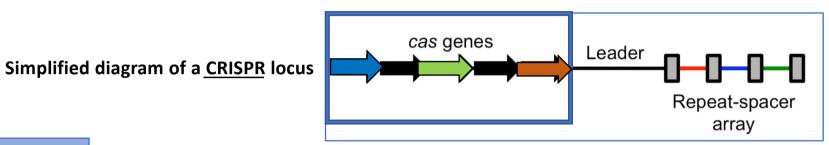
LOSS OF FUNCTION - theory

How does this genetic material in CRISPR locus then manage to kill bacteria?





The system can be slighty different in different types of bacteria but the best studies one is *Streptococcus pyogenes* so we will focus on that one

For the sake of simplicity lets focus on the 3 Cas genes (now colored arrows) most important for genetic engineering;



Codes for a **trans-activating CRISPR RNA (tracrRNA**) that will help in the process of ensuring the whole process only cuts bacteriophage DNA



Codes **for a protein** that is a nuclease that cuts DNA but only if it is given a very specific set of signals to do so (otherwise it would potentially damage the bacteria's own DNA). The most common one used in genetic engineering approaches is called Cas9. ; additional Cas1 and Cas2 are responsible for spacer geration

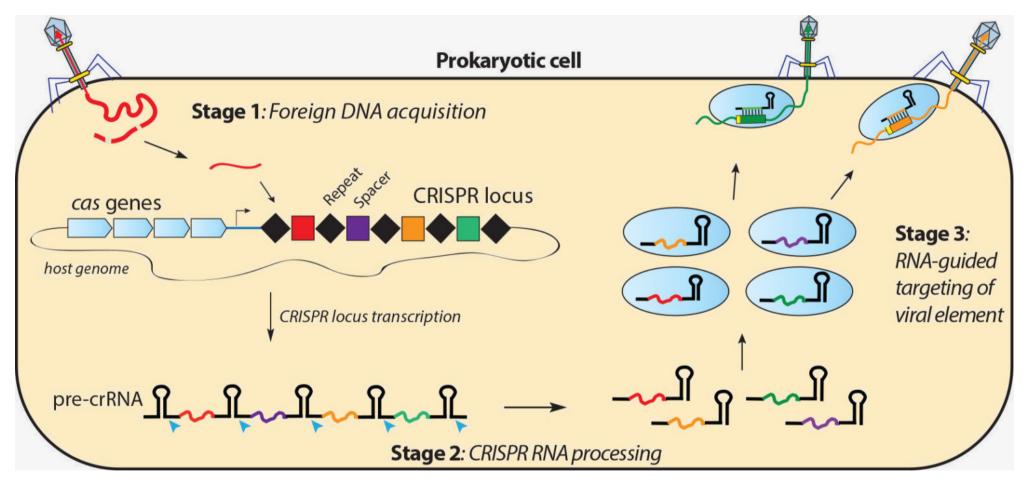


Codes for a very specific piece of RNA (**crRNA or guide RNA**) that will help in the process of ensuring the whole process only cuts bacteriophage DNA

For now lets not worry about the other genes in the Cas locus

Acquisition of immunity

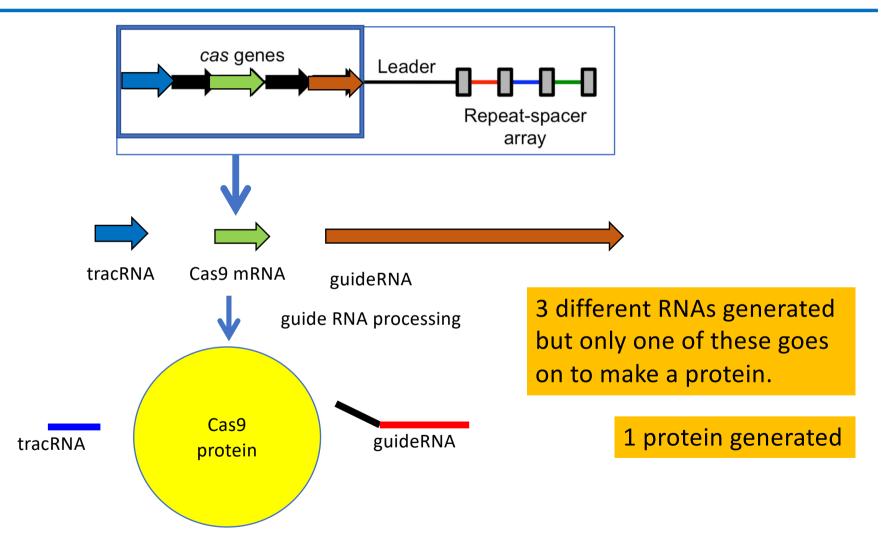
Adaptive immunity

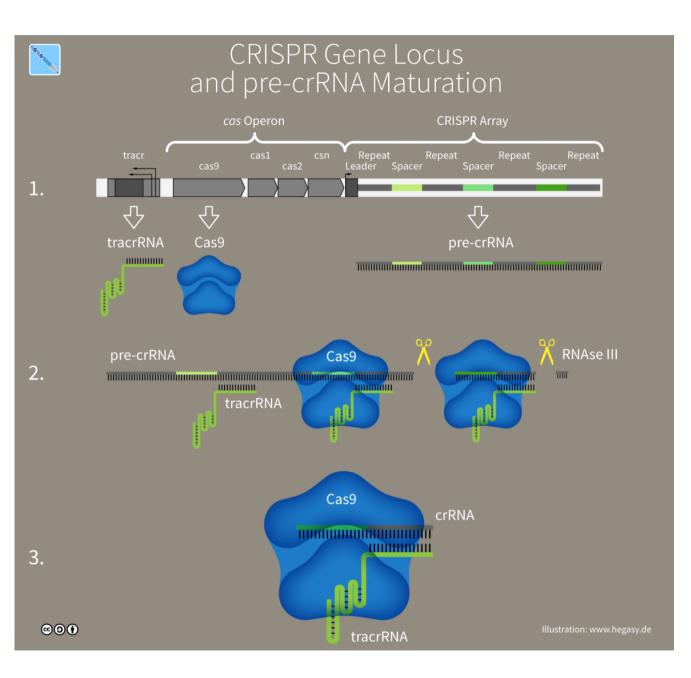


Cas1 and Cas2 are responsible for spacer generation: Bioinformatic analysis of regions of phage genomes that were excised as spacers (termed protospacers) revealed that they were not randomly selected but instead were found adjacent to short (3 - 5 bp) DNA sequences termed protospacer adjacent motifs (PAM).

LOSS OF FUNCTION - theory

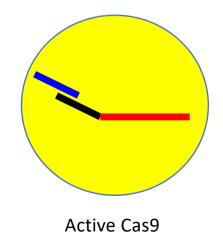
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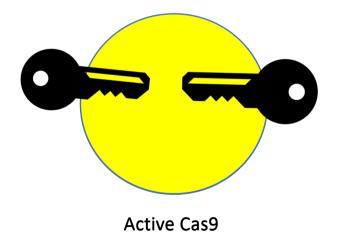
What is Cas9?

- Cas9 is an endonuclease that can cut double stranded DNA
- Cas 9 is only activated when the tracRNA and the guide RNA are associated with it (i.e it is a nucleoprotein). Imagine this a bit like the fail safe mechanism they use to prevent accidental launch of nuclear missiles where 2 people have to insert keys at exactly the same times
- In fact the tracRNA and the guide RNA have a short overlapping sequence that means they actually have to bind to each other in this complex for this to work properly

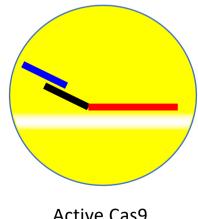


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- Cas9 has a channel that DNA can fit into.
- It scans the DNA looking for sequence that match the guide sequence

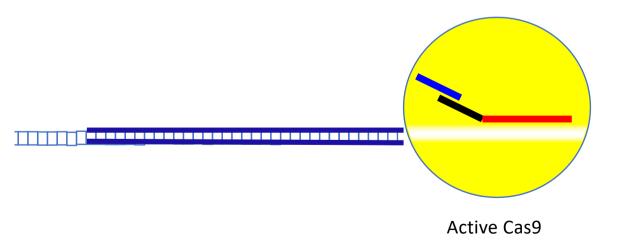




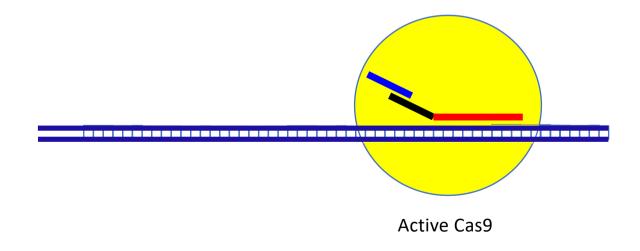
Active Cas9

How Cas9 works?

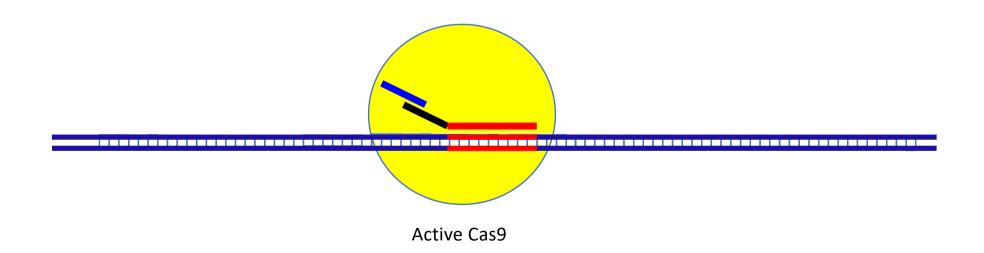
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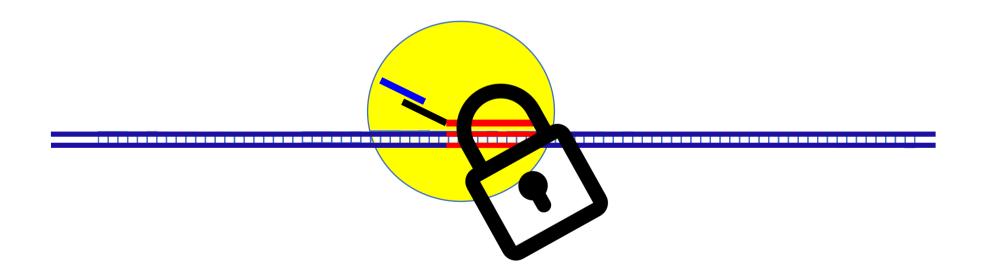
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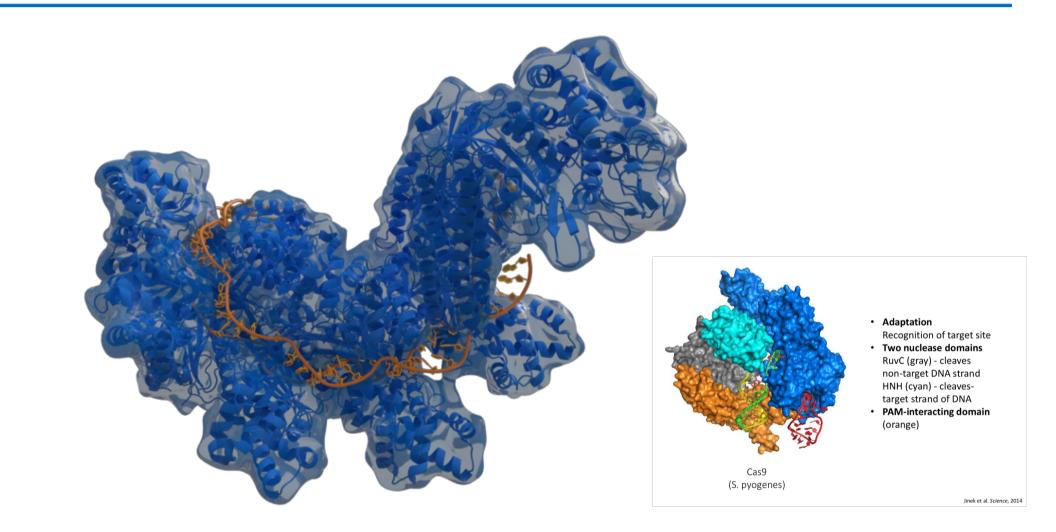
• When a DNA sequence complementary to the guide RNA is found the scanning stops



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Structure of DNA bound to a Cas enzyme

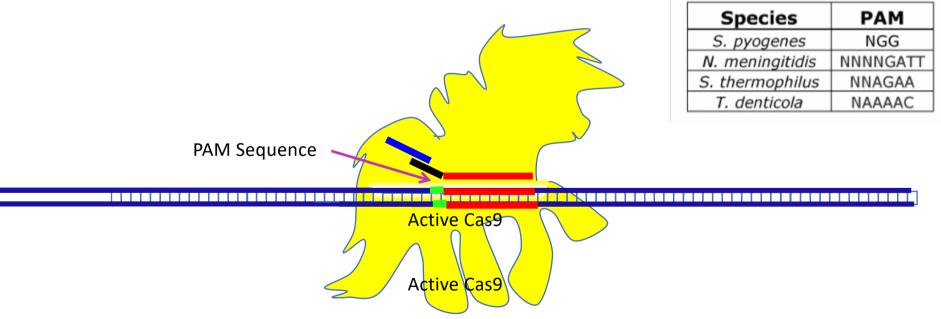


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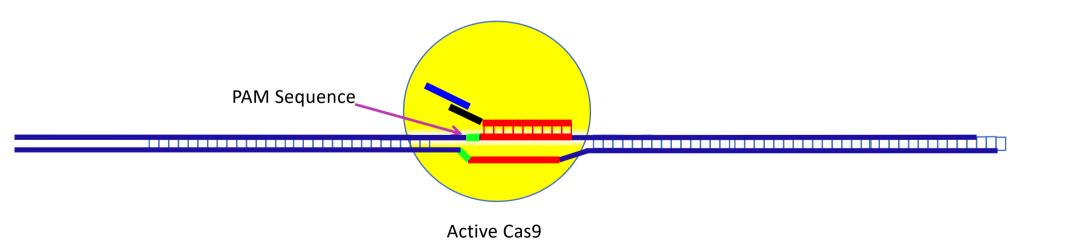
Completely irrelevant aside



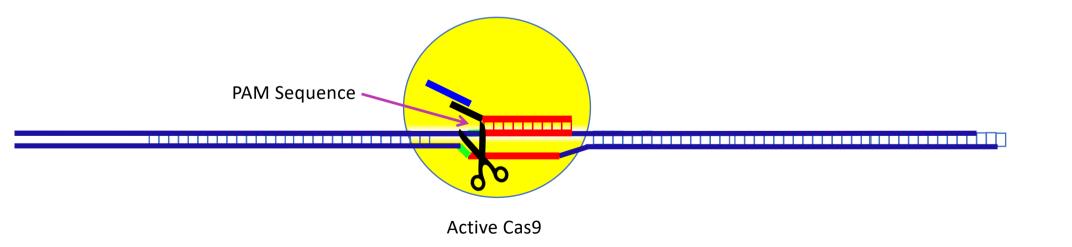
- There is one additional check
- In this control step the target site in the bacteriophage DNA needs to have the PAM sequence (Protospacer Adjacent Motif)
- PAM sequences DO NOT APPEAR in the bacterial genome
- PAM sequences are required for Cas9 endonucease activity
- PAM sequences are specific for bacterial strains and protect the Cas locus from beeing cut by Cas9



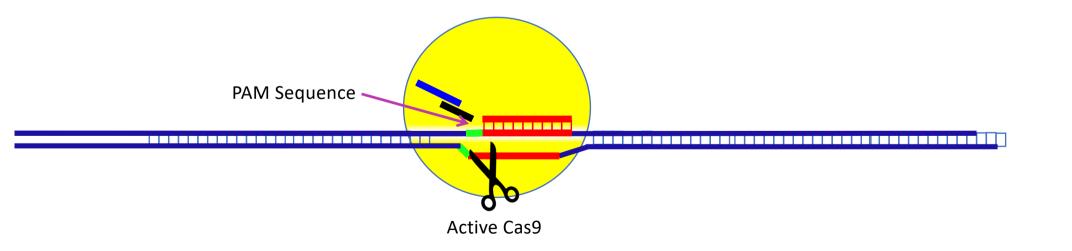
• Now the RNA binds to the complementary strand of the DNA and opens up the DNA helix



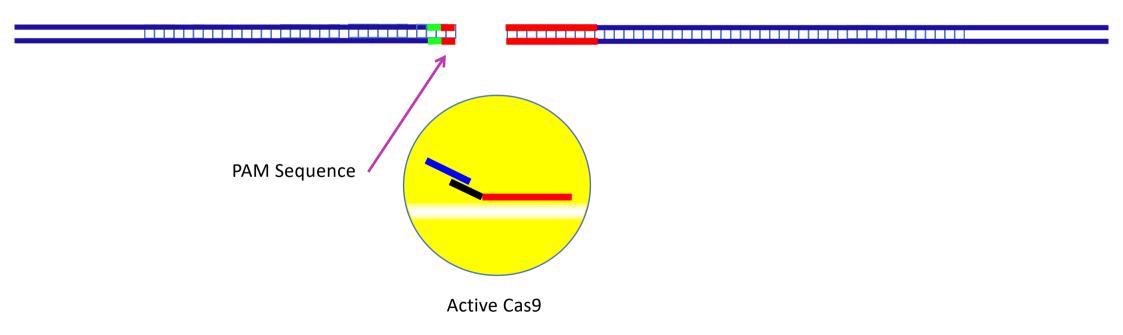
• Now the bacteriophages DNA gets cut very close to the PAM site

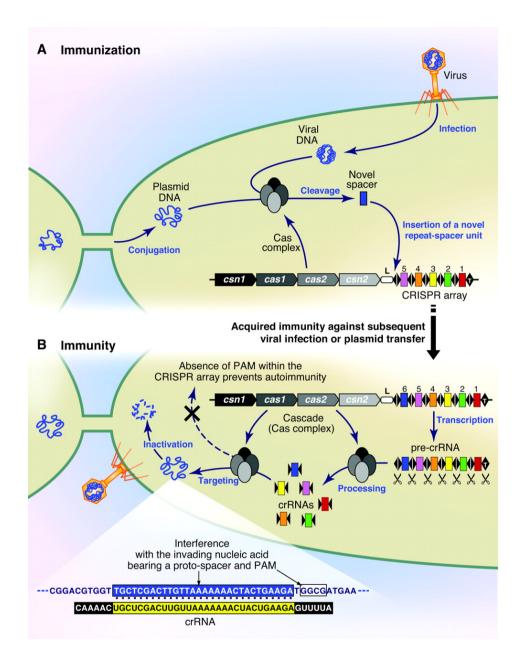


• Now the bacteriophages DNA gets cut very close to the PAM site



• Now the bacteriophages DNA gets cut very close to the PAM site, it looks like this and the bacteriophage is essentially inactivated



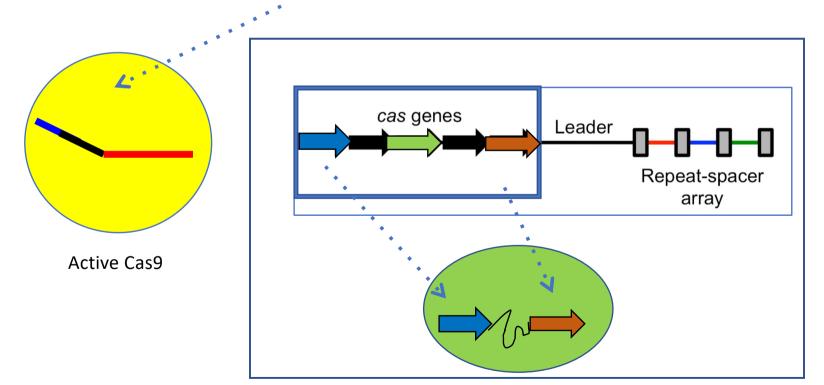


Protospacer adjacent motif (PAM) is a 2-6 base pair DNA sequence immediately following the DNA sequence targeted by the Cas9 nuclease in the CRISPR bacterial adaptive immune system.

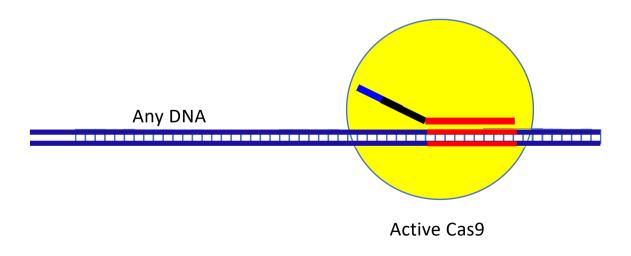
PAM is a component of the invading virus or plasmid, but is not a component of the bacterial CRISPR locus. Cas9 will not successfully bind to or cleave the target DNA sequence if it is not followed by the PAM sequence.

PAM is an essential targeting component (not found in bacterial genome) which distinguishes bacterial self from non-self DNA, thereby preventing the CRISPR locus from being targeted and destroyed by nuclease.

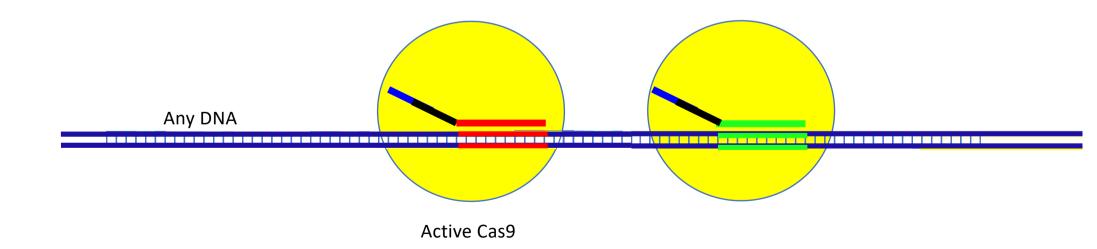
 Jennifer Doudna and Emmanuelle Charpentier re-engineered the Cas9 endonuclease into a more manageable two-component system by fusing the two RNA molecules tracRMA and guide RNA (or crRNA) into a "SINGLE-GUIDE RNA" (sgRNA) that, when combined with Cas9, could find and cut the DNA target specified by the guide RNA



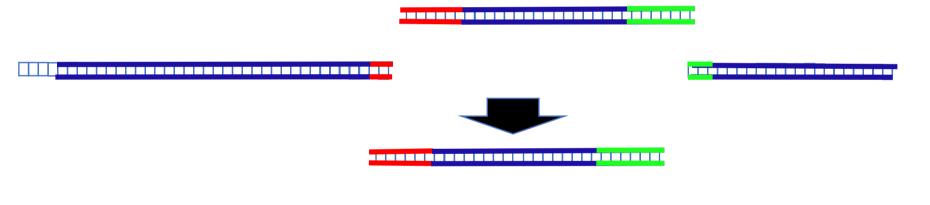
- This means we can artificially make a sgRNA that can be designed to target **any part of the genome** (as long as it has an appropriate PAM sequence nearby)
- All we have to do is artificially express the Cas9 and the sgRNA together



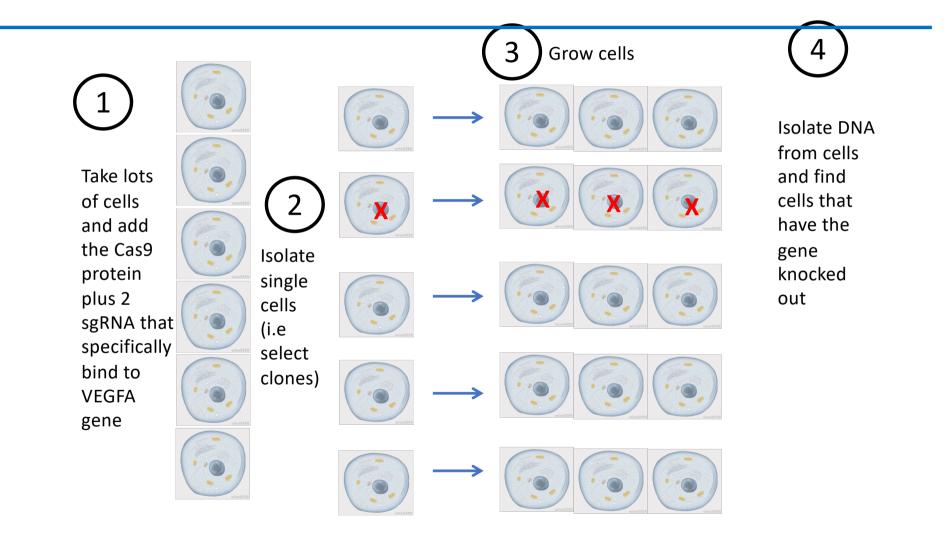
 We can put two different sgRNA into the same protein and cut at 2 places in the genome . — we can cut out large regions of DNA.



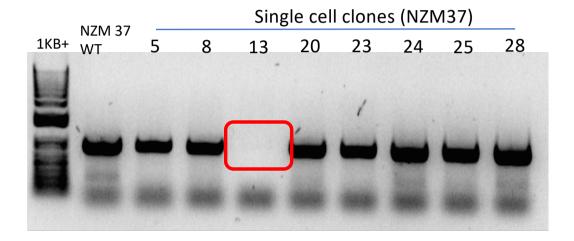
This allows us to selectively "knock out" regions of the genome



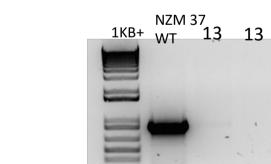
Just an example: Knockout of VEGF-a gene



Just an example: Knockout of VEGF-a gene

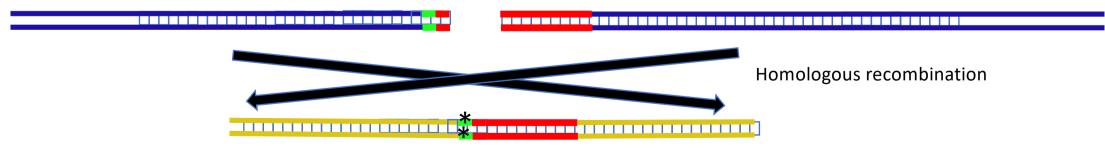


Here is an example of PCR of the VEGFA gene of melanoma cells where we have tried to use CRISPR to "knockout the VEGFA gene (achieved in clone 13)



Repeat PCR

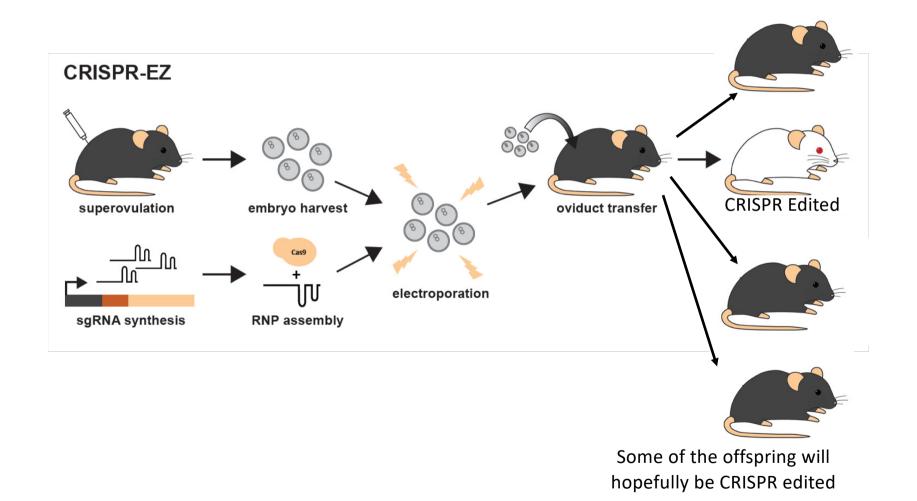
- We can use an artificial piece of DNA that is identical to the cleaved region of DNA (with "corrected" sequence -*-) -----
- when the cell tries to repair its own chromosomal DNA it will sometimes accidentally incorporate this into its own DNA by homologous recombination!
 - \rightarrow Bring in CRISP-Cas9 components to make a specific cut
 - ightarrow Bring in a DNA fragement that contains the desired genetic alteration
 - (wt \rightarrow mutant; mutant \rightarrow wt)
 - \rightarrow Strand invasion by cut sequence and HR



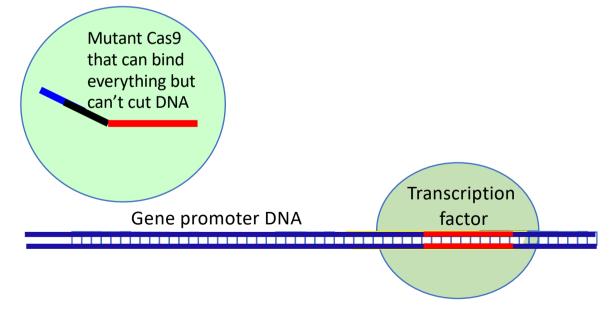
• Now the artificially produced piece of DNA is "knocked in" to the genome



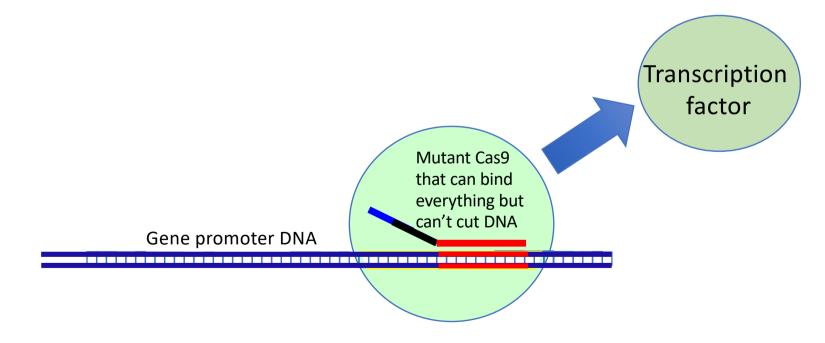
Making mice where genes are knocked out is easier and cheaper



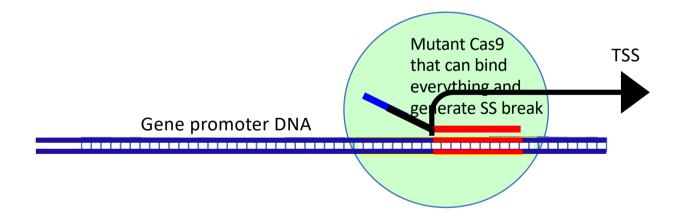
- Uses a mutant Cas9 that can bind everything but can't cut DNA
- This means it locks on tightly to the DNA that matches the guide sequence and stops on that region of DNA
- An example of how this can be used is by having a big Cas9 protein sitting at specific transcription factor binding site: we can block the transcription factor from coming into the gene promoter and therefore we switch off the expression of that specific gene controlled by TF in a highly targeted way.
- Negative aspect: keeps DNA in a short RNA:DNA hybrid state



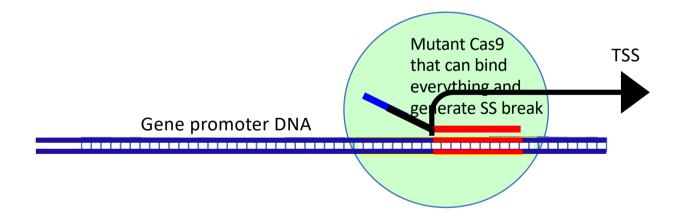
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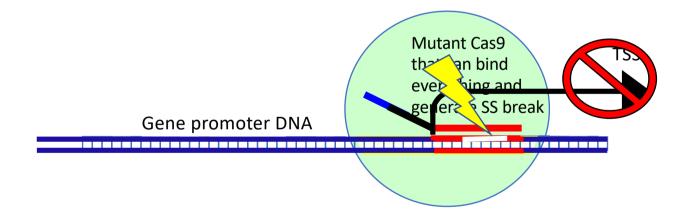
- Alterantively we can use a mutant Cas9 that can bind everything but generates a single strand cut DNA
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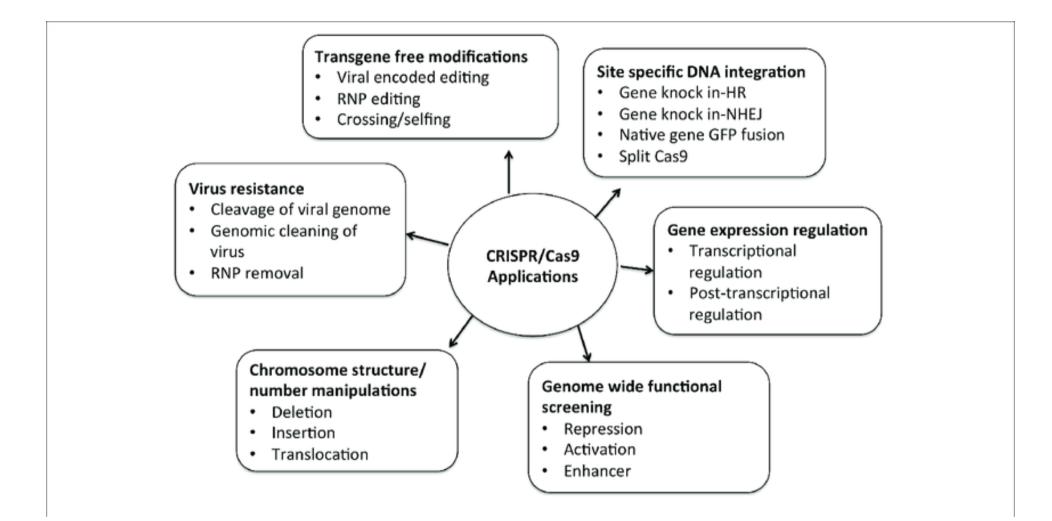


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Using CRISPR a weapon to wipe out mosquitos

A gene drive is a genetic engineering technology that can propagate a particular suite of genes throughout a population. Gene drives can arise through a variety of mechanisms. They have been proposed to provide an effective means of genetically modifying specific populations and entire species.

