

M2D7: Induce CRISPRi system

1. BE Communication workshop
2. Pre-lab
3. Analyze sequencing results
4. Prep for dCas9 induction and mixed-acid fermentation

Major assignments for M2

- **Research Article**
 - Due by 10pm on Mon., November 12th

Research Article content

1. Title
2. Abstract
3. Introduction
4. Materials and Methods
5. Figures and Results
6. Discussion
7. References

- Blog post for journal club due 10pm, 11/2
- Blog post for Mod 2 due 10pm, 11/13

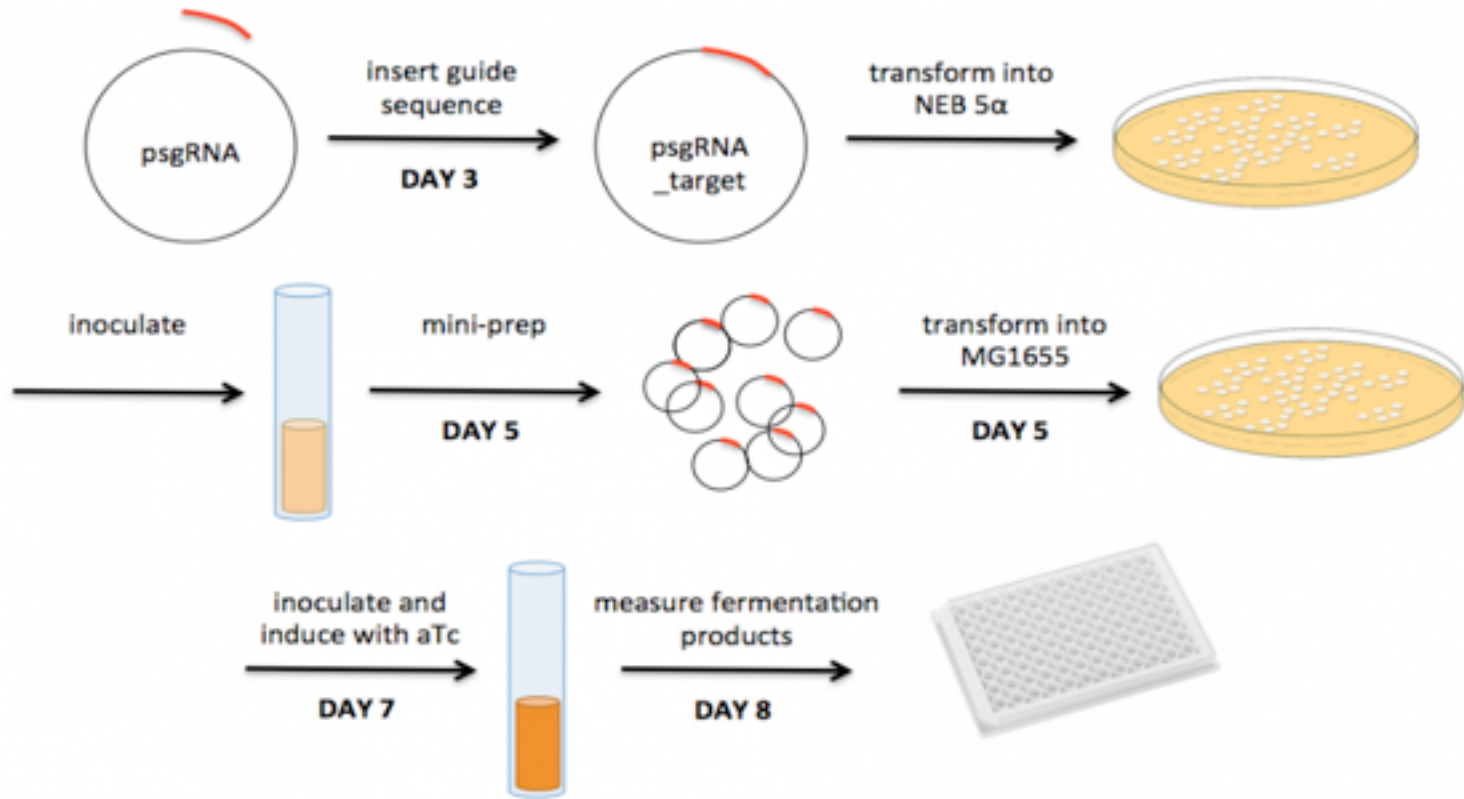
Extra office hours

11/10 (Sat): *late afternoon*
11/11 (Sun): *2-5 Noreen*

Regular office hours

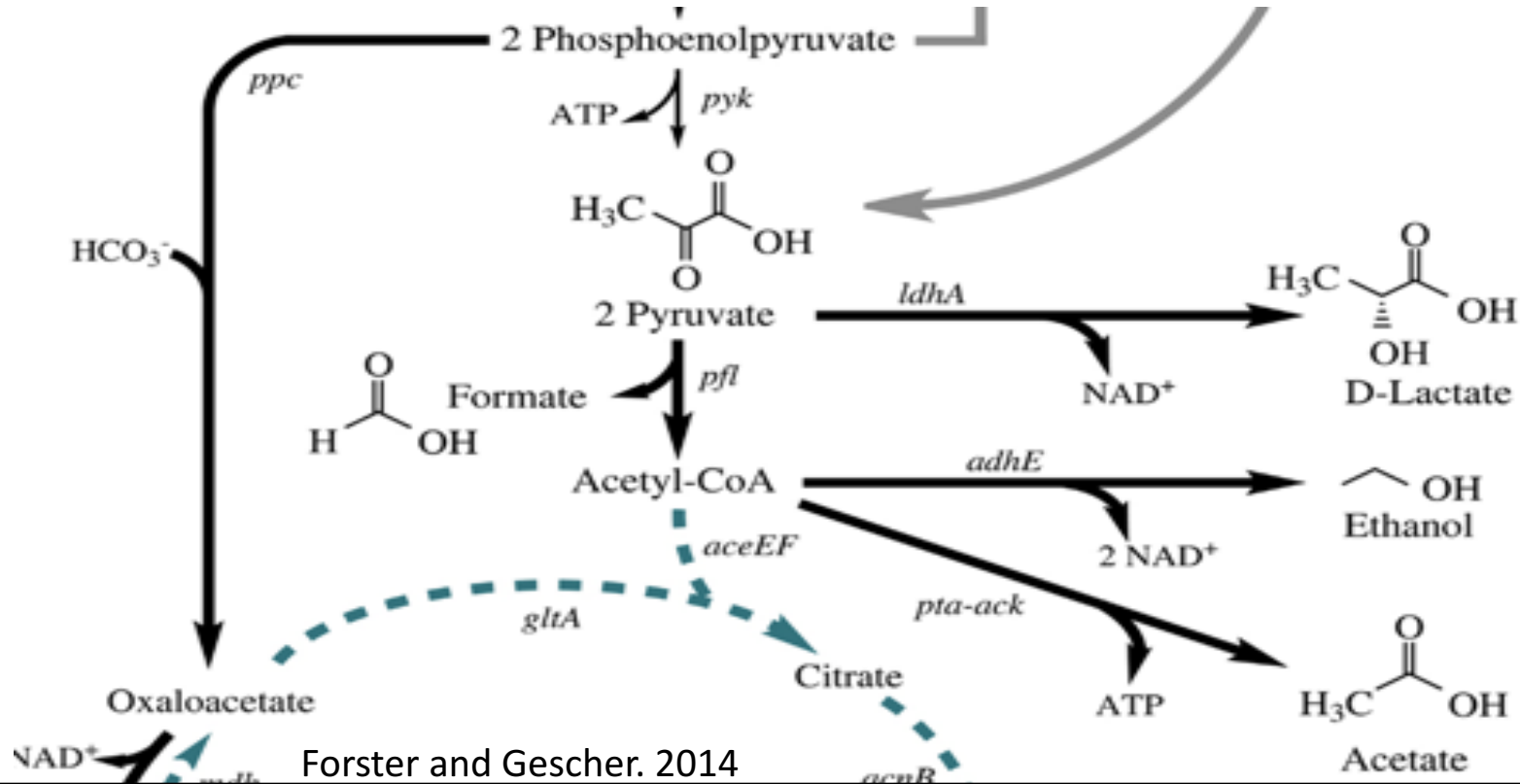
- Noreen: Mon. 2-5pm (16-317)
- Leslie: Th 2-3pm, Fr 12-1pm (56-341c)
- Josephine: W 12-1pm, Fr 2-3pm (56-341c)
- Email us to schedule a different time

M2 experimental overview

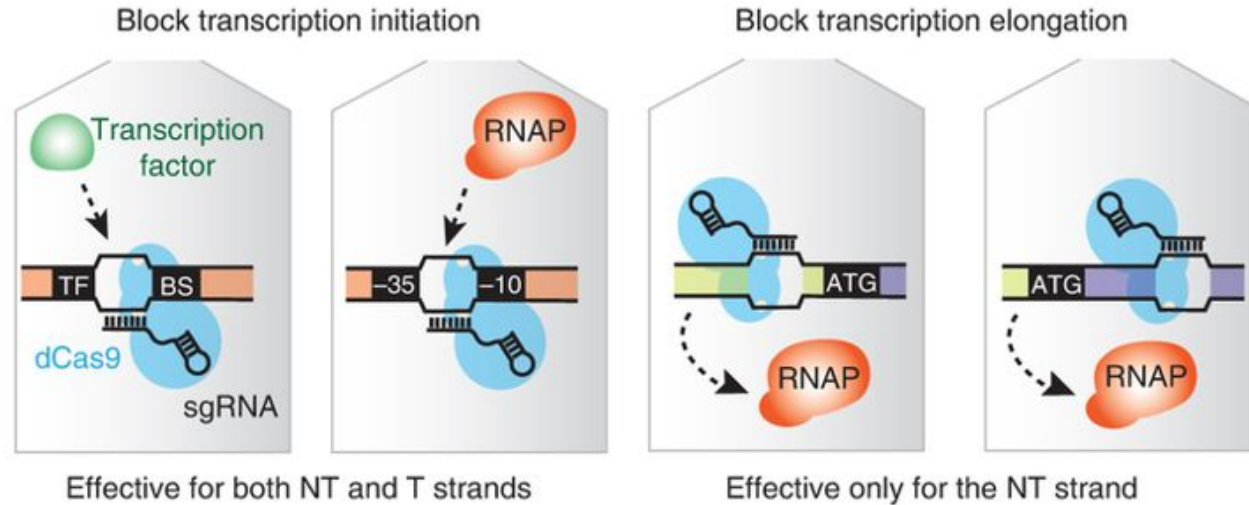
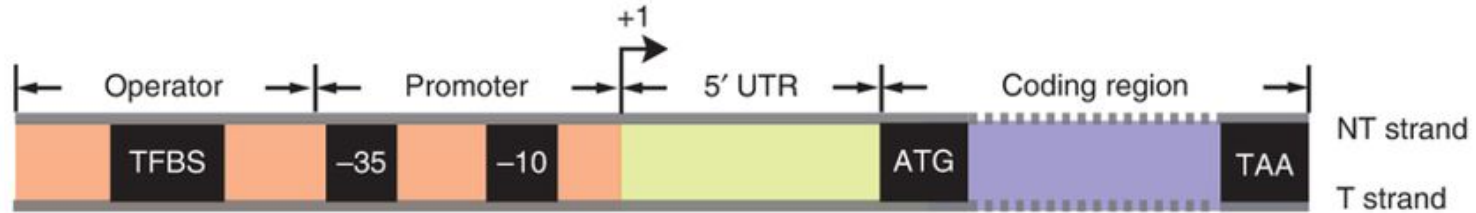


Note: sgRNA = gRNA

Using **CRISPRi** to manipulate the *E. coli* fermentation pathway

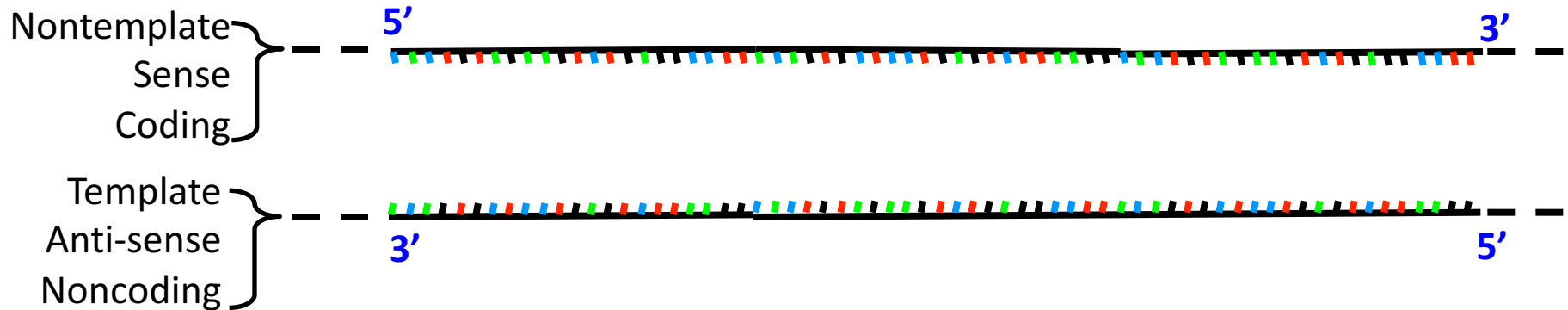


Design of gRNA for CRISPRi system



Design of gRNA for CRISPRi system

- (1) If you target the template DNA strand, the gRNA (DNA) sequence will be the same as the transcribed (nontemplate) sequence.
- (1) If you target the nontemplate strand, the gRNA (DNA) sequence will be the reverse-complement of the transcribed (template) sequence.



Please add your targeting info to the wiki today

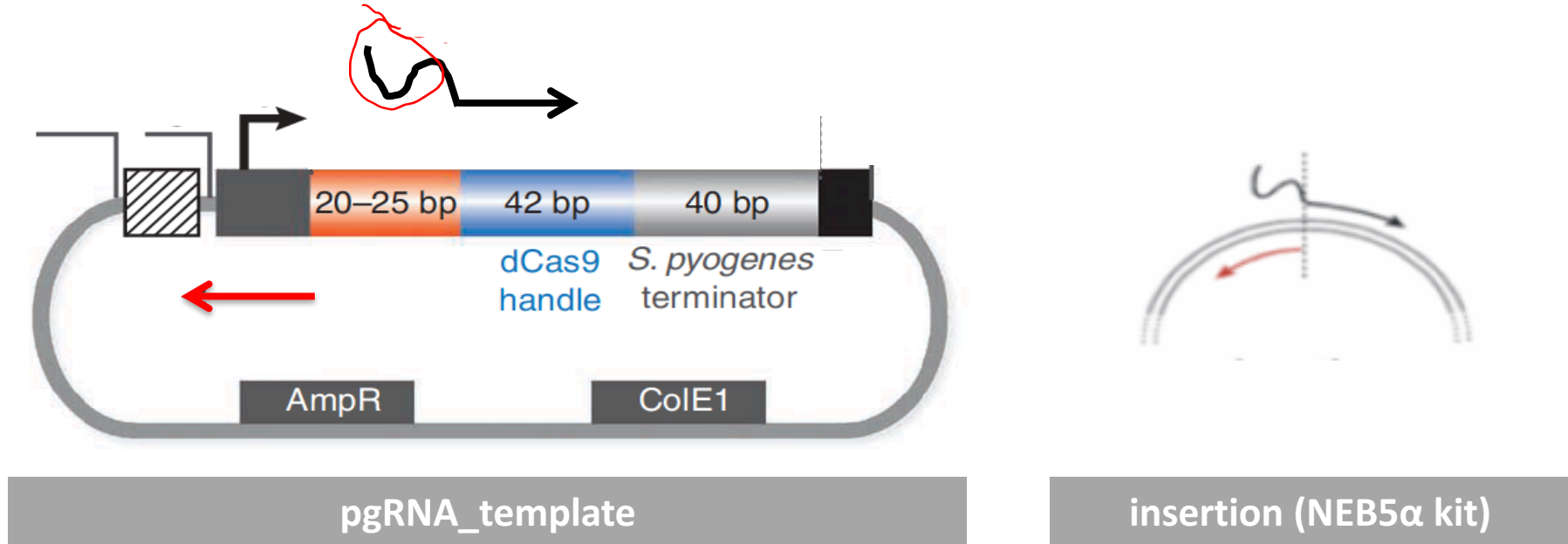
On the [Class Data](#) page



Team	Ethanol (E) or Acetate (A)	Gene targeted by CRISPRi gRNA	gRNA sequence end)
red	Ethanol	ack (indirectly, pta)	GTTTTTTTAGCC
orange	Ethanol	ldhA	ATTCAACATCAC
blue	Ethanol	ackA	TTTTTAGCCACG

...

	Locus targeted (eg. beginning of gene, putative promoter, -35 region)	Target template or nontemplate strand
TT		

M2D3: Generated pgRNA_target by SDM



← CRISPRi universal *amplification* reverse primer
forward primer including crRNA to be inserted ()
dCas9 handle ()

Analyzing Sequence Information

- Reverse and complement your reverse primer sequence before alignment
- Check whether your target sequence was successfully incorporated into the pgRNA_target plasmid

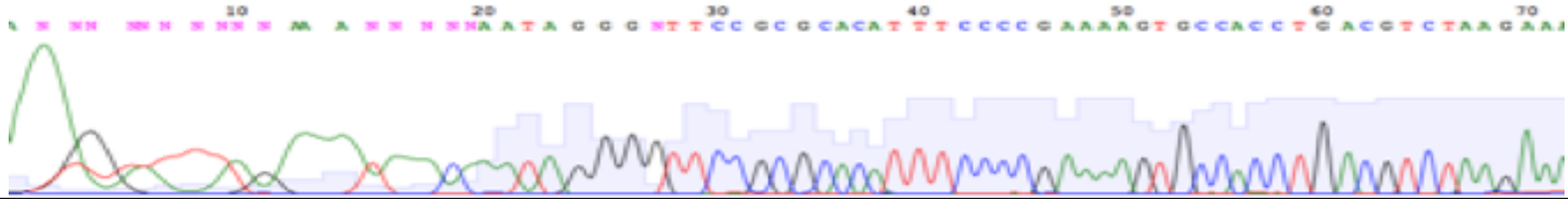
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1>---gaattctaagaatctttgacagctagctcagtcctaggtataatactagct-----gttttagagctagaaatagcaag>73
73>---GAATTCTAAGATCTTTGACAGCTAGCTCAGTCCTAGGTATATACTAGTAAATCCACTTAAGAGGTAAGGTGTGTTTTAGAGCTAGAAATAGCAAG>269
01>CTGGAATTCCTAAGATCTTTGACAGCTAGCTCAGTCCTAGGTATATACTAGTAAATCCACTTAAGAGGTAAGGTGTGTTTTAGAGCTAGAAATAGCAAG>500
1>---gaattctaagaatctttgacagctagctcagtcctaggtataatactagct-----gttttagagctagaaatagcaag>73

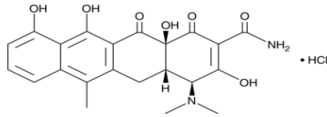
* * * * *
74>ttaaaataaggctagtcggttatcaacttgaaaaagtggcaccgagtcgggtgcttttttgaagcttgggcccgaacaaaaactcatctcagaagaggat>173
70>TTAAATTAAGGCTAGTCGGTTATCACTTGAAAGTGGCACCGAGTCGGTGCTTTTTTGAGCTTGGGCCCGAACAAAACCTCATCTCAGAGAGAGAT>369
01>TTAAATTAAGGCTAGTCGGTTATCACTTGAAAGTGGCACCGAGTCGGTGCTTTTTTGAGCTTGGGCCCGAACAAAACCTCATCTCAGAGAGAGAT>600
74>ttaaaataaggctagtcggttatcaacttgaaaaagtggcaccgagtcgggtgcttttttgaagcttgggcccgaacaaaaactcatctcagaagaggat>173

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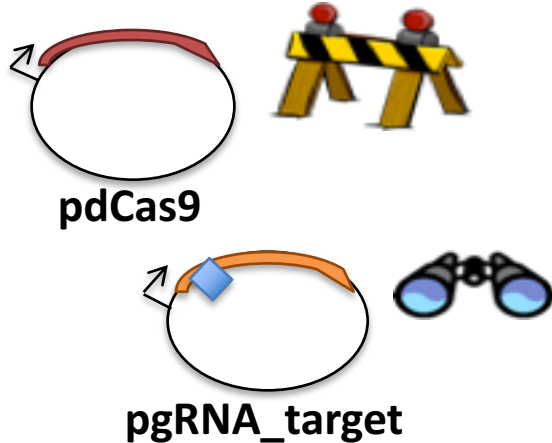
- Sanger sequencing traces are also on wiki for your reference



Induction of CRISPRi system with aTc



Anhydrotetracycline (aTc)



E. coli MG1655 genome

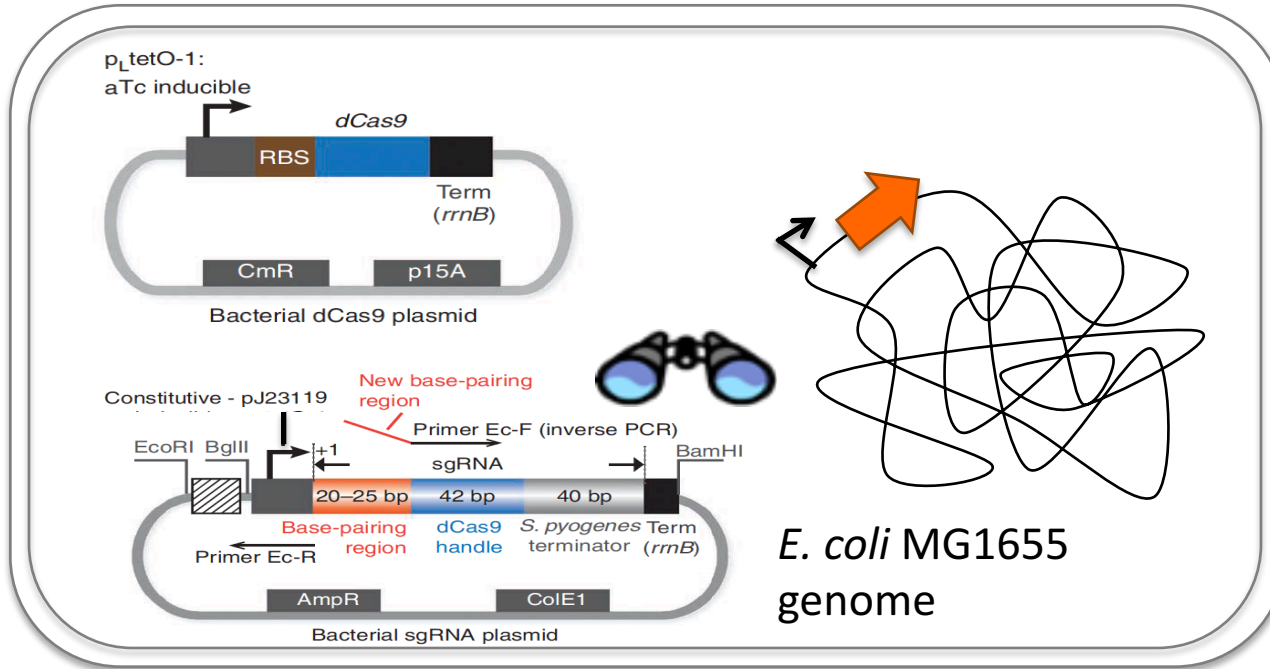
- Expressed constitutively:

gRNA

- Expression induced with aTc:

dcas9

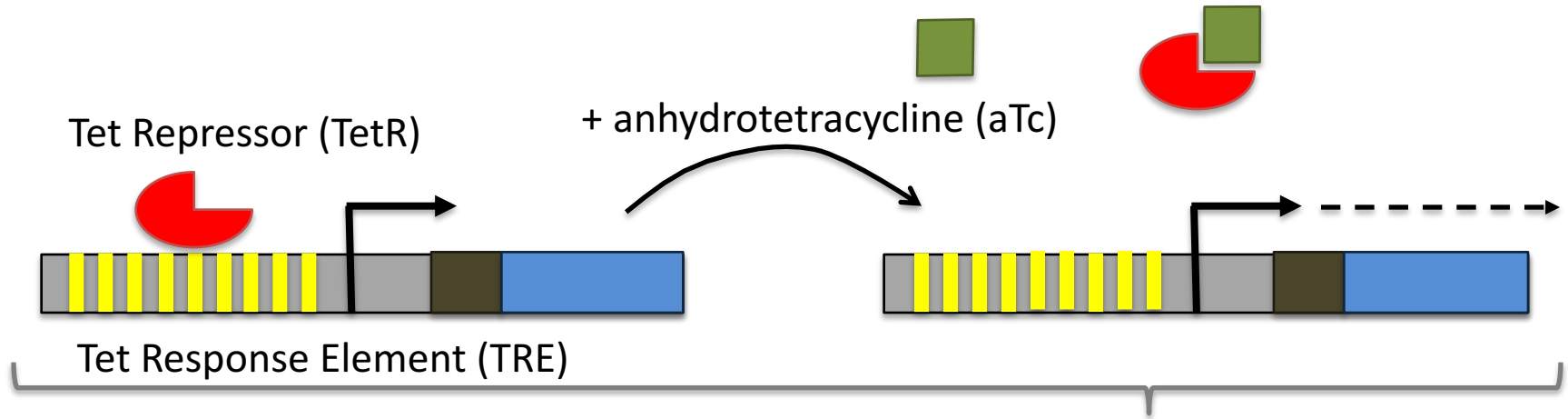
CRISPRi 'inactive' in absence of inducer



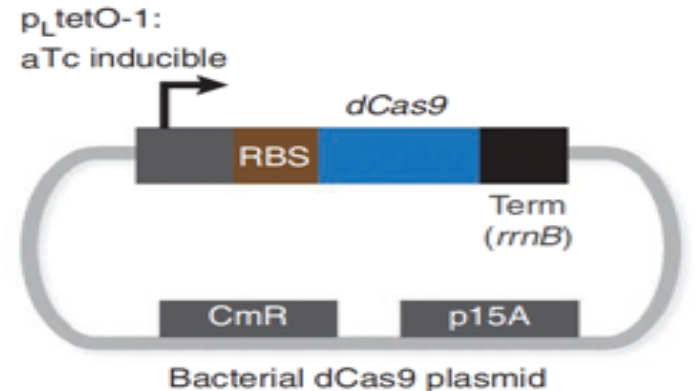
Without aTc

- Only gRNA present
- No (or little) dCas9

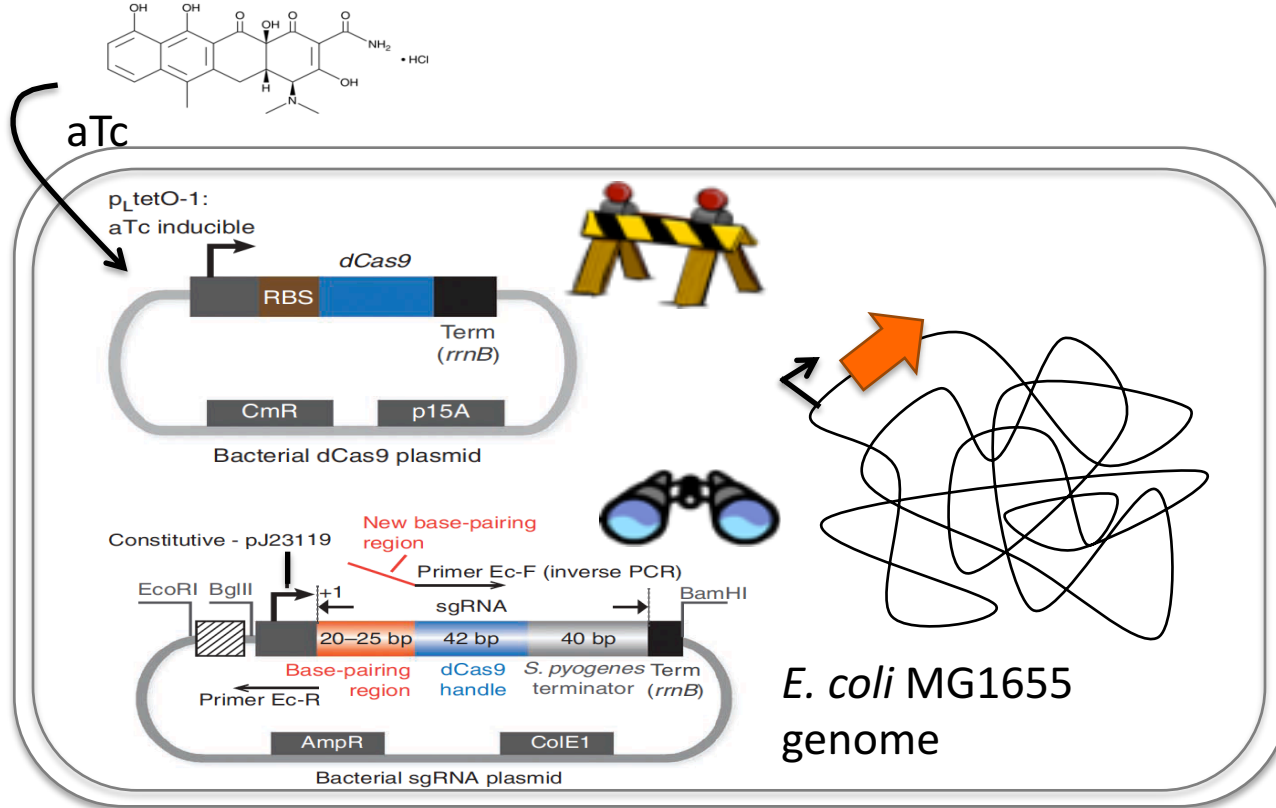
aTc induction of pdCas9



- Tet promoter regulates expression of dCas9 gene



CRISPRi 'blocks' gene expression in presence of inducer



Set up liquid cultures for mixed-acid fermentation and pdCas9 induction

- What are the necessary components?

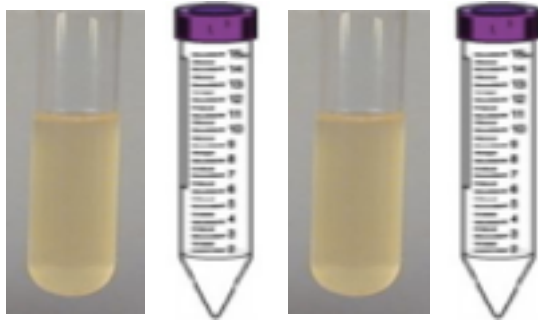
~~E. coli w/ crISPR & w/out crISPR~~
E. coli: w/ CRISPR & w/out CRISPR
media + antibiotics ATC

- What control conditions will we have?

anaerobic vs aerobic
a/c vs no a/c
transform vs no transform

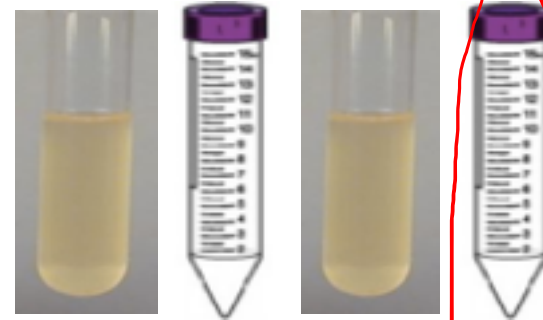
Set up liquid cultures for mixed-acid fermentation and pdCas9 induction

- Where do we expect most ethanol if hypothesis confirmed?



+ O₂ - O₂
- aTc - aTc
+ aTc + aTc

MG1655



+ O₂ - O₂
- aTc - aTc
+ aTc + aTc

MG1655 with CRISPRi

Form
FERMENTATION
+
CRISPRi

M2D8 Assignments

- Quiz on M2D8
- Peer-review methods
 - Do not leave today before receiving Methods to peer-review (soft or hard copy?)
 - Either print it out and manually indicate which part each comment refers to, type out comments, and scan to submit online, or hand in hard copy.
 - Or make digital comments with related highlights in Adobe Acrobat

Today in lab...

- Upload gRNA design information to wiki
- Download your sequencing data from discussion tab and align (using ApE software)
- Prepare media for mixed-acid fermentation inoculations