


# MID2: Clean and Cut DNA

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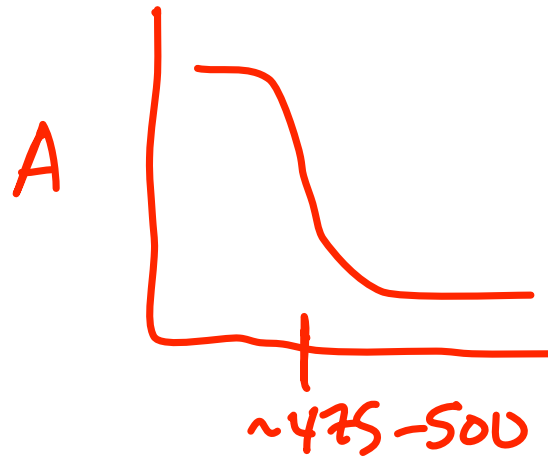
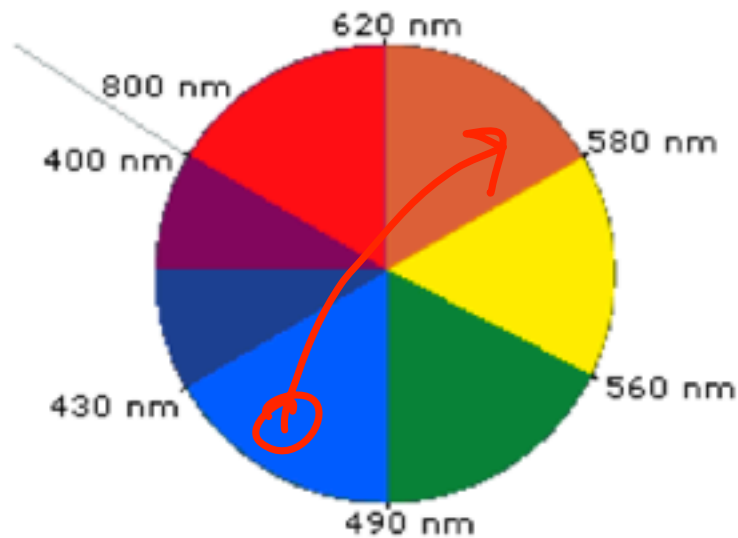
9/12/13

1. Pre-lab discussion
2. Clean-up PCR reaction
3. Digest vector & PCR rxn  APÉ
4. 4pm -- visit from Leslie

Note: Practice notebook option -- see wiki!

(PLEASE GRADE!) <sup>5pm</sup>  
Friday

# Lab practical review:



ROYGBIV

\* gloves  $\Rightarrow$  benchtop/burn box

0.03% XC  $\rightarrow$  1% stock

$$C_1 V_1 = C_2 V_2$$

$$\frac{(10 \text{ mL})(0.0003)}{0.01} = 0.3 \text{ mL}$$




# Module I major assessments:

5% ① Methods → FNT draft  
MIDI-PCR  
MIDI-RE dig

15% ② Data Summary — WAC here

- figures + captions
- summary
- abstract

# Methods Section Tips

- Divide into sub-sections! 
  - Put in a logical order -- list primer THEN list PCR reagents, etc
  - Start with a overview sentence “EGFP was amplified using polymerase chain reaction (PCR).”
- Methods are clear and concise explanations
  -  **The methods section is not a benchtop protocol** 
  - Space-wise, avoid tables/lists when a sentence will do
  - Sentence-wise, avoid extra words
  - Content-wise, cover what’s needed and only that needed to understand and replicate your work
- Think about the most flexible units -- concentration vs volume. (For replication!)

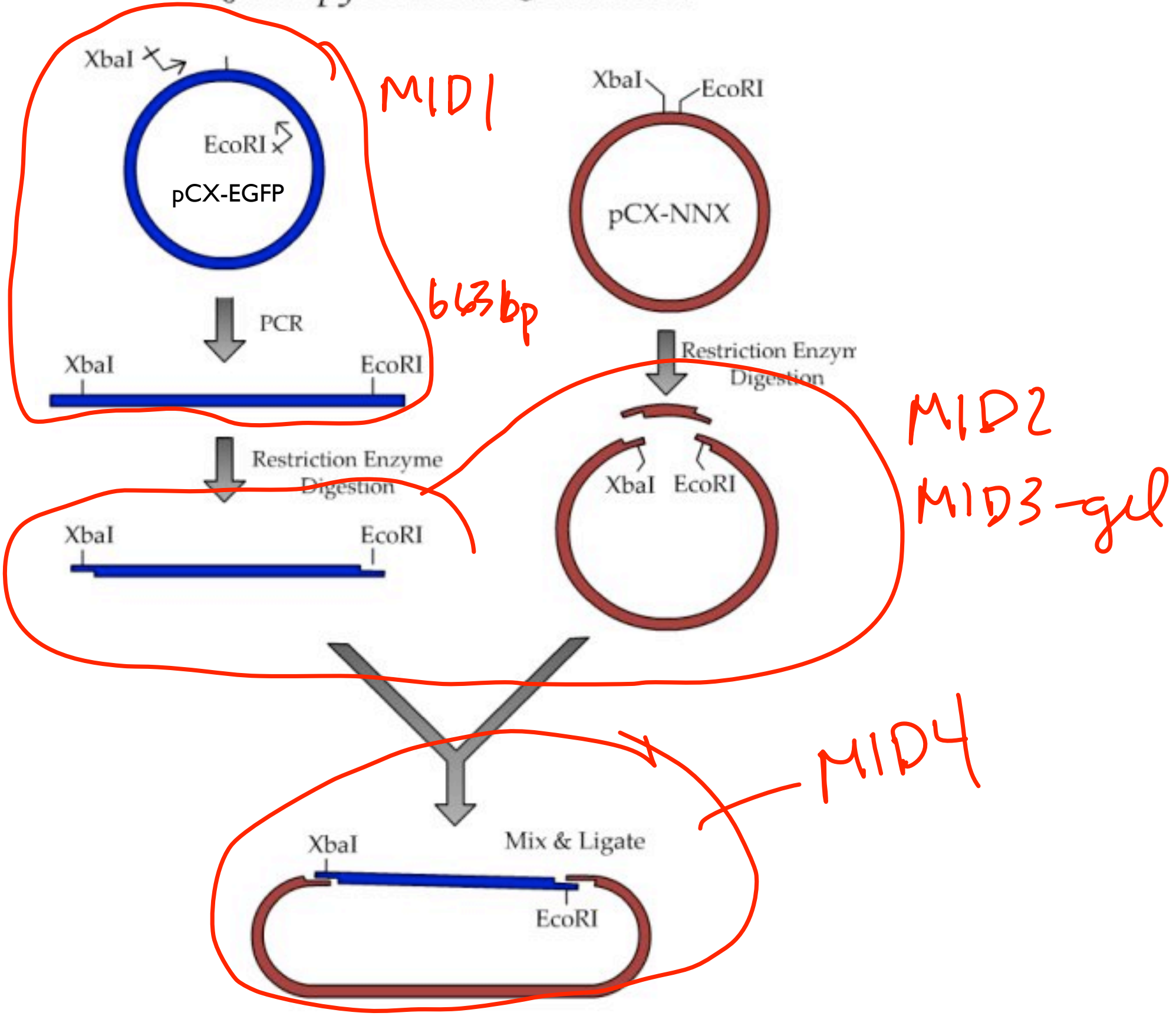
# Methods section exercise

- Consider the following passage: *pcx-EGFP plasmid* *concentration* *primers / seq* *what is it?* “Template DNA (5 ng) and primers were mixed with 20  $\mu$ L of 2.5X Master Mix in a PCR tube. Water was added to 50  $\mu$ L. A tube without template was prepared and labeled control.”  
*[final]*
- What information is missing?
- What information can be cut?

Thanks to Agi for this slide!

# Step I: Build the system!

Roadmap for Plasmid Construction



Last time:



XbaI + EcoRI - Sub cloning tools

pCX-EGFP

XbaI  
EcoRI

NNX

BamHI + EcoRV - troubleshooting/  
confirming our success

Oh my, so G/C rich -- what can you do to improve your chances?

~ 70% G/C  
T<sub>m</sub> = 69.3°C

DMSO - untold  
glycerol - NMP: ↓ T<sub>m</sub>  
T<sub>A</sub>

# First we'll clean up our PCR product:

① PB → ↑ salt ↓ pH  
chaotropic

bind

② Wash w/  
ethanol

(PE)

Silica  
membrane



③ Elute

↓ salt ~ pH

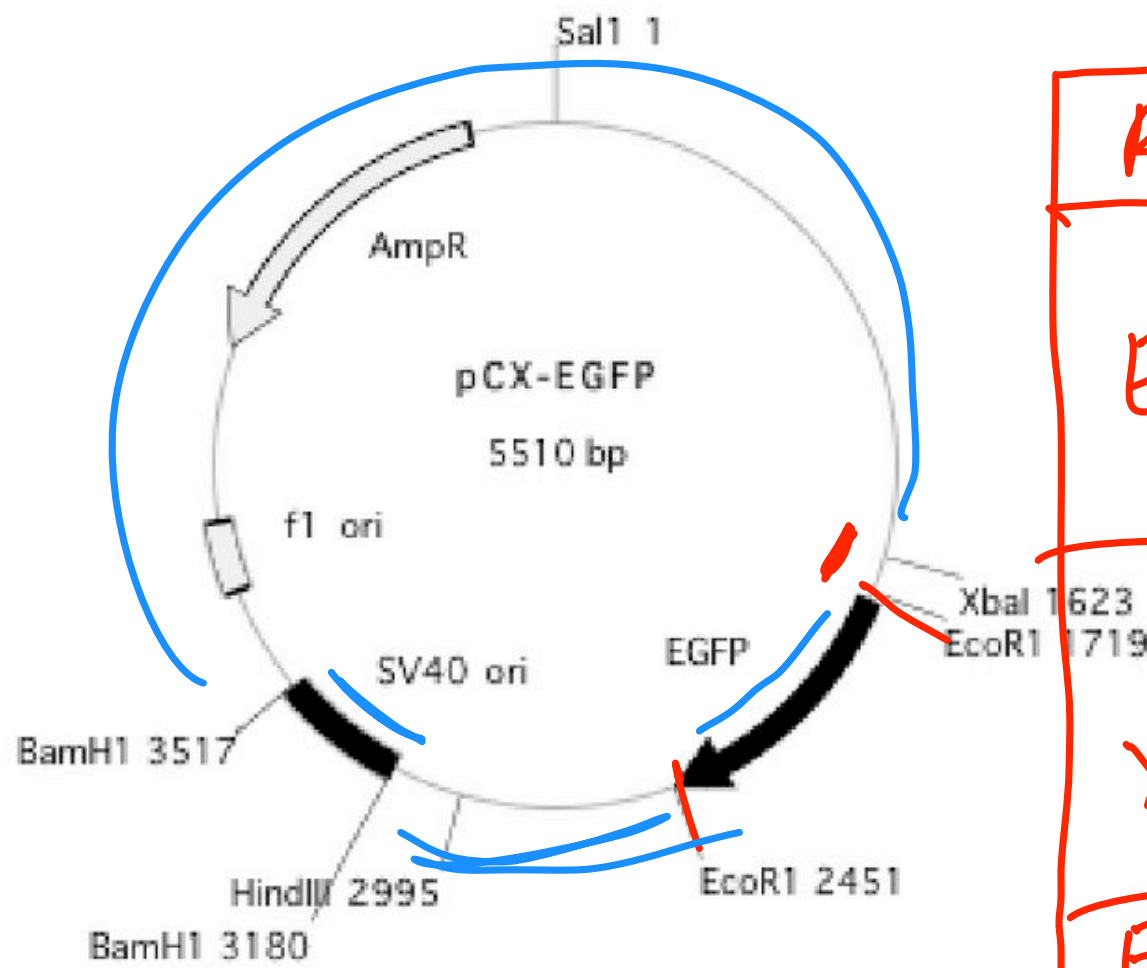
(H<sub>2</sub>O) → EB





# Today we will cut:

## Thought experiment:



RE	# of bands	Size
EcoRI	2	~ 700 ~ 4800
EcoRI + XbaI	3	~ 100 ~ 700 ~ 4700
EcoRI + BamHI	4	~ 700 ~ 300 ~ 700 ~ 3700

# Today we will cut:

XbaI

EcoRI

TCT AGA  
AGA TCT

G AATTC  
CTT AAG

↓ ≈ 37°C

C T A G A  
T

G  
C T T A A

Total volume of RE digest = 25  $\mu$ L ★ Label your tubes ★

HF

Component	Details	Order
Enzymes	EcoRI ✓ XbaI ✓ ]	④
DNA	PCR (1) - single digest PCR-NNX (3) - double digest	③
2.5 $\mu$ L Buffer	Cutsmart (10x) $\Rightarrow$ 1x	② } mix
Water	to 25 $\mu$ L	① } mix

Rxn temp: 37°C

Total Rxns: 4

## Today in the lab:

- PCR purify
- Set-up digests (4)
- ApE tutorial
- WAC visit at 4pm

## Next time in the lab:

- Agarose gel visualization
- Purification of DNA
- Longer FNT

