



# **Module 2: Manipulating Metabolism**

DNA engineering

10/13/15

# Wrapping up Mod 1

- Data summary draft submitted!!



- Comments to be returned on Oct. 18
- Revision due by 5 pm on Oct. 24
- Mini presentation due by 10 pm on Oct. 15
- Blog post due by 5 pm on Oct. 25

# Mod 2 assignment overview

- Journal club presentation
  - Communicate the key findings from a peer-reviewed published article
  - Completed individually
- Research article
  - Describe your results and analysis in a written 'formal' document
  - Completed individually

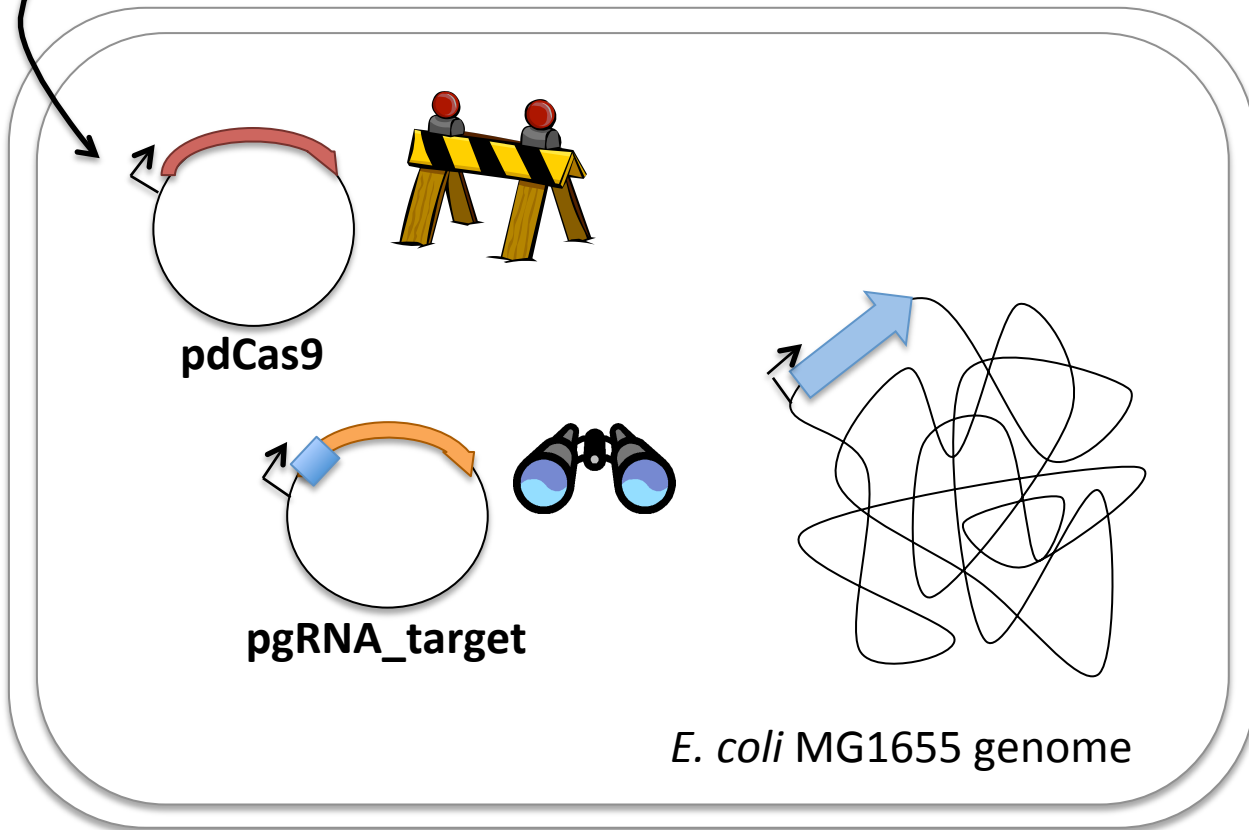
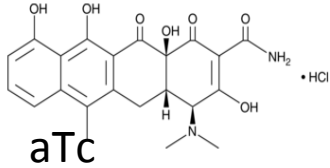
Your engineering task in Mod 2:

Increase production of ethanol or lactate  
in *E. coli* MG1655 by manipulating the  
fermentation pathway

**WHENEVER I'M NEAR YOU  
I UNDERGO ANAEROBIC  
RESPIRATION**

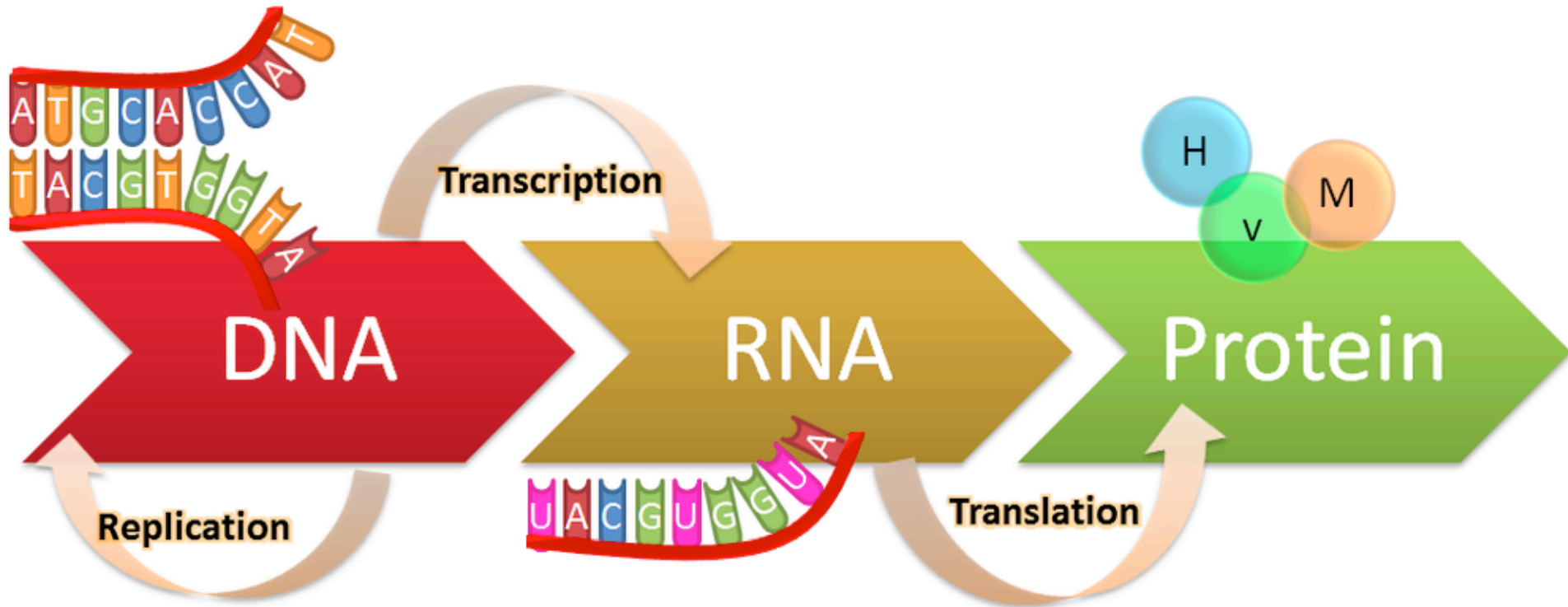
**BECAUSE YOU TAKE  
MY BREATH AWAY**

# CRISPRi system overview

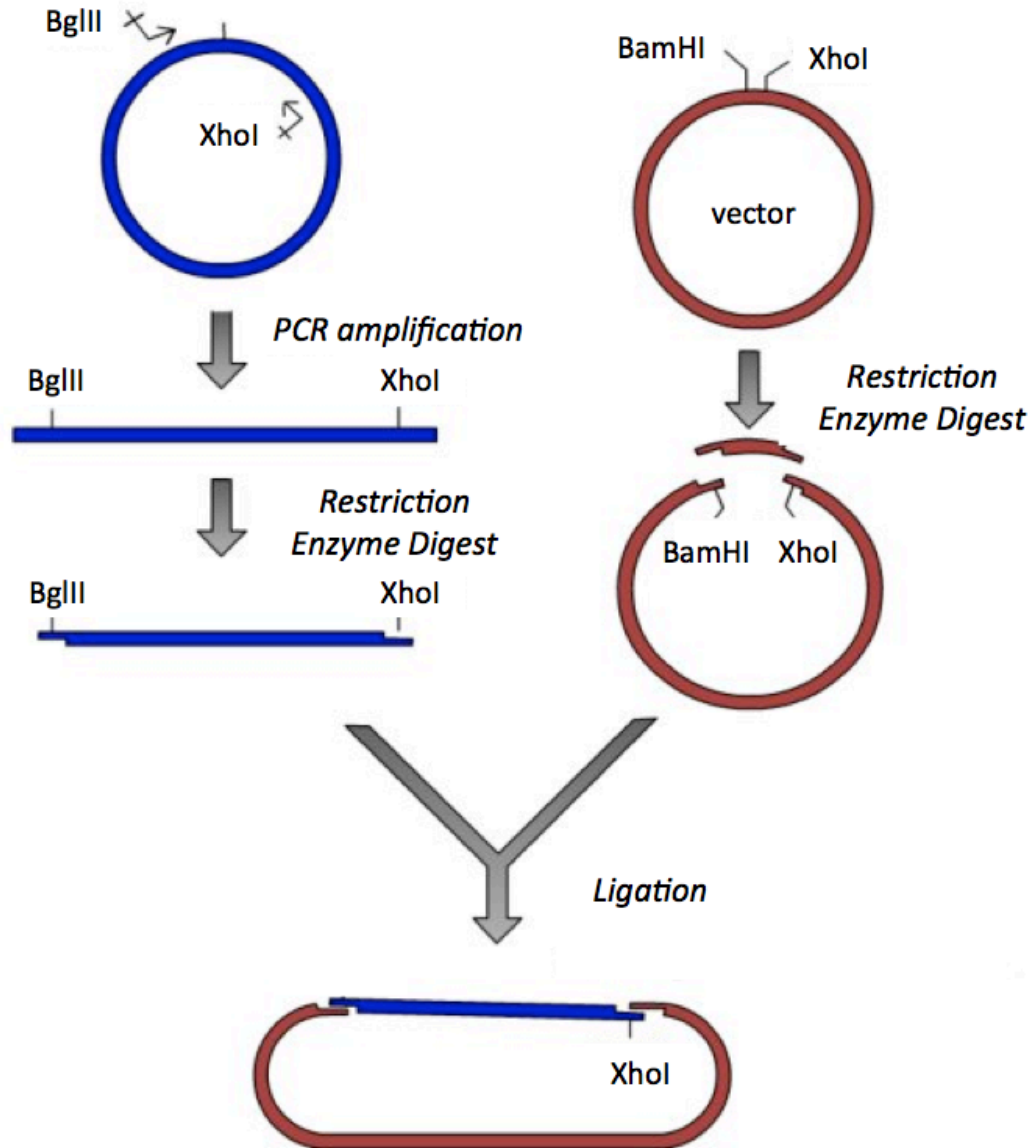


- Target gene
- pgRNA\_target
- pdCas9

# The central dogma



# How do we engineer DNA?



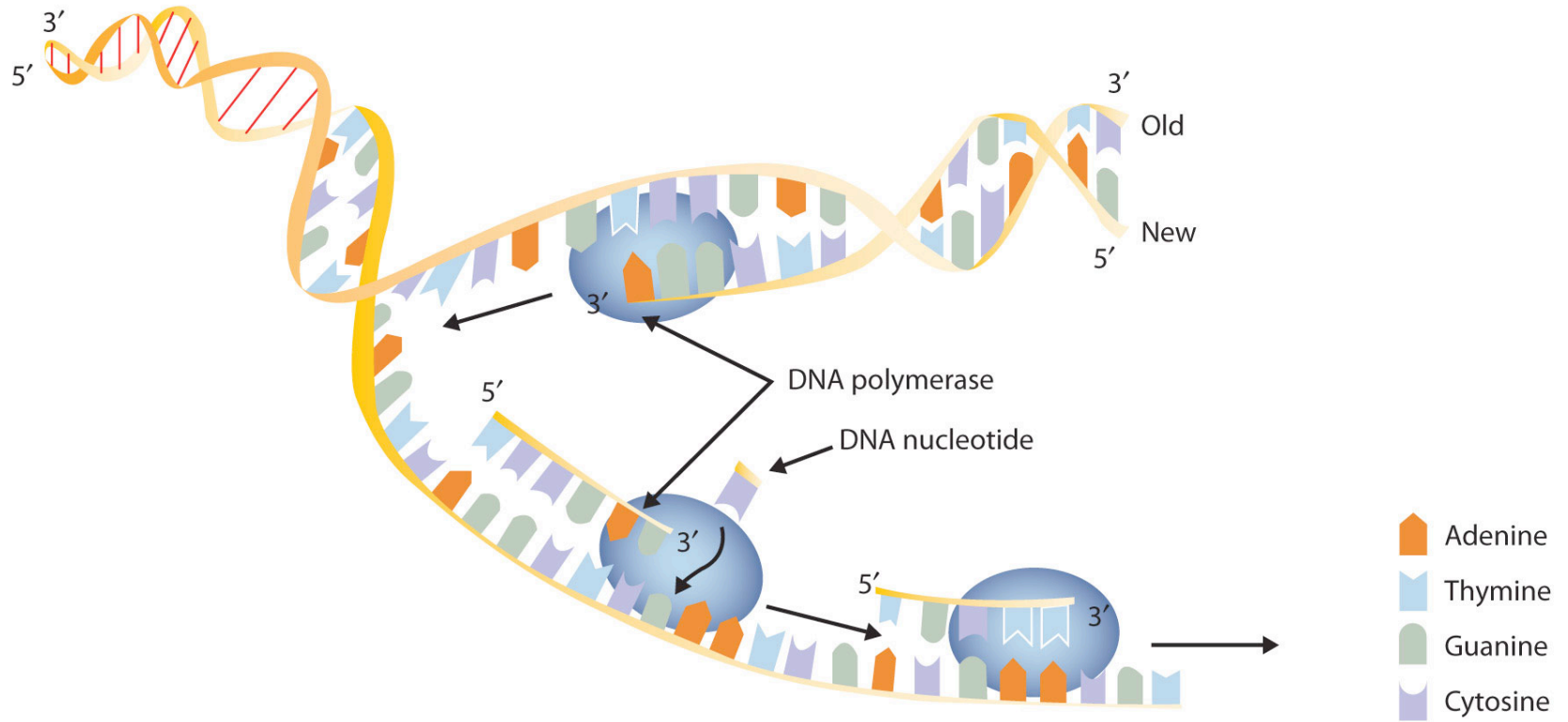
1. Amplification

2. Digestion

3. Ligation

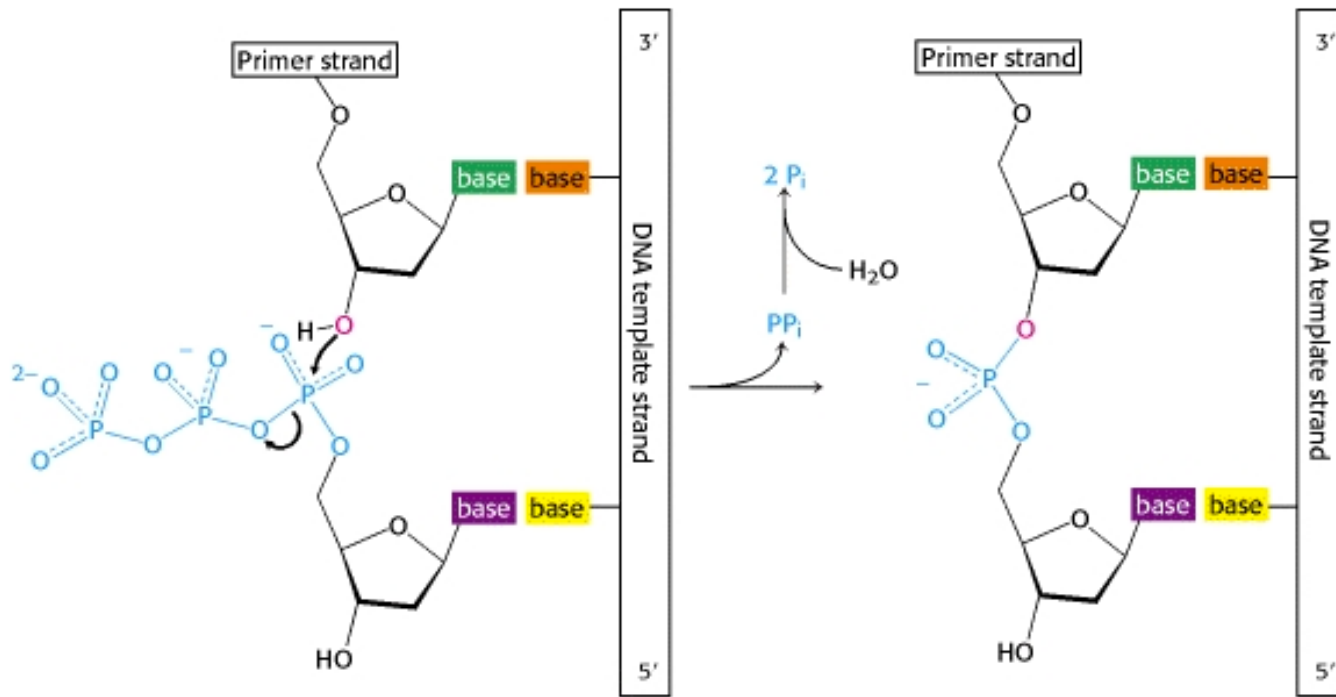


# Amplification



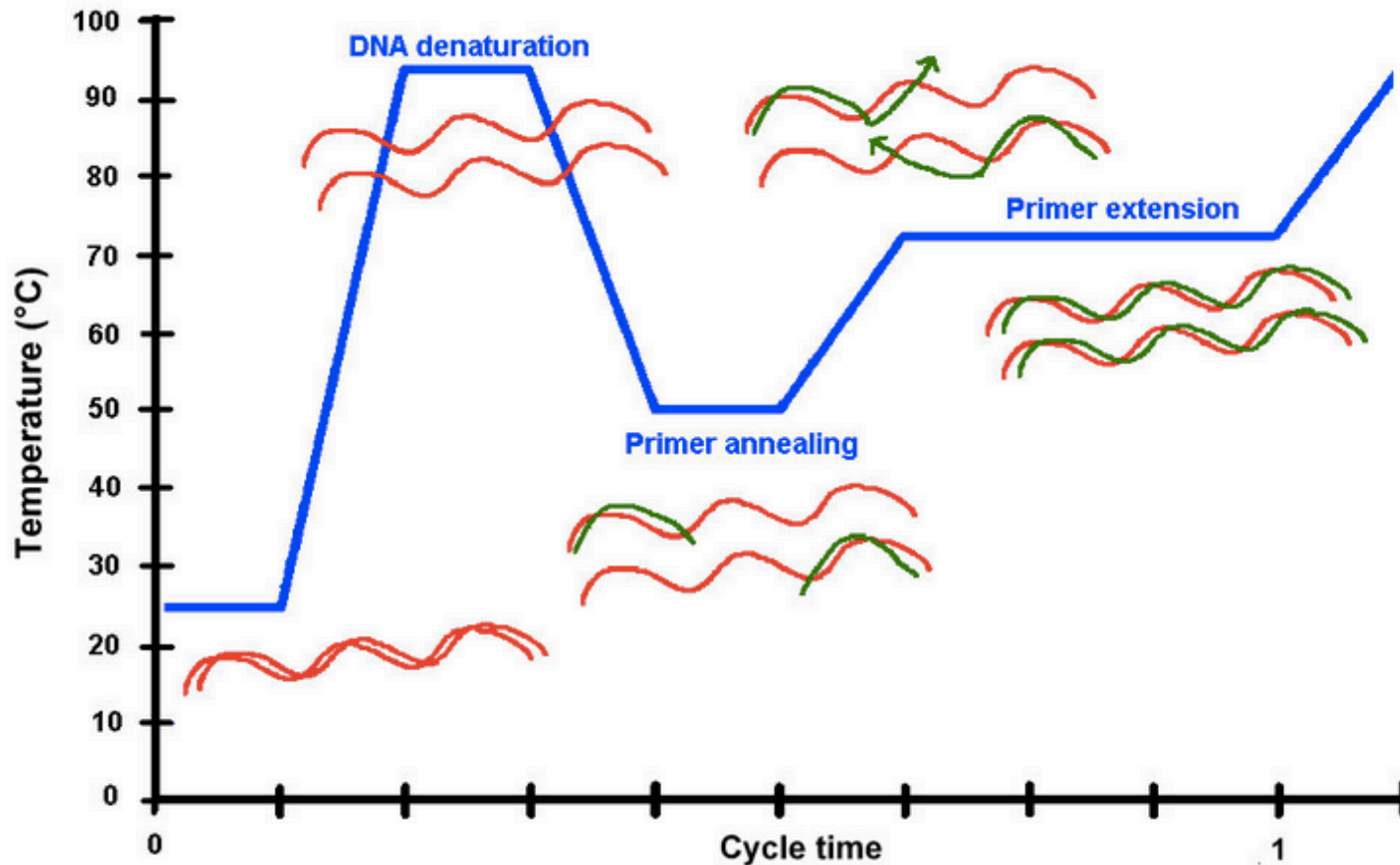
Who are the key players?

# DNA polymerase



- Catalyzes formation of polynucleotide chains
- Requires a primer base-paired to template

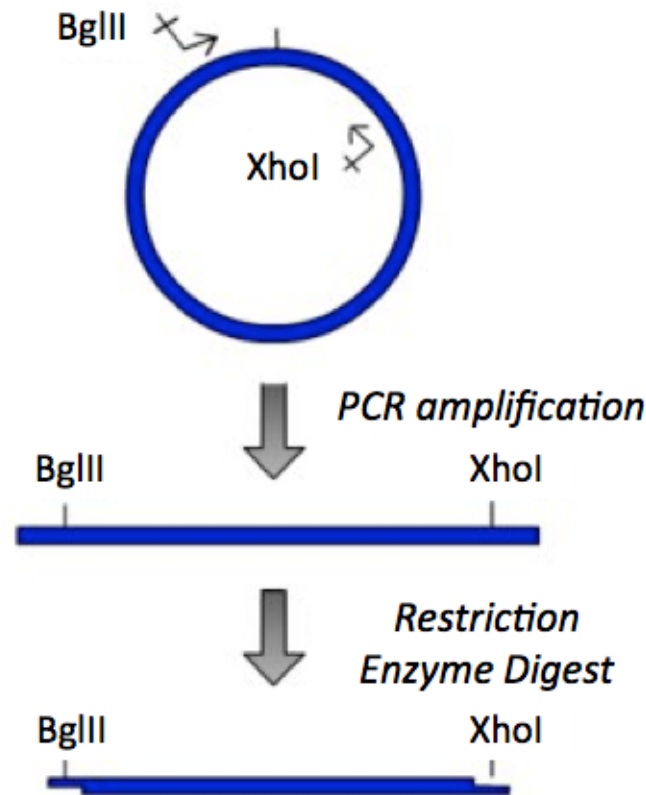
# Polymerase chain reaction (PCR)



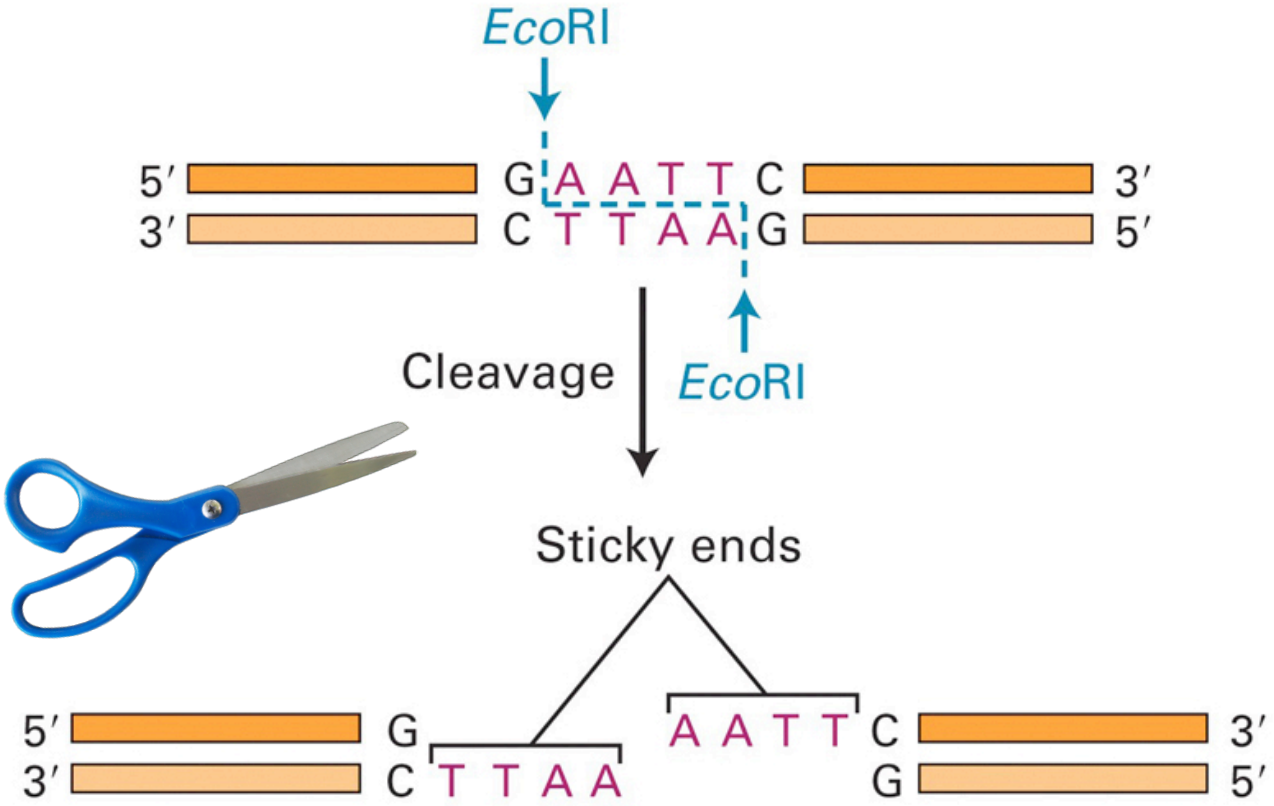
How many cycles until your product is generated?

# What are we amplifying?

- Primers enable you to specify which region of DNA is amplified by polymerase

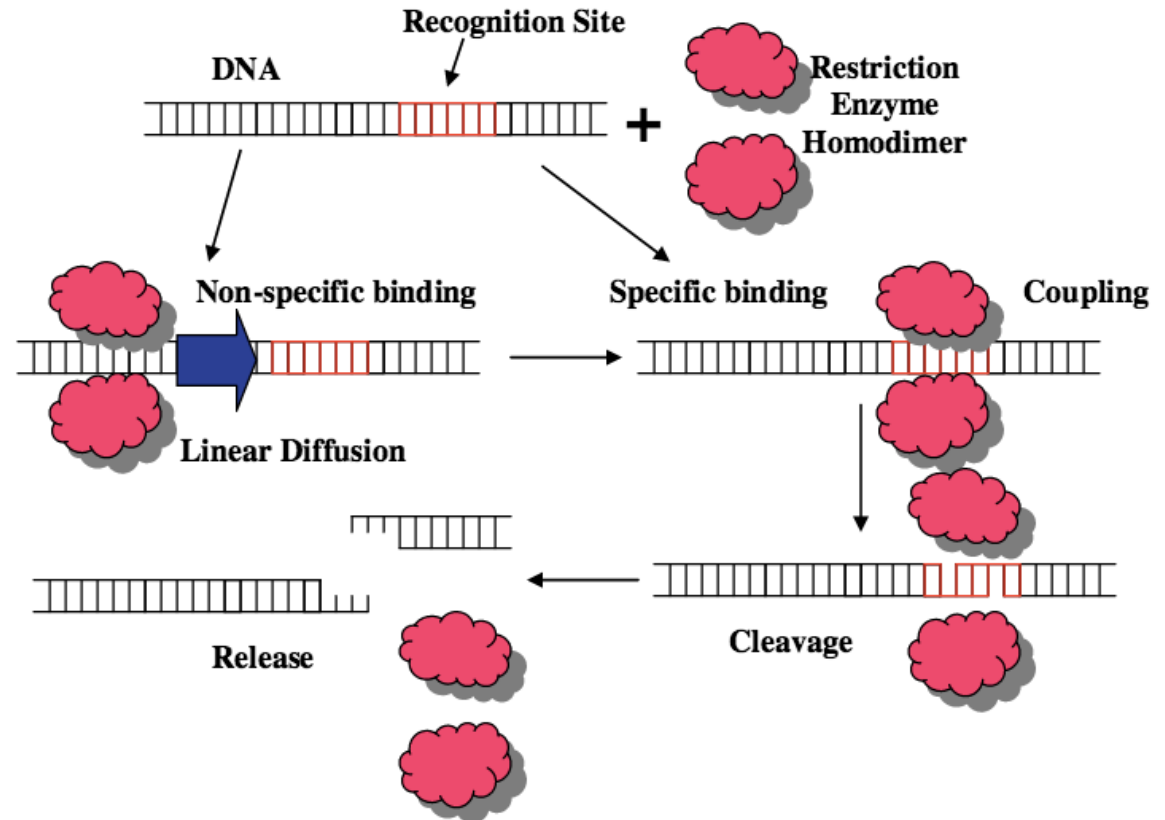


# Digestion

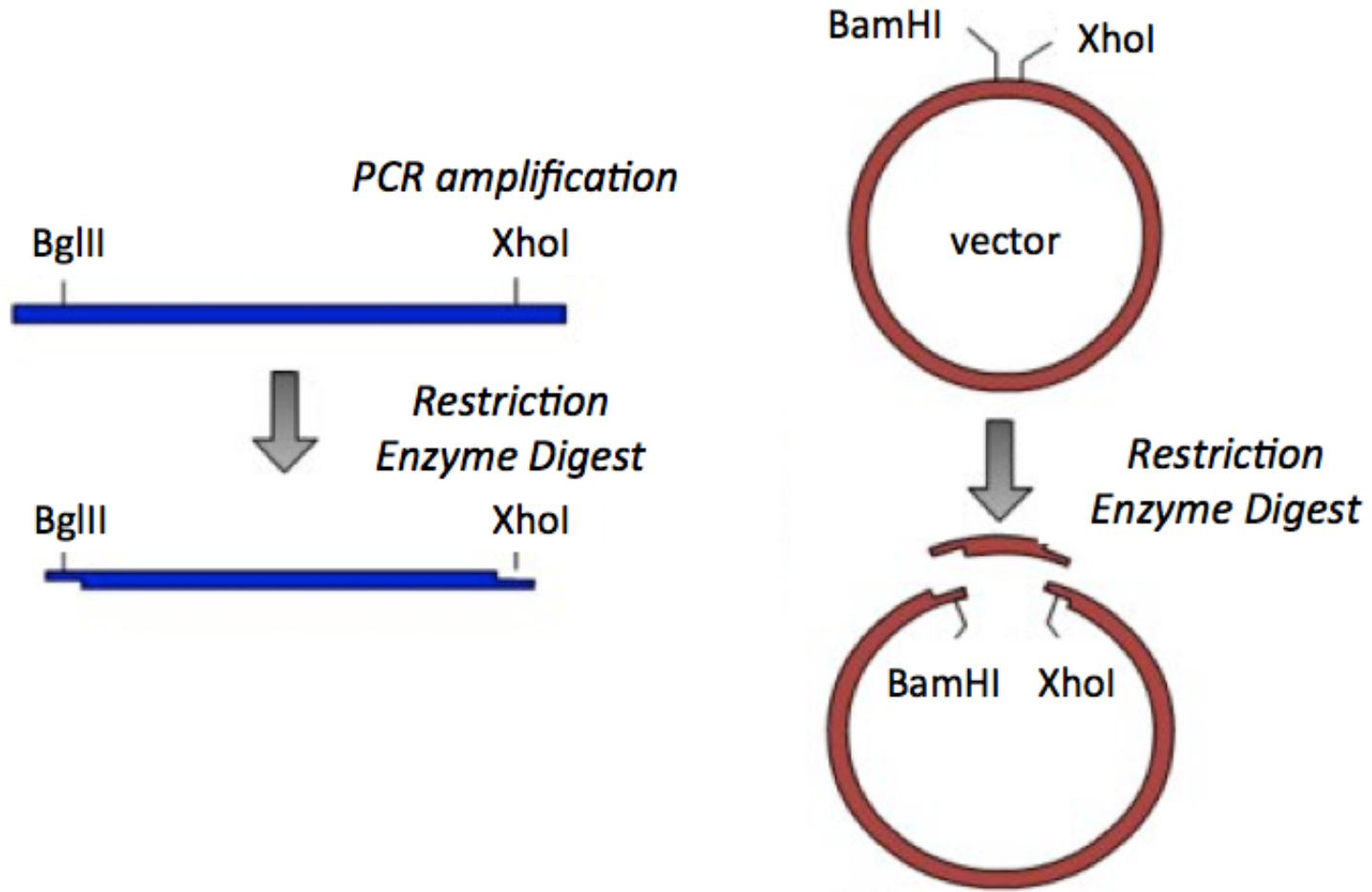


# Restriction enzymes

- Function as homodimers
  - Each dimer cleaves backbone at site of palindromic recognition sequence

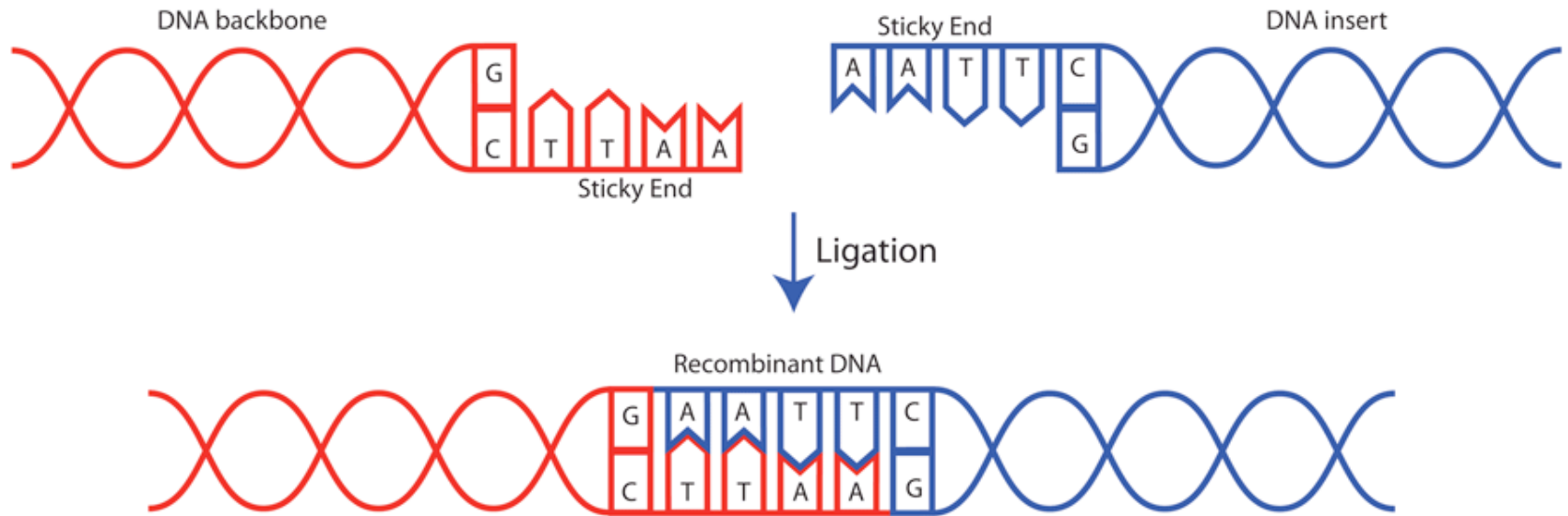


# What are we digesting?



What should we consider when performing a double digest?

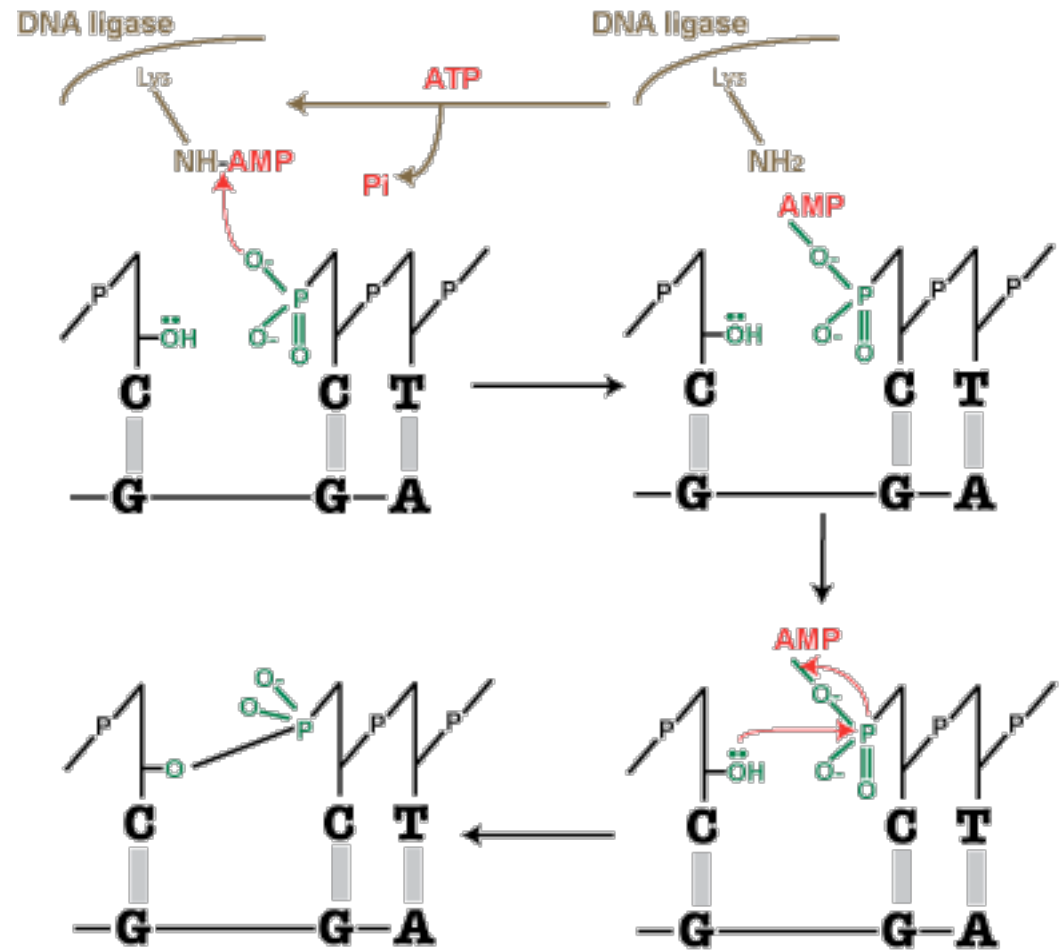
# Ligation



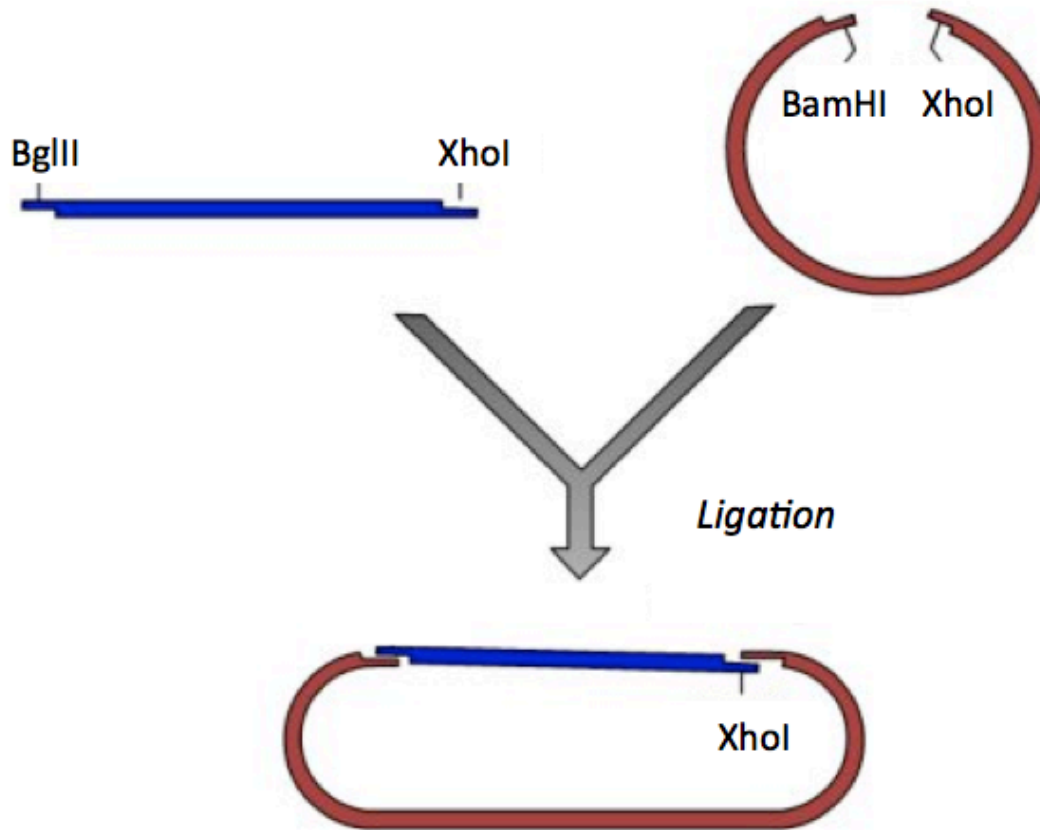


# DNA ligase

- Forms covalent phosphodiester bond between 3' OH acceptor and 5' phosphate donor
- Requires ATP

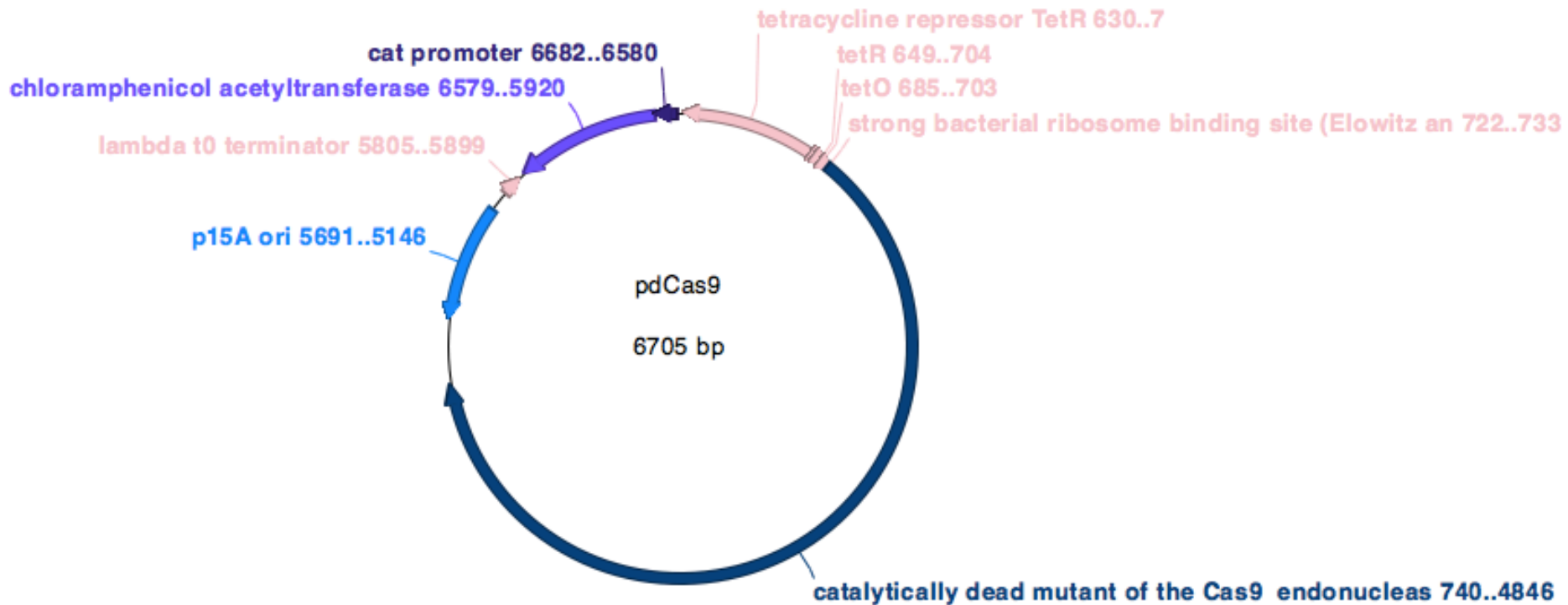


# What are we ligating?



Note: in your laboratory exercise only the 'top' DNA strand is represented...remember this when determining basepair sites of digestion and ligation.

# How do we confirm our product?

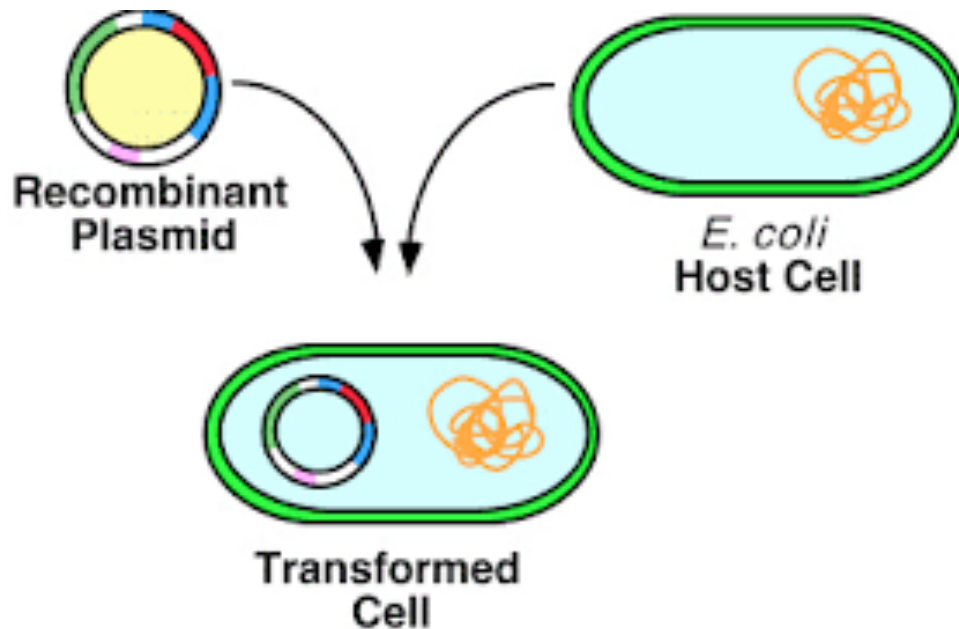


1. Transformation

2. Purification

3. Digestion

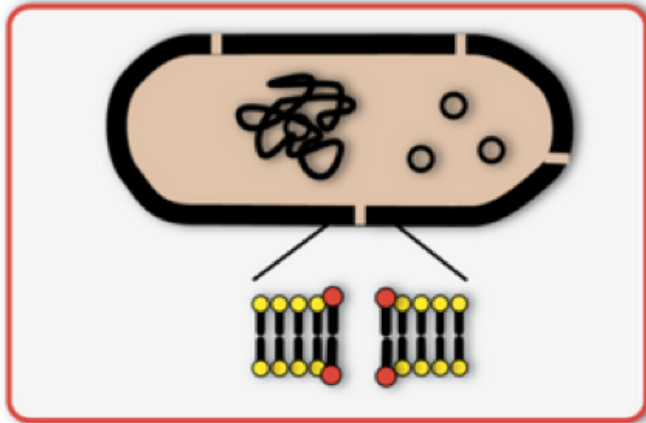
# Transformation



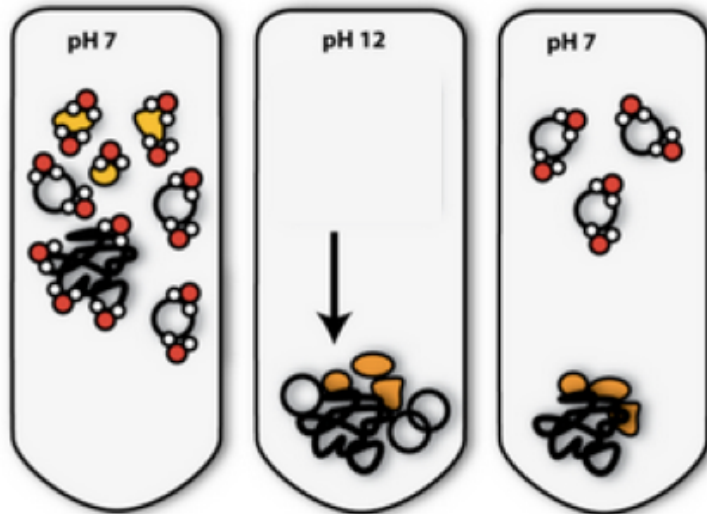
1. Incubation
2. Heat shock
  - DNA taken in by competent cells
3. Recovery
4. Selection

Why do we transform the ligation product?

# Purification



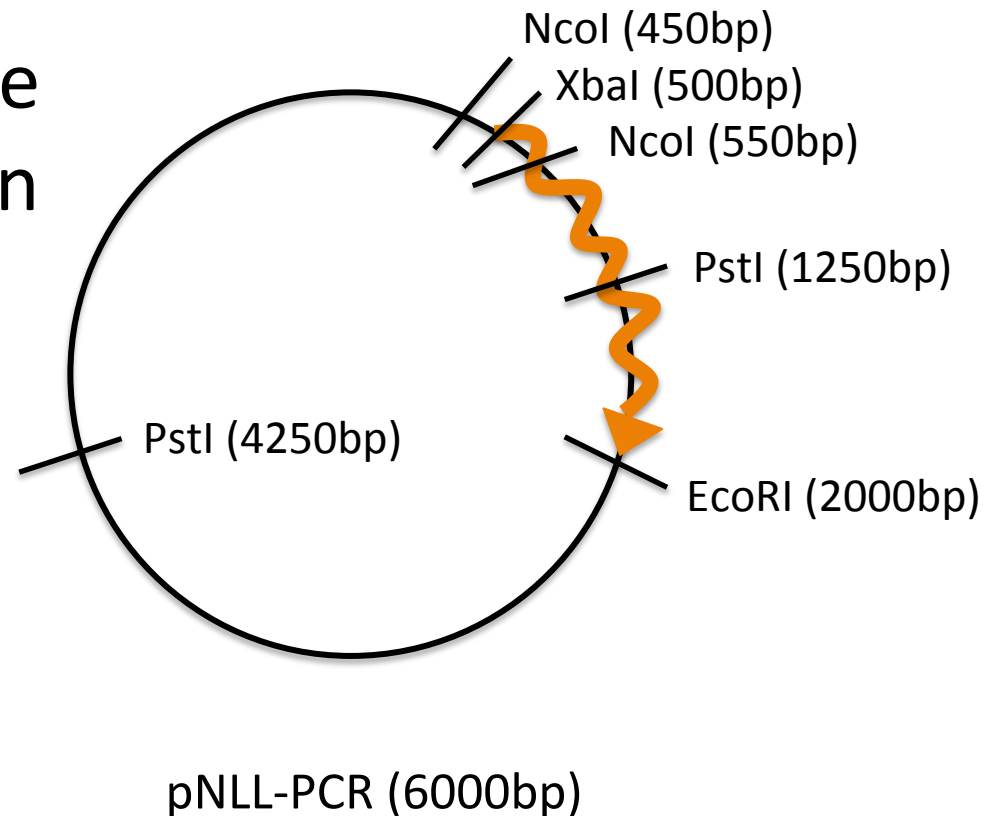
1. Resuspend cells
2. Lysis
3. Neutralization
  - Separates chromosomal DNA from plasmid DNA
4. Wash
5. Resuspend or elute DNA



Why do we purify (mini-prep) the ligation product?

# Digestion, again

- Confirmation digests
- Ideally, will cut once in insert and once in vector
  - XbaI and EcoRI?
  - PstI?
  - NcoI?



# In the laboratory...

1. Engineer pdCas9 construct
  - In silico 'cloning' of expression vector and dCas9 gene insert
2. Confirm pdCas9 construct
  - Digestion of pdCas9 to confirm presence of dCas9 gene insert