Module 2: Manipulating Metabolism

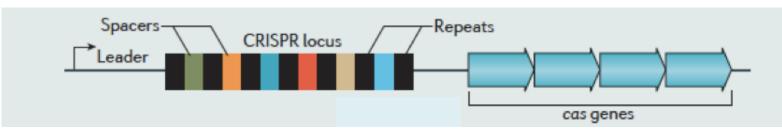
CRISPR and genetic engineering

11/1/15



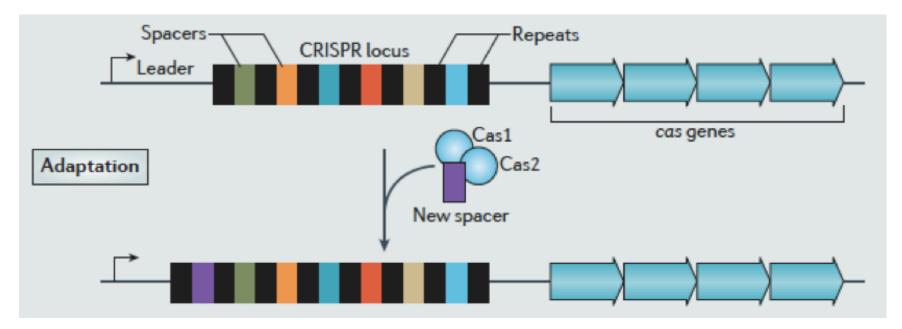
Native CRISPR system cleaves phage DNA

 Adaptive immune response encoded by CRISPR loci and Cas genes



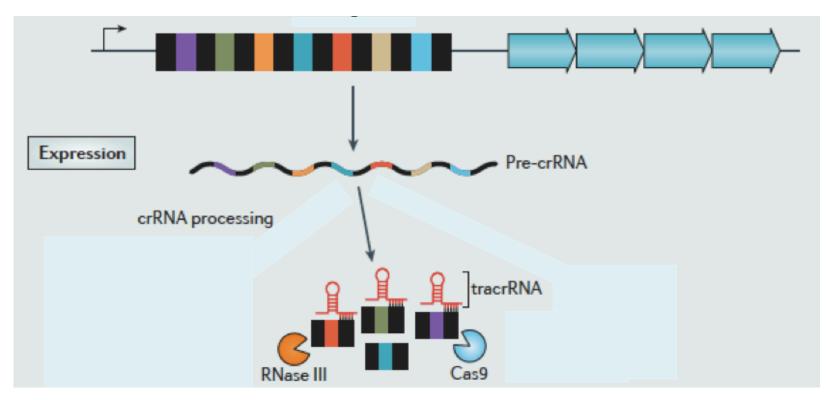
- Mechanism involves three stages:
 - Adaptation
 - Expression
 - Interference

CRISPR system: adaptation



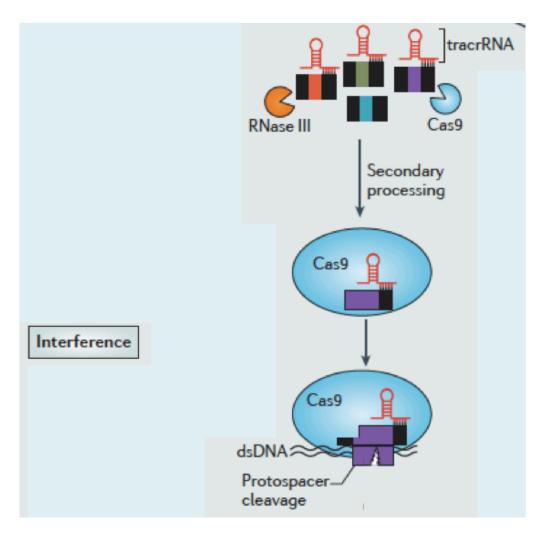
Phage DNA recognized and fragmented by restriction enzyme system, then 'spacers' incorporated into bacterial genome

CRISPR system: expression



Cas9 and Rnase III involved in processing precrRNA, then Cas9 forms complex with crRNA and tracrRNA

CRISPR system: interference



Cas9 / tracrRNA / crRNA bind invading phage DNA and cleave at target sequence that is complementary to 'spacer' sequence

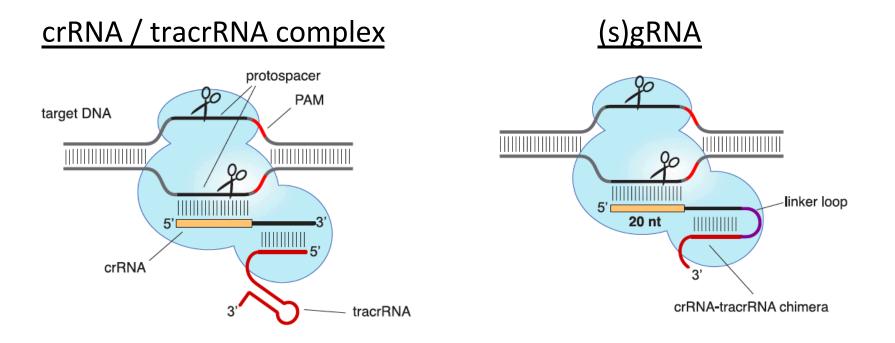
Engineered CRISPRi system

• Modifications to crRNA / tracrRNA complex?

• Modifications to Cas9?

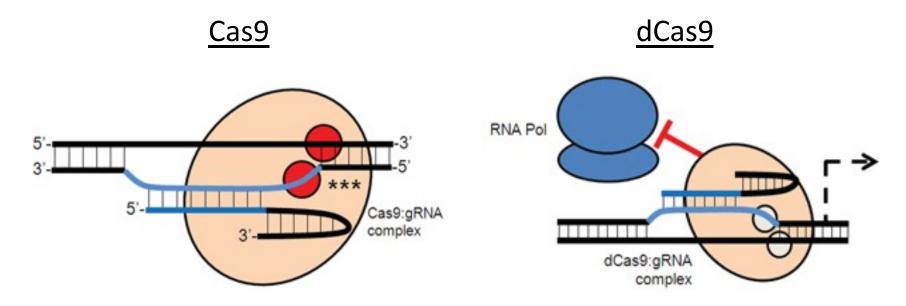
CRISPRi system: (s)gRNA

 (s)gRNA molecule is a target sequence and tracrRNA fused by a linker loop such that a single transcript used to direct Cas9 cleavage

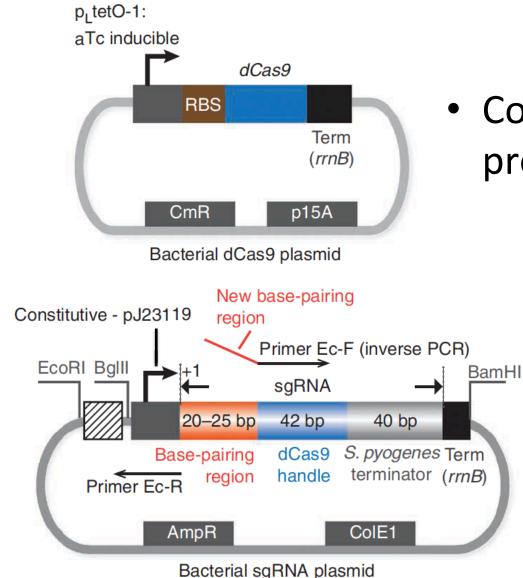


CRISPRi system: dCas9

 dCas9 protein contains mutated residues D10A and H840A that render it catalytically inactive and unable to cleave DNA, but still able to bind DNA



Closer look at pgRNA and pdCas9



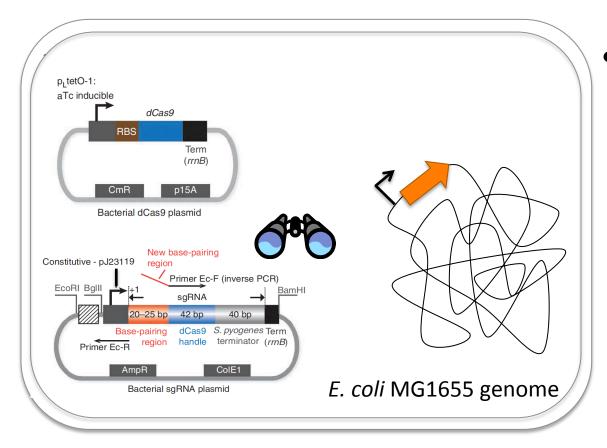
 Confirmation digest prepared on D1

> Insert (gRNA target sequence) designed on D2

Closer look at aTc induction of pdCas9 + anhydrotetracycline (aTc) Tet Repressor (TetR) Tet Response Element (TRE) p₁ tetO-1: aTc inducible Tet promoter regulates dCas9 RBS expression of dCas9 gene Term (rrnB) p15A CmF

Bacterial dCas9 plasmid

CRISPRi 'inactive' in absence of inducer

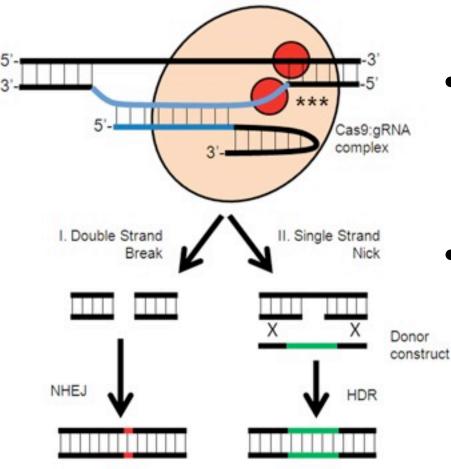


pgRNA_target
expressed
constitutively
Always
transcribed and
binding to
target gene

CRISPRi 'blocks' gene expression in presence of inducer HCI aTc pdCas9 p₁ tetO-1: aTc inducible expressed when dCas9 RBS aTc added Term (rrnB) CmR p15A – When Bacterial dCas9 plasmid New base-pairing transcribed Constitutive - pJ23119 region Primer Ec-F (inverse PCR) EcoRI Ball BamHI saRNA associates with 42 bp 40 bp dCas9 S. pyogenes Term Base-pairing handle terminator (rrnB) region pgRNA target / Primer Ec-R CoIE1 AmpR E. coli MG1655 genome Bacterial sgRNA plasmid target gene

What if we want to engineer a permanent genetic mutation?

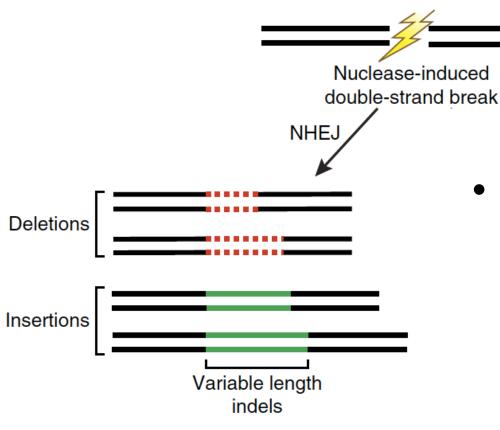
Mammalian cells able to repair dsDNA breaks



 Non-homologous end joining (NHEJ)

 Homology-directed repair (HDR) or
homologous recombination (HR)

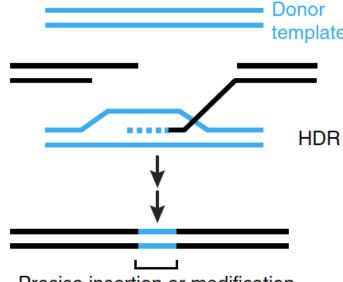
NHEJ repair generates random insertions / deletions



 Indels (insertions / deletions) result in frameshift mutations and loss of protein function

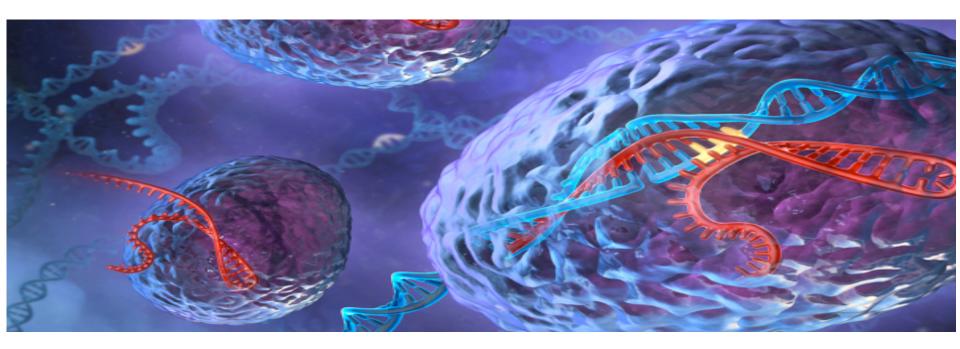
H(D)R repair enables specific sequence insertions Nuclease-induced double-strand break Insertions used to Donor template incorporate protein tags / new gene sequences and to modify

the native DNA sequence



Precise insertion or modification

Final overview of CRISPR / Cas9



https://www.youtube.com/watch?v=k99bMtg4zRk

In the *laboratory*...

- Journal club presentations
 - Meet at 1p in 16-336 for M2Q2

