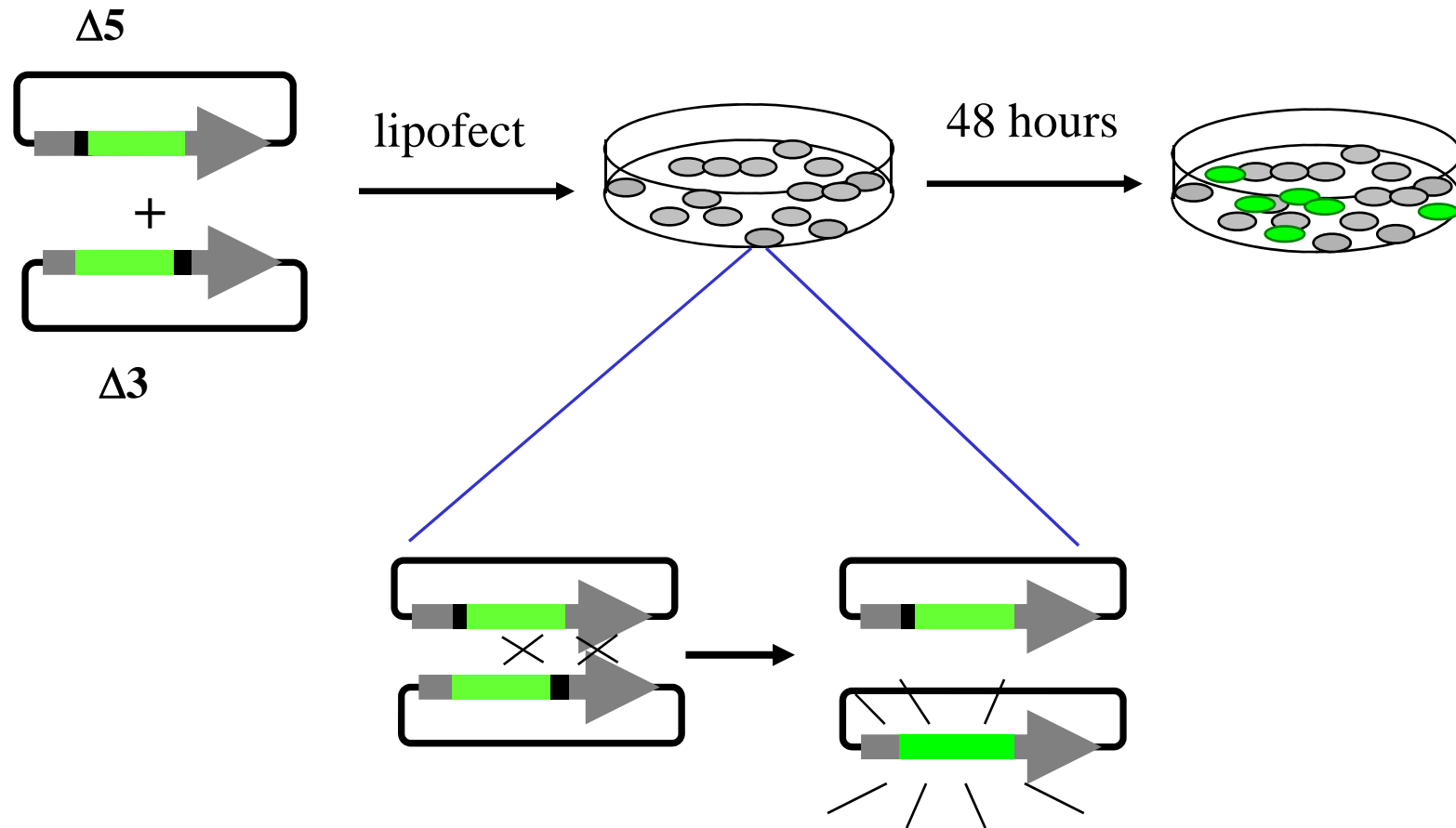
  - tertiary structure

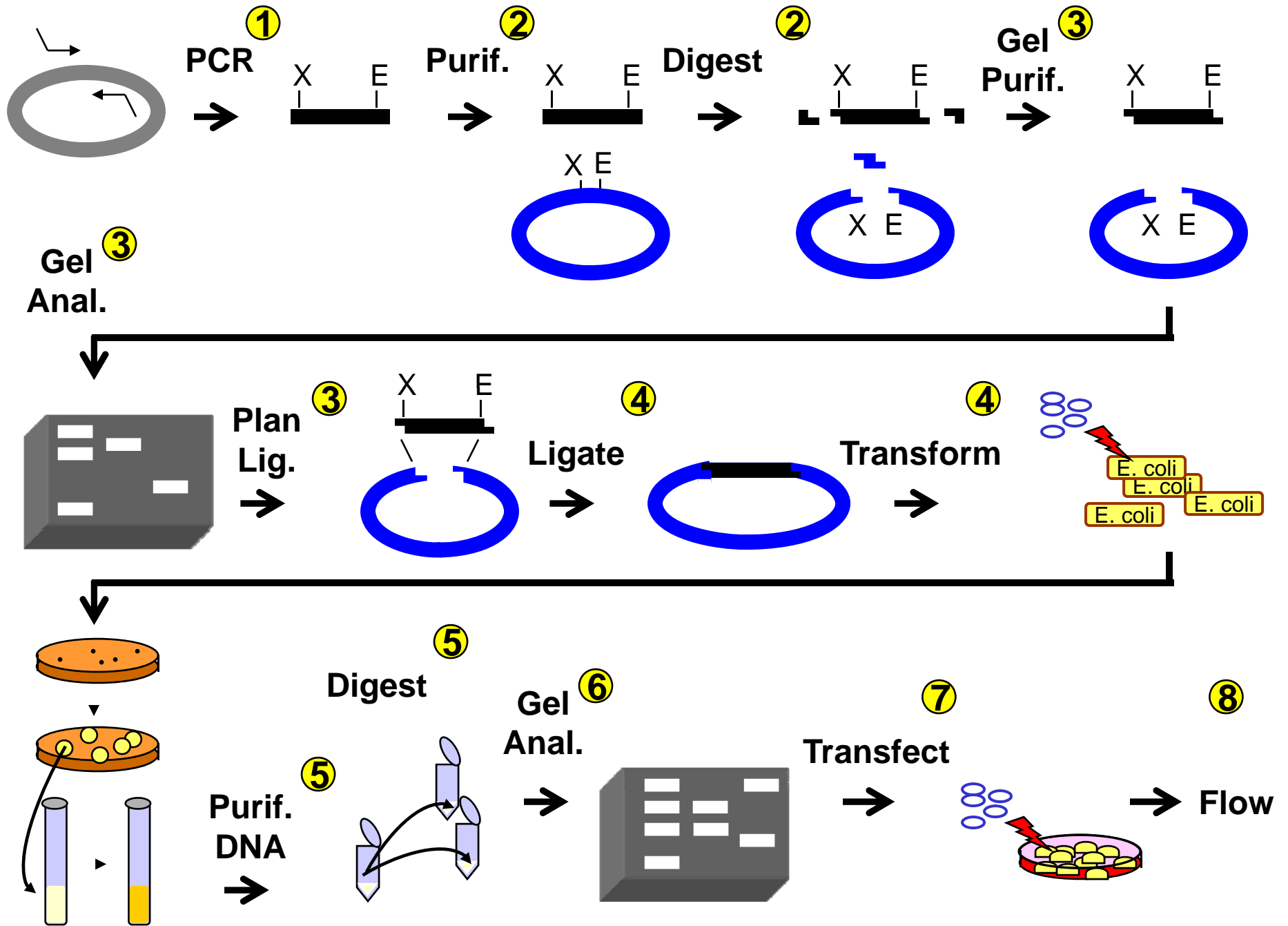
  - effects of overloading

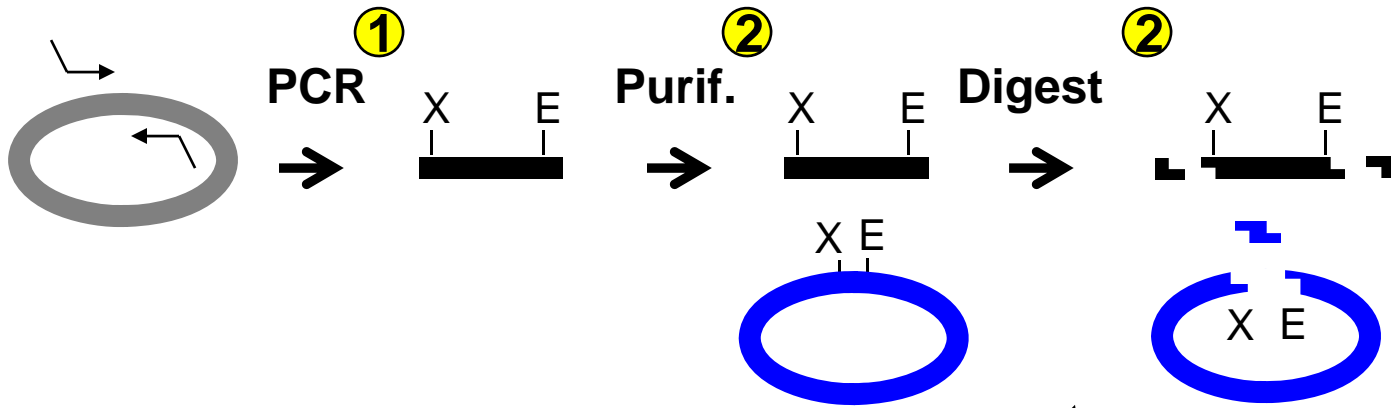
# Overview of the Experiments in Mod1

Where you are,  
and where you are going

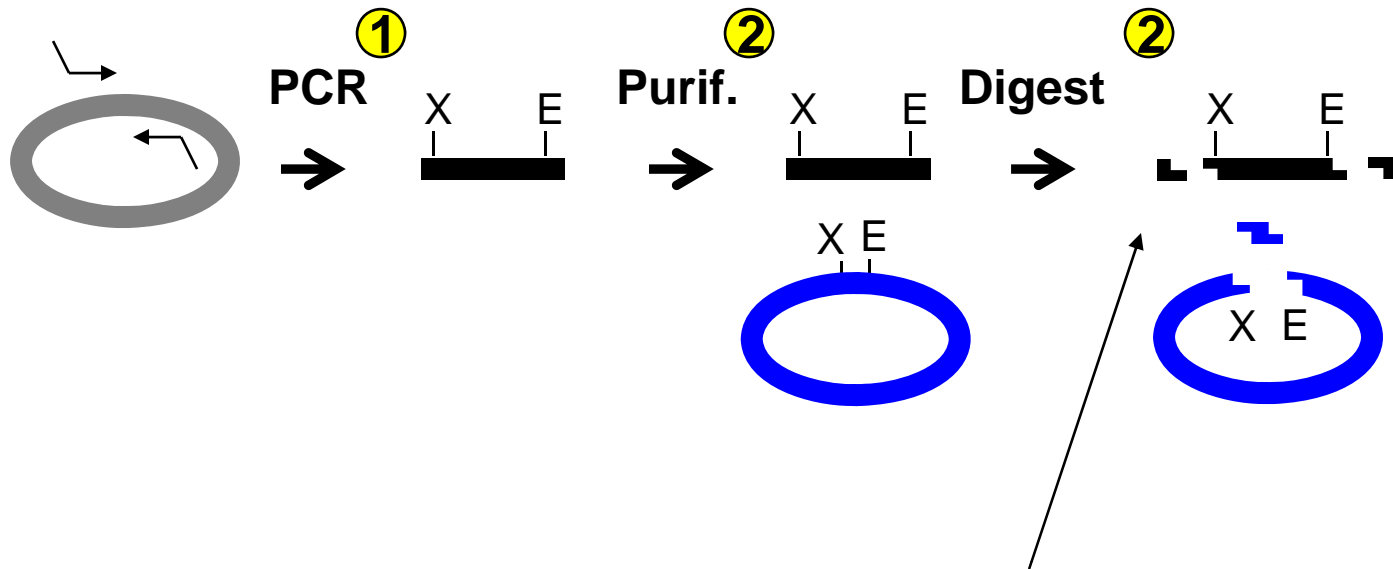
# A Plasmid-Based Assay for Homologous Recombination in Mammalian Cells





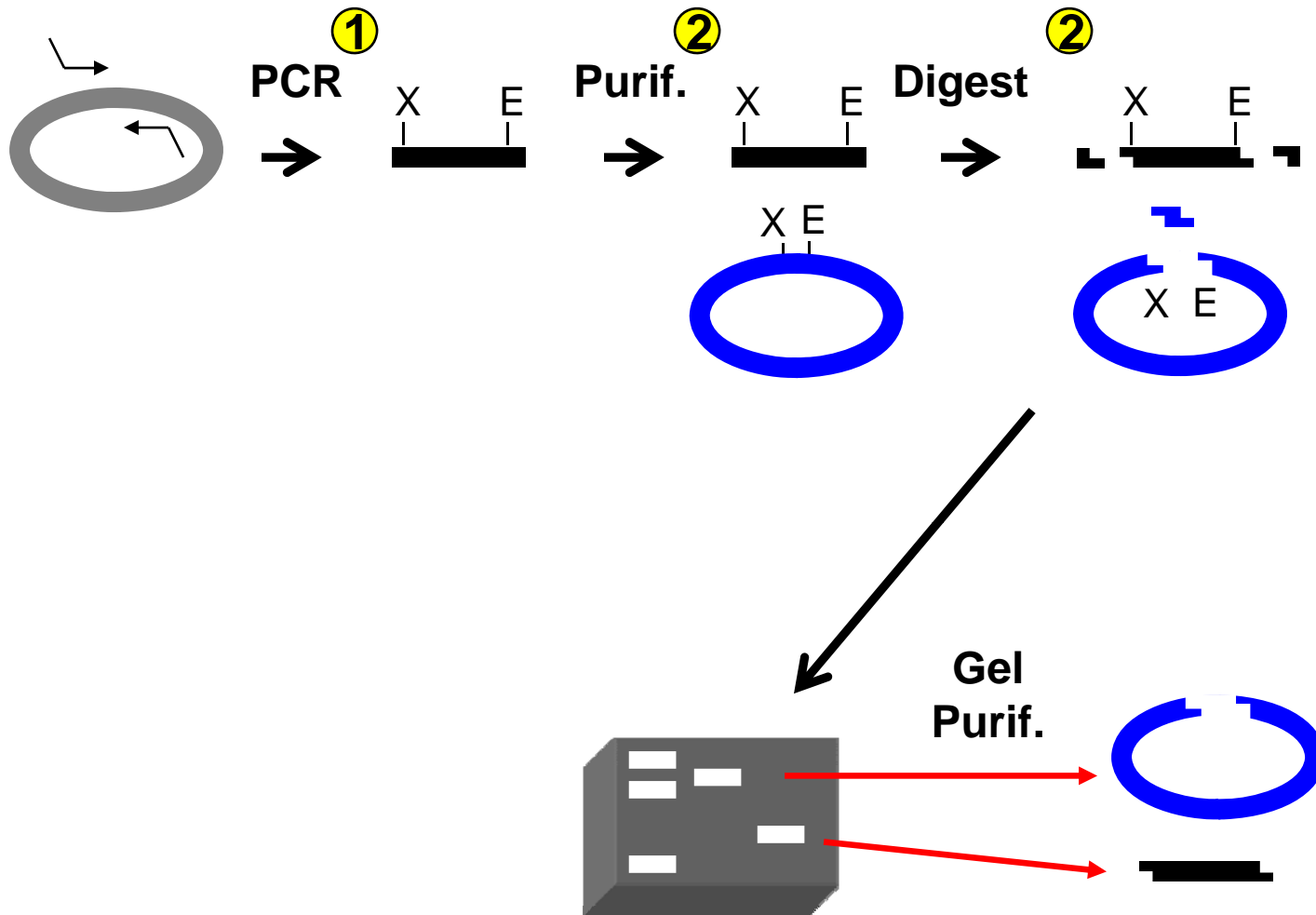


How do you know that your restriction enzymes actually cut the DNA?



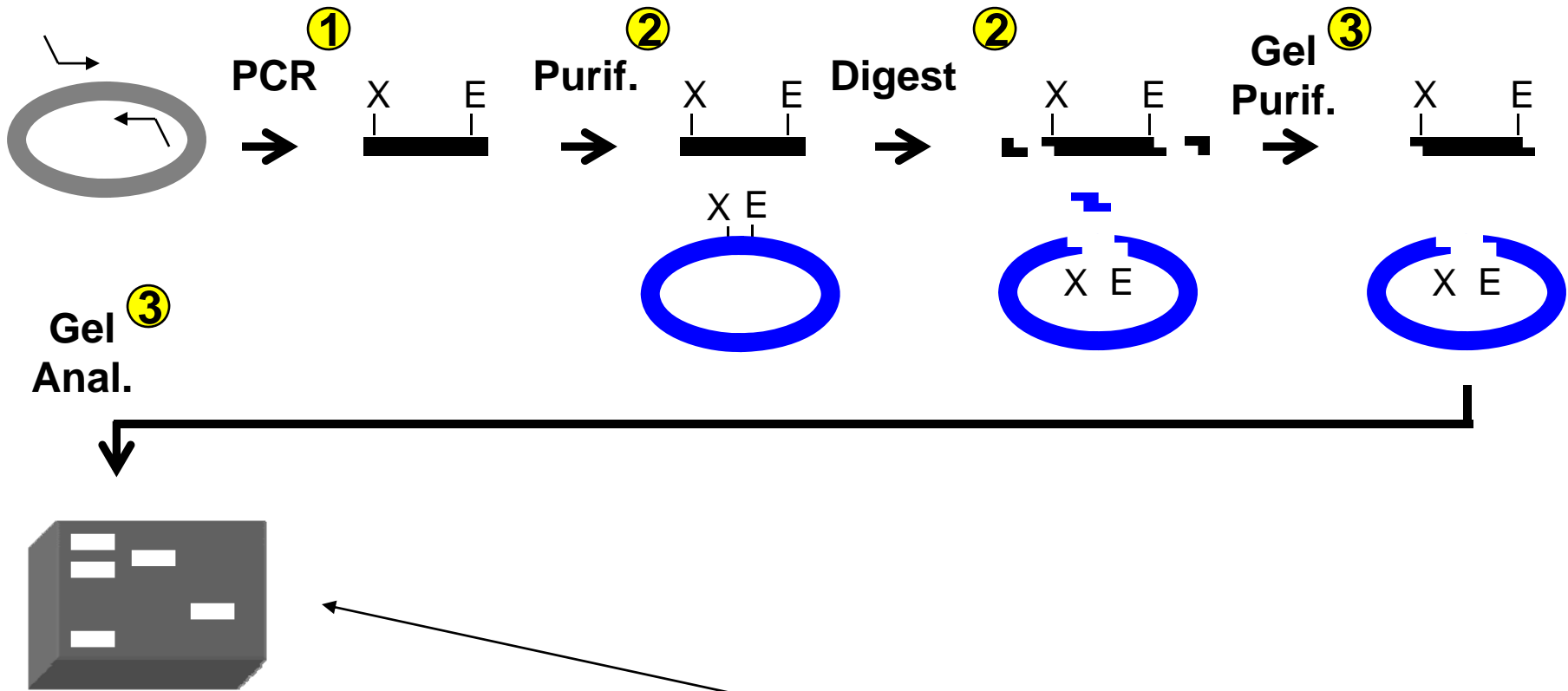
What else is in the reaction with the digested PCR product?

What effect could it have?

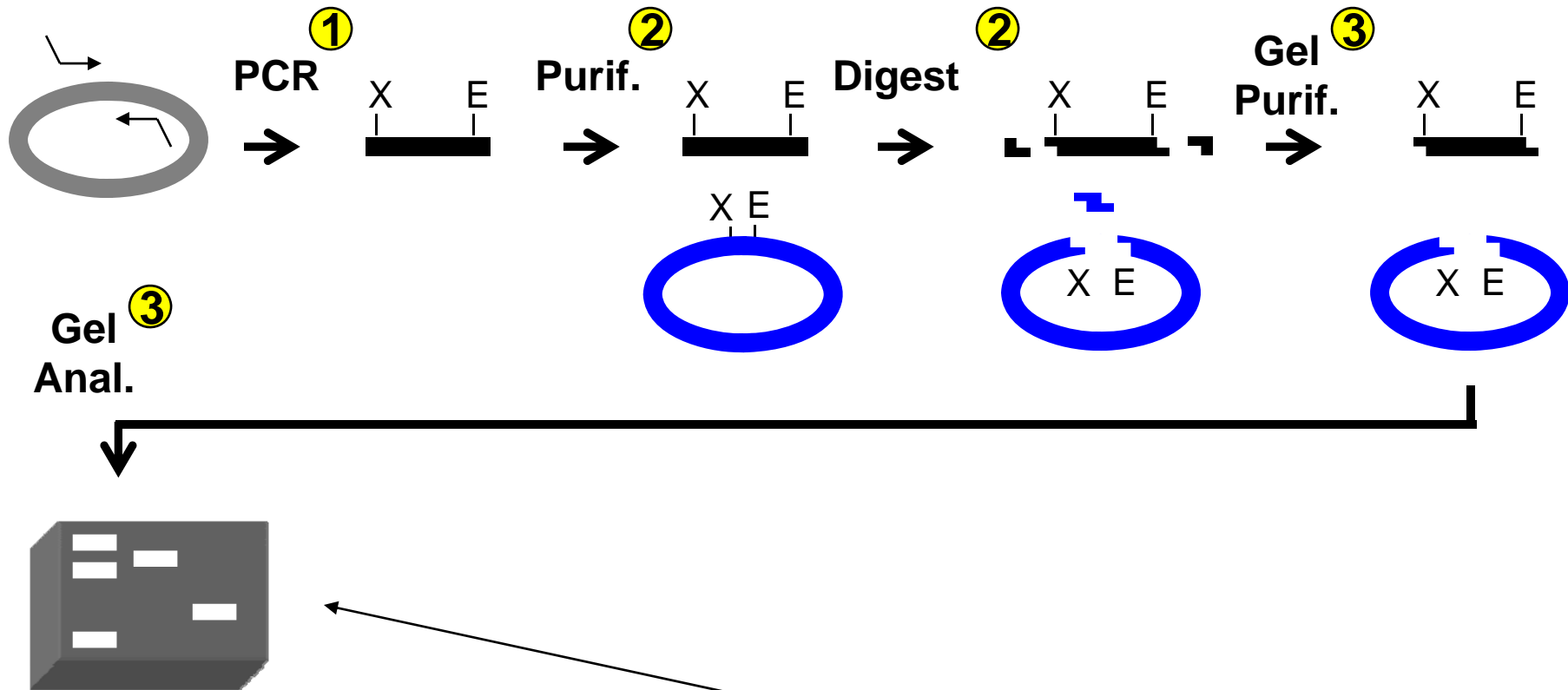


Why is it important to excise the DNA from the gel relatively quickly?



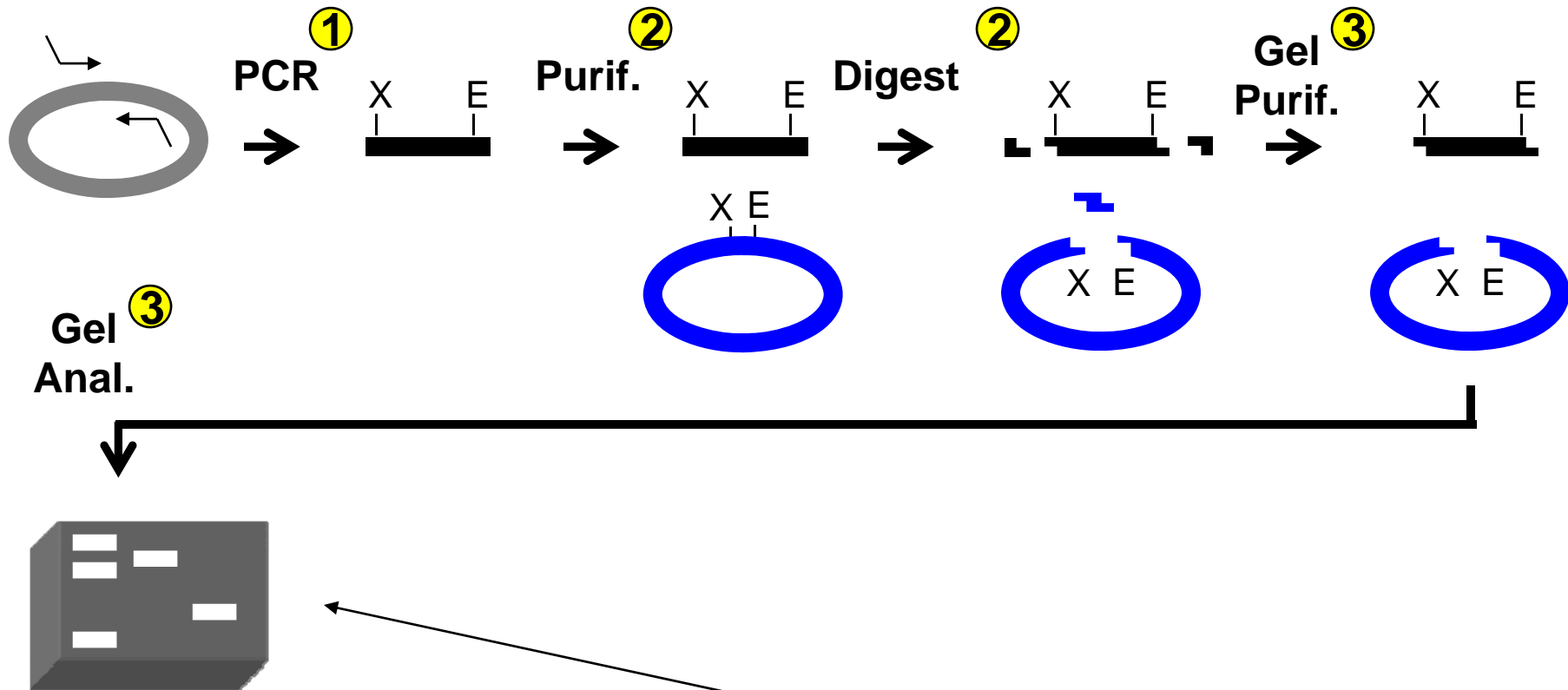


Why run this gel?



Your objective is a 1:4 vector:insert ratio – Why?

What if it was 1:100?  
What if it was 100:1?



How do you figure out how to get a 1:4 molar ratio?

## **About the experiments in Mod1**

- how is recombination used to fix double strand breaks
- how your two-plasmid assay works
- overview of the experiments you will be doing

## **Agarose Gels – How do we ‘look’ at DNA?**

- what is a gel and how do you load it?
- what happens to your DNA when it is exposed to UV?
- parameters that affect migration

## **Anticipating Potential Problems & Pitfalls**

- Getting the right DNA ratios