

M2D1: Complete *in silico* cloning of dCas9 & actual confirmation digest

10/12/18

1. Design primers to dCas9
2. *In silico* PCR amplification, digest, and ligation
3. *Actual* (wet-lab) diagnostic digest of pdCas9

(Almost) done with Mod1!

- Mini-presentation
 - due 10pm on Saturday, October 13
 - You don't have to be exhaustive;
tell a focused story
- Data summary
 - ✓ draft due 10pm on Wednesday, October 10
 - revision due 10pm on Saturday, October 20th
- Blog post
 - due 10pm on Thursday, October 11th



How is DNA engineered?

1. PCR amplification of DNA:

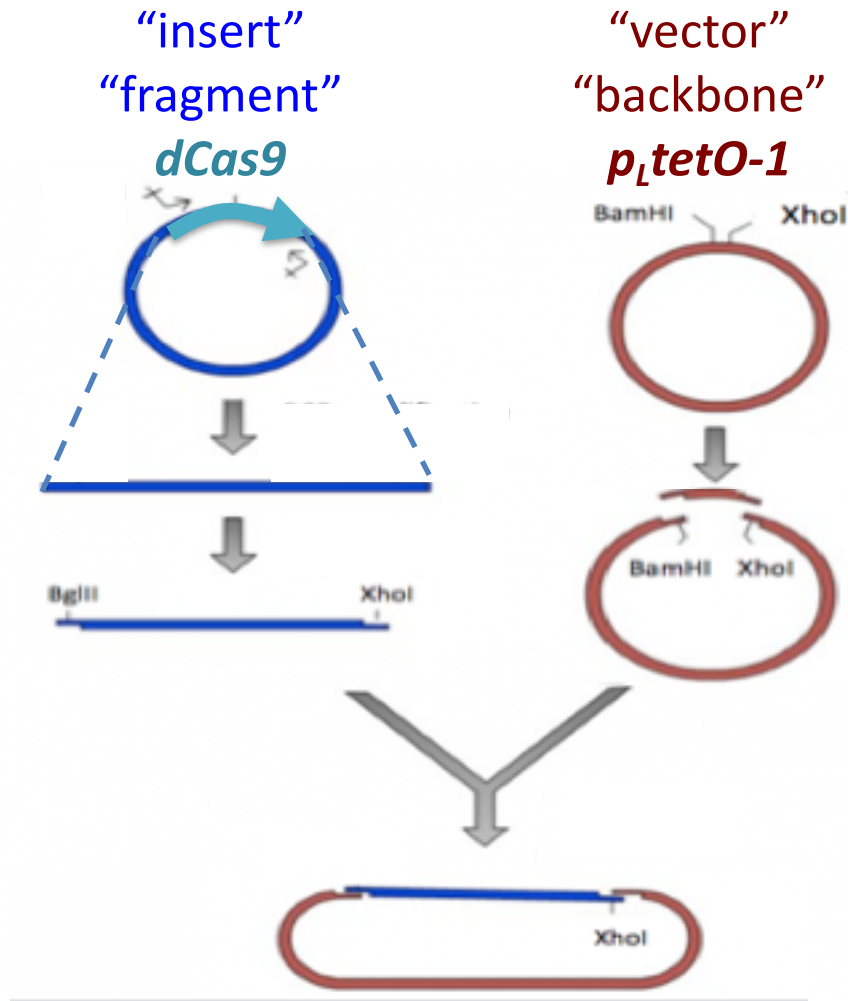
DNA polymerase
primers specific to gene seq.
nucleotides (dNTPs)

2. Digestion:

restriction enzymes
↳ endonucleases
• sticky end or blunt

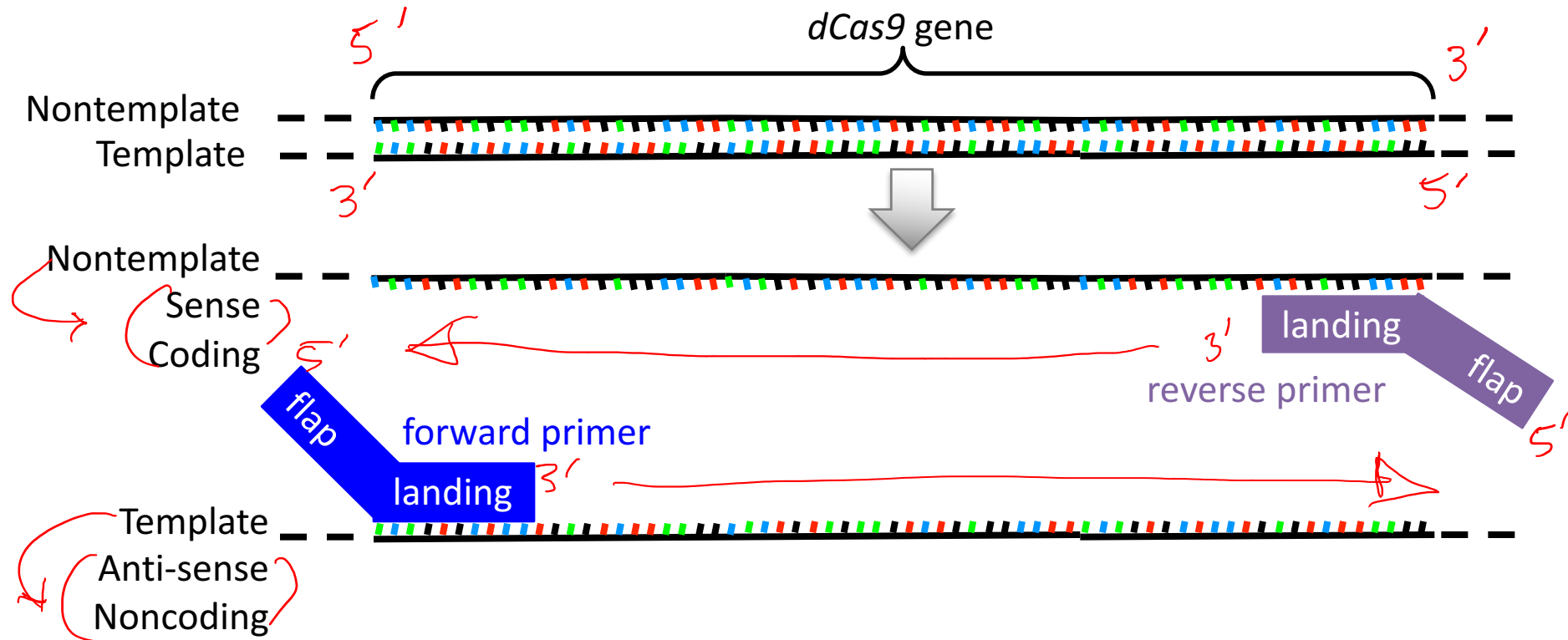
3. Ligation:

DNA ligase
↳ seals phosphodiester
bond in backbone



1. PCR amplification of DNA

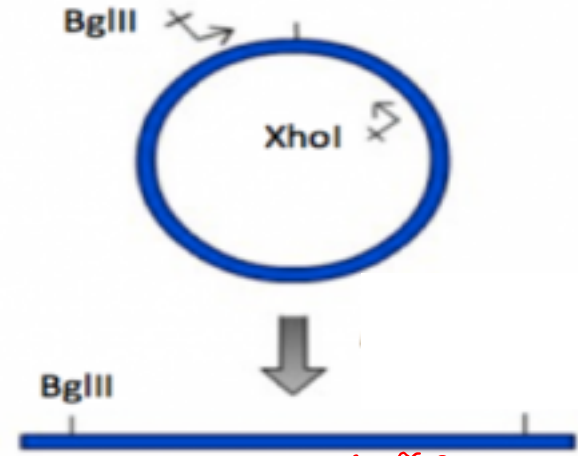
Defining terminology



1. PCR amplification of DNA

Designing primers

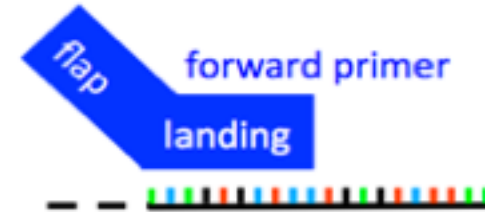
- **Landing sequence:** match to dCas9
- **Flap sequence:** contains endonuclease recognition sequence and junk DNA



1. PCR amplification of DNA

Primer design guidelines

- Length: 17-28 base pairs
 - long enough to be specific*
 - short enough for easy annealing*
- GC content: 40-60%
 - GC has 3 hydrogen bonds; AT has 2 hydrogen bonds
 - GC clamp at ends
- $T_m(\text{primer}) < 65^\circ\text{C}$ *ideal for specific priming*
- Avoid secondary structures
 - hairpins
 - complementation w/in primer sequence
- Avoid repetitive sequences
 - Max of 4 di-nucleotide repeats (ex. ATATAT)
 - Max of 4 bp in a run (ex. GATGGGG)



1. PCR amplification of DNA

Three major PCR steps—which temperature & why?

- Melt

- 95 °C

- hydrogen bonds break

- Anneal

- $T_m(\text{primer}) = 1/2$ primer annealed to target

- $T_{\text{anneal}} \sim T_m(\text{primer}) - 5^\circ\text{C}$ ↑ probability primer bound

- Extend

- 72 °C (for Taq)

- 1000 bases/min

Leslie's favorite PCR animation

<http://learn.genetics.utah.edu/content/labs/pcr/>

cycle # 2

The temperature is lowered so the primers will attach.

50° C

<< BACK

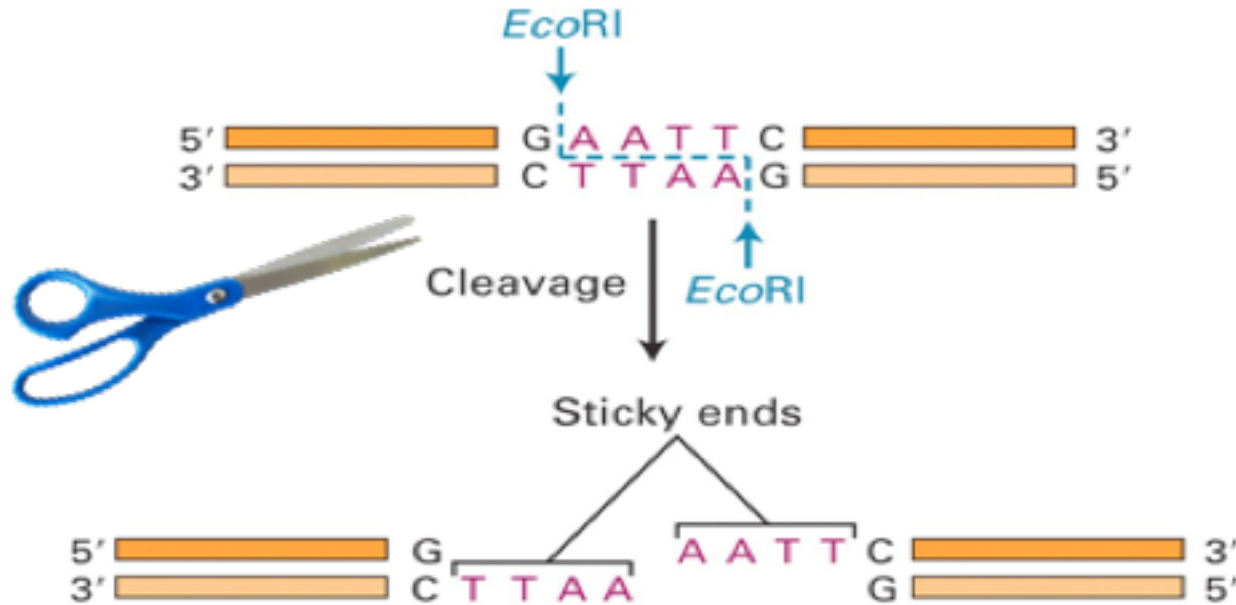
NEXT >>

PCR

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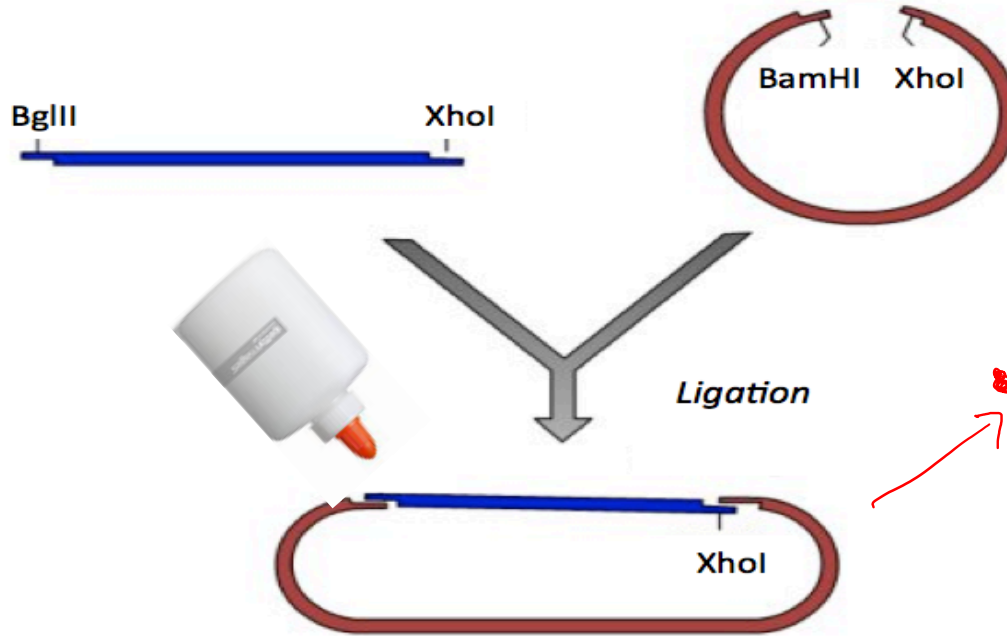
2. Digestion

Restriction endonucleases create sticky ends on dCas9 insert and plasmid backbone



3. Ligation

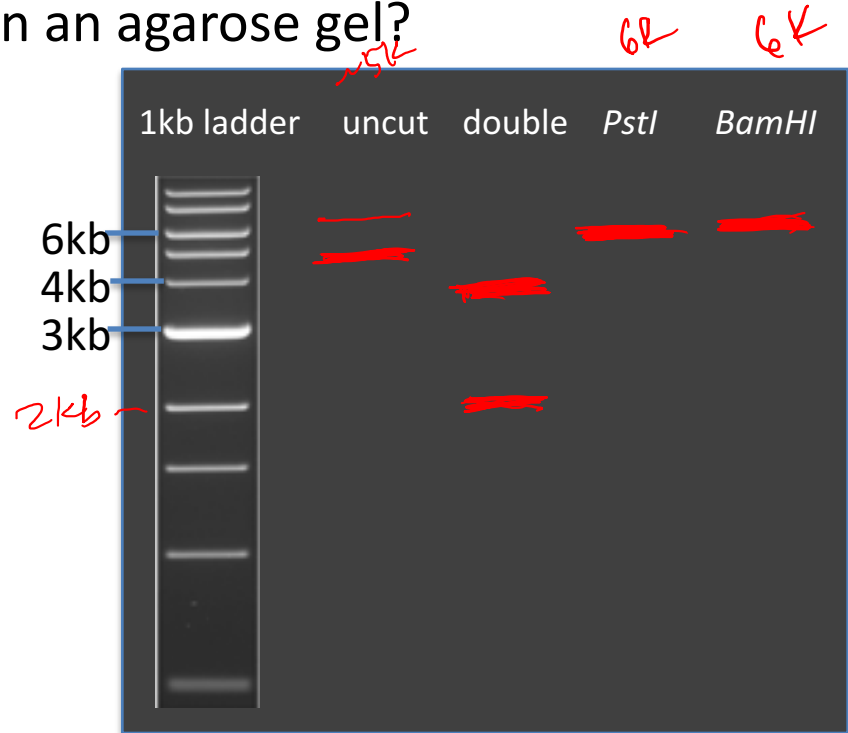
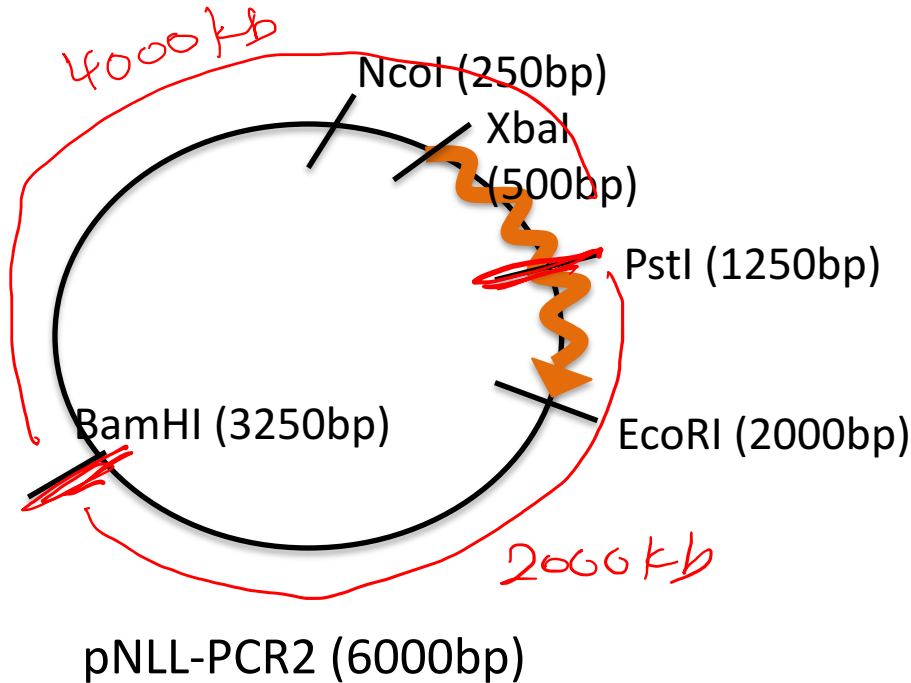
Insert dCas9 into expression vector (backbone) to create new plasmid (pdCas9)



transform into bacteria
• grow in liquid culture + purify
• conf. digest seq

Confirmation digest considerations

- Do you have access to the enzymes?
- Are the two enzymes compatible?
- Are fragments easily distinguished on an agarose gel?



M2D2 homework—Sign up for Journal Club

- Sign up on wiki for which day you will present:
M2D4 (October 24th) or M2D6 (October 31th)
- Pick 1 of 20 papers, or suggest your own
- Reserve paper by adding name next to it [LMM/WF/Red]
 - First come, first served!
 - Only one T/R and one W/F student per article
 - Don't pick a paper randomly

Slot	Day 4 (T/R)	Day 6 (T/R)	Day 4 (W/F)	Day 6 (W/F)
1				
2				
3				
4				

M2D2 homework—Make a presentation slide

To help you prepare for the Journal Club presentation, you will craft 1-2 slides using this article by Ji. et al. to present the data from Figure 2.

- Your slide(s) should show the data and highlight the key finding(s). *bullet text*
- The information should be clear and large enough to read.
- Keep text to a minimum. *No caption*
- The title should state the take-home message of the data that are shown.

Today in lab

1. Reproduce *in silico* (in Benchling) the cloning of pdCas9
 - Design primers that would amplify the gene dCas9
 - Depict PCR amplification product
 - Digestion of dCAS9 PCR product and vector by restriction enzymes
 - Ligation of insert and vector
2. Set up confirmation digests of pdCas9 for agarose gel electrophoresis (start at 3pm)
 - Choose restriction enzymes for diagnostic digest
 - Calculate volumes of digest components
 - Set-up digest and leave overnight at 37°C