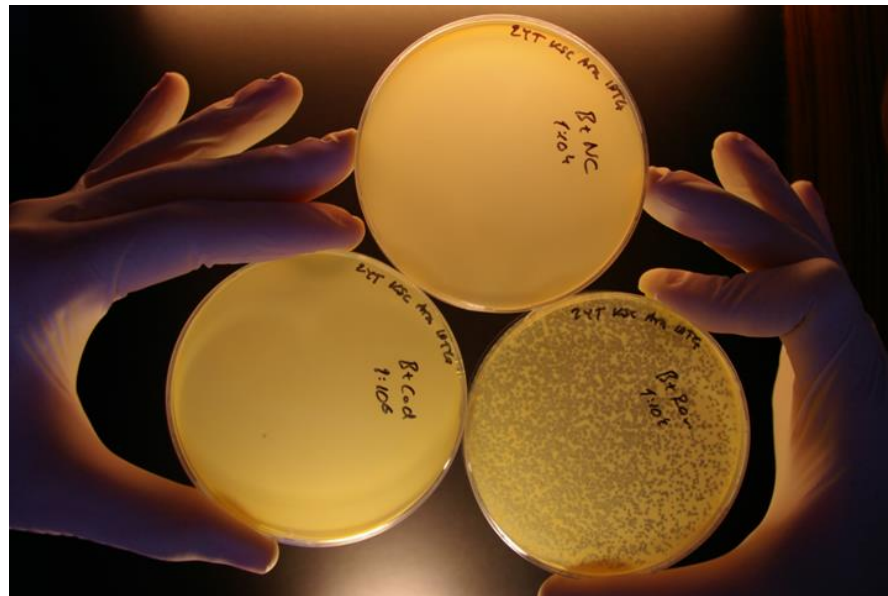


# CRISPR-Cas systems

*history, biology & applications*



John van der Oost

Laboratory of Microbiology  
Wageningen University

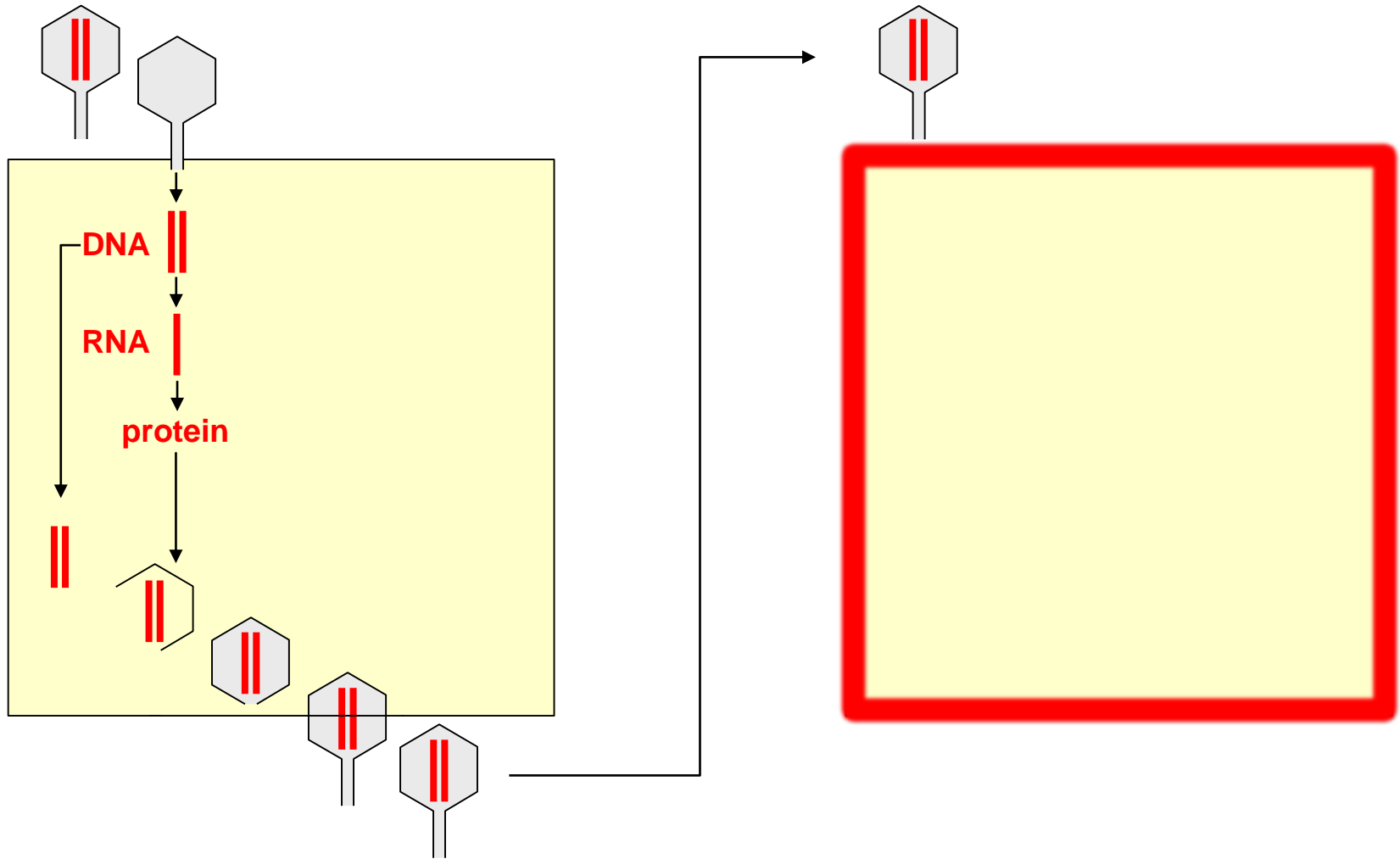




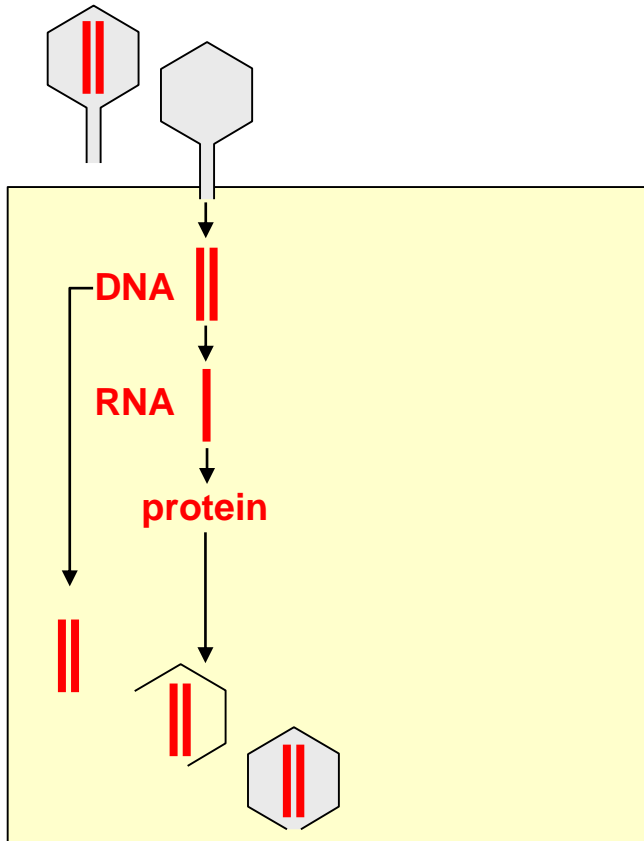
## History - *discovery & diversity*

- Biology - *anti-virus defence mechanism*
- Applications - *genome editing & gene therapy*

# virus infection of bacteria



# anti-virus systems of bacteria



## known anti-virus systems

- attachment
- inhibit DNA injection
- degradation of alien DNA
- suicide system

## new anti-virus system

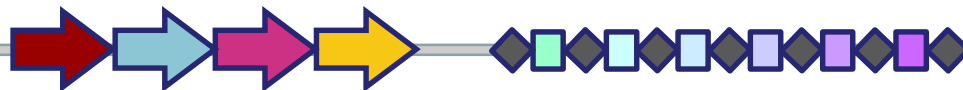
- CRISPR-Cas

# CRISPR-Cas - *discovery*



```
GAGTTCCCCGCGCCAGCGGGGATAAACCGCTTTCGCAGACGCGCGGGCGA  
TACGCTCACGCA GAGTTCCCCGCGCCAGCGGGGATAAACCGCAGCCGAA  
GCCAAAGGTGATGCCGAACACGCTGAGTTCCCCGCGCCAGCGGGGATAA  
ACCGGGCTCCCTGTTCGGTTGTAATTGATAATGTTGAGAGTTCCCCGCGC  
CAGCGGGGATAAACCGTTTGGATCGGGTCTGGAATTTCTGAGCGGTTCGC  
GAGTTCCCCGCGCCAGCGGGGATAAACCGCGAATCGCGCATACCCTGCG  
CGTCGCCGCCTGC GAGTTCCCCGCGCCAGCGGGGATAAACCGTCAGCTT  
TATAAATCCGGAGATACGGAAACTA GAGTTCCCCGCGCCAGCGGGGATA
```

- CRISPR – clustered regularly interspaced palindromic repeats
- Cas – CRISPR-associated genes & proteins

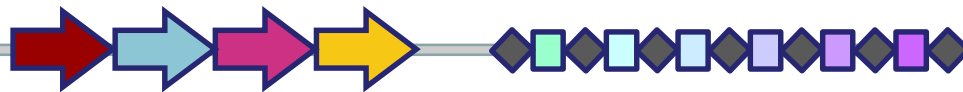


# CRISPR-Cas system – *adaptive immunity*

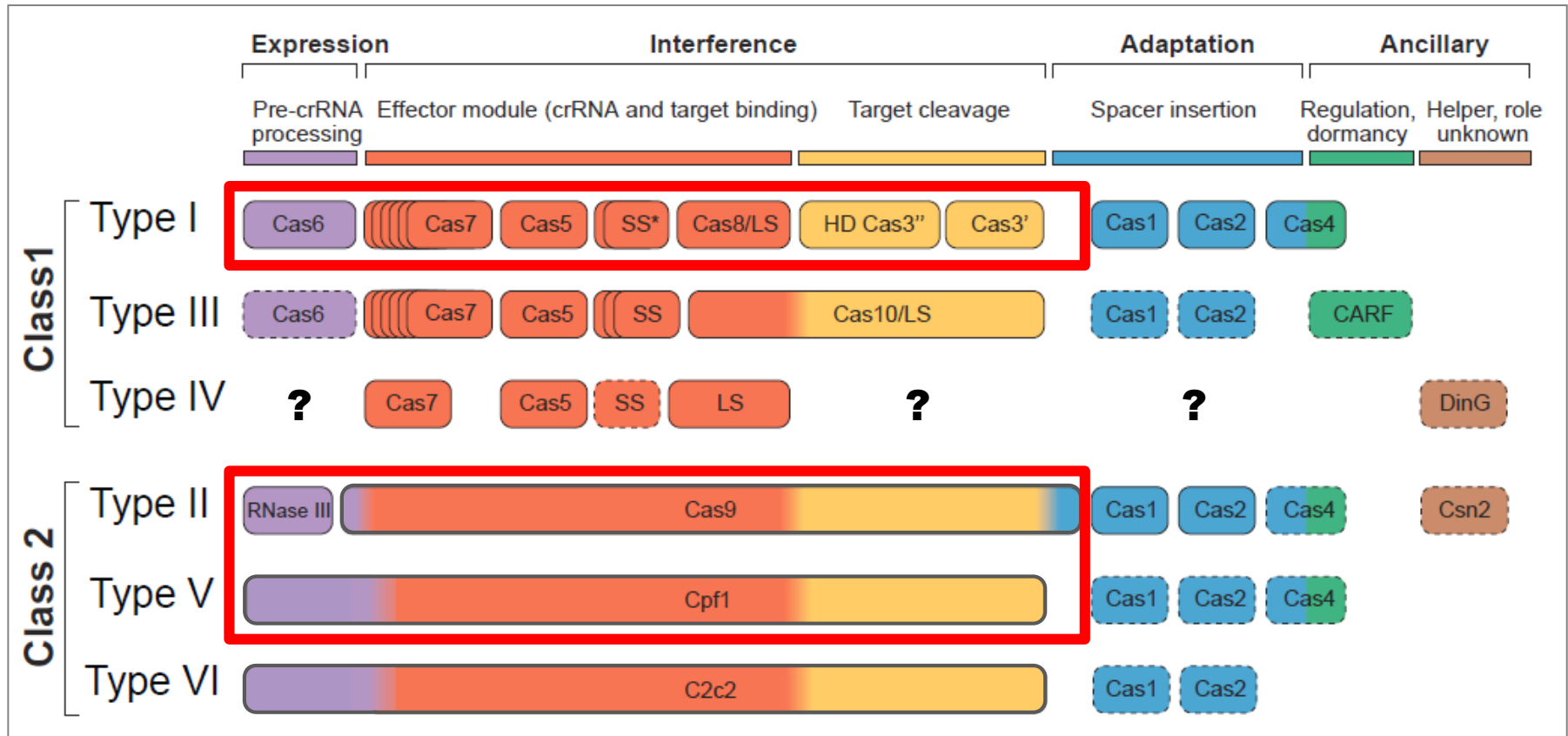


```
GAGTTCCCCGCGCCAGCGGGGATAAACCGCTTTCGCAGACGCGCGGGCGA  
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CAGCGGGGATAAACCGTTTGGATCGGGTCTGGAATTTCTGAGCGGTTCGC  
GAGTTCCCCGCGCCAGCGGGGATAAACCGCGAATCGCGCATACCCTGCG  
CGTCGCCGCCTGC GAGTTCCCCGCGCCAGCGGGGATAAACCGTCAGCTT  
TATAAATCCGGAGATACGGAAACTA GAGTTCCCCGCGCCAGCGGGGATA
```

- many CRISPR spacers are homologous to viruses or plasmids
- hypothesis: novel prokaryotic defence system – RNA interference ?
- present in genomes of 40% of bacteria and 85% of archaea



# CRISPR-Cas diversity – 2 classes / 6 types



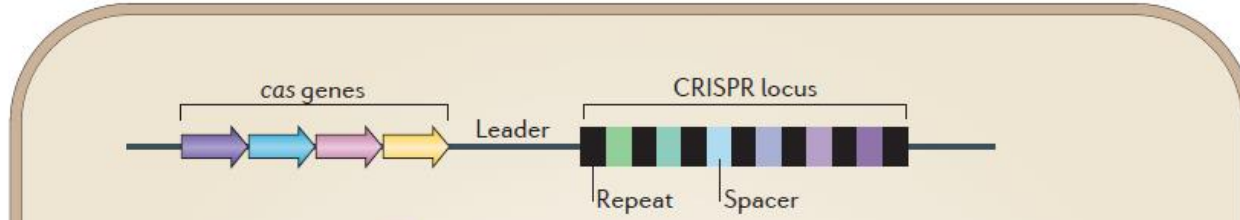
# outline



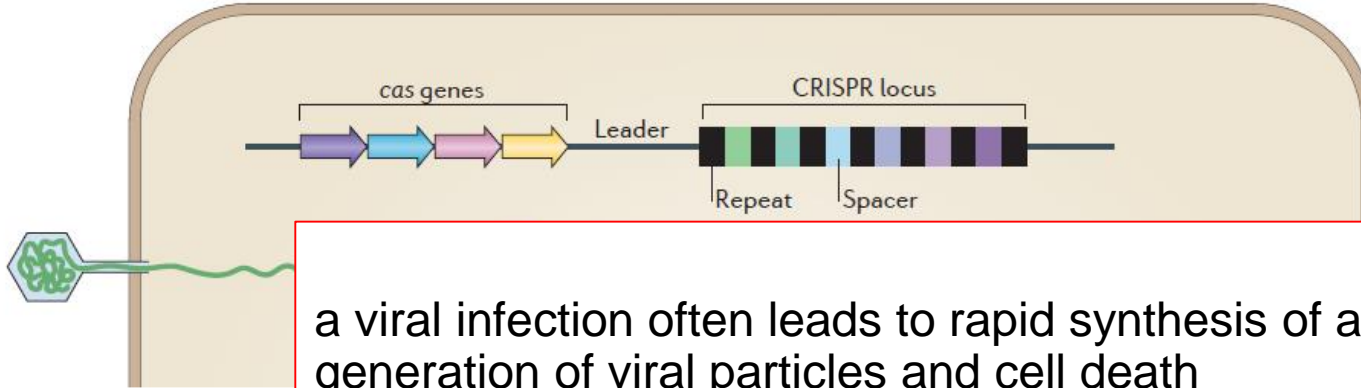
- History - *discovery & diversity*
- ➔ Biology - *anti-virus defence mechanism*
- Applications - *genome editing & gene therapy*



# CRISPR-Cas mechanism



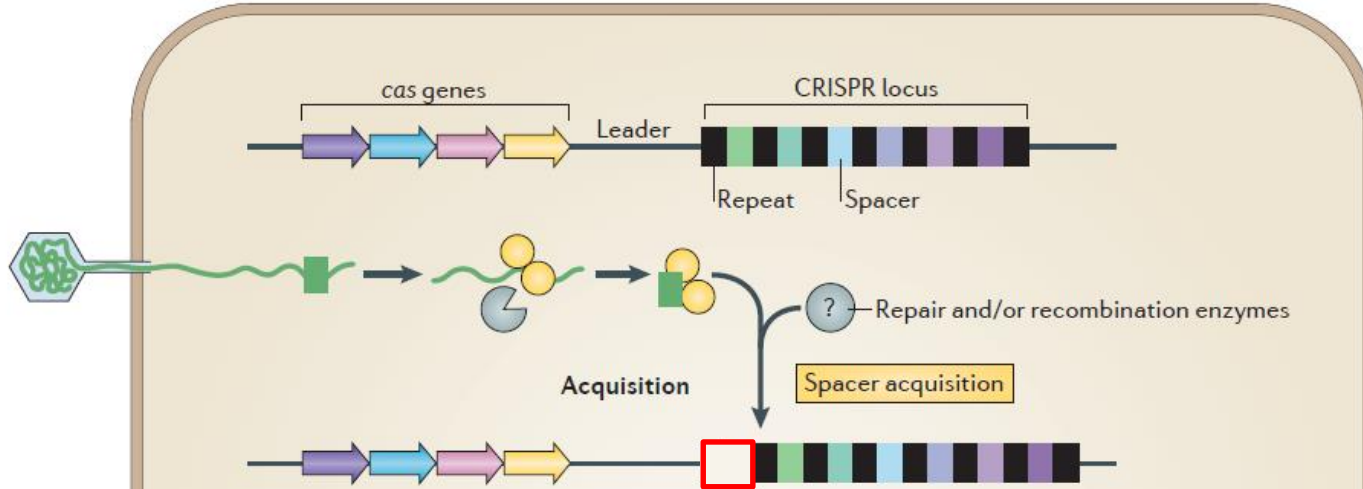
# CRISPR-Cas mechanism



a viral infection often leads to rapid synthesis of a new generation of viral particles and cell death

in a sub-fraction of a bacterial population, for instance due to infection by an inactive virus mutant, immunity can be gained through CRISPR-Cas

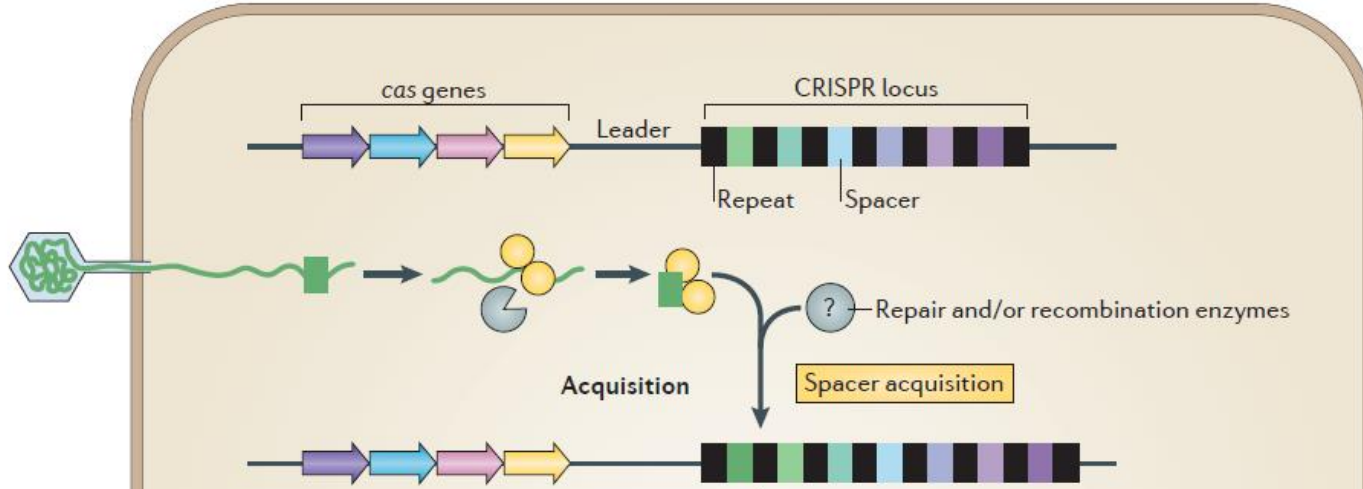
# CRISPR-Cas mechanism – step 1



Spacer acquisition

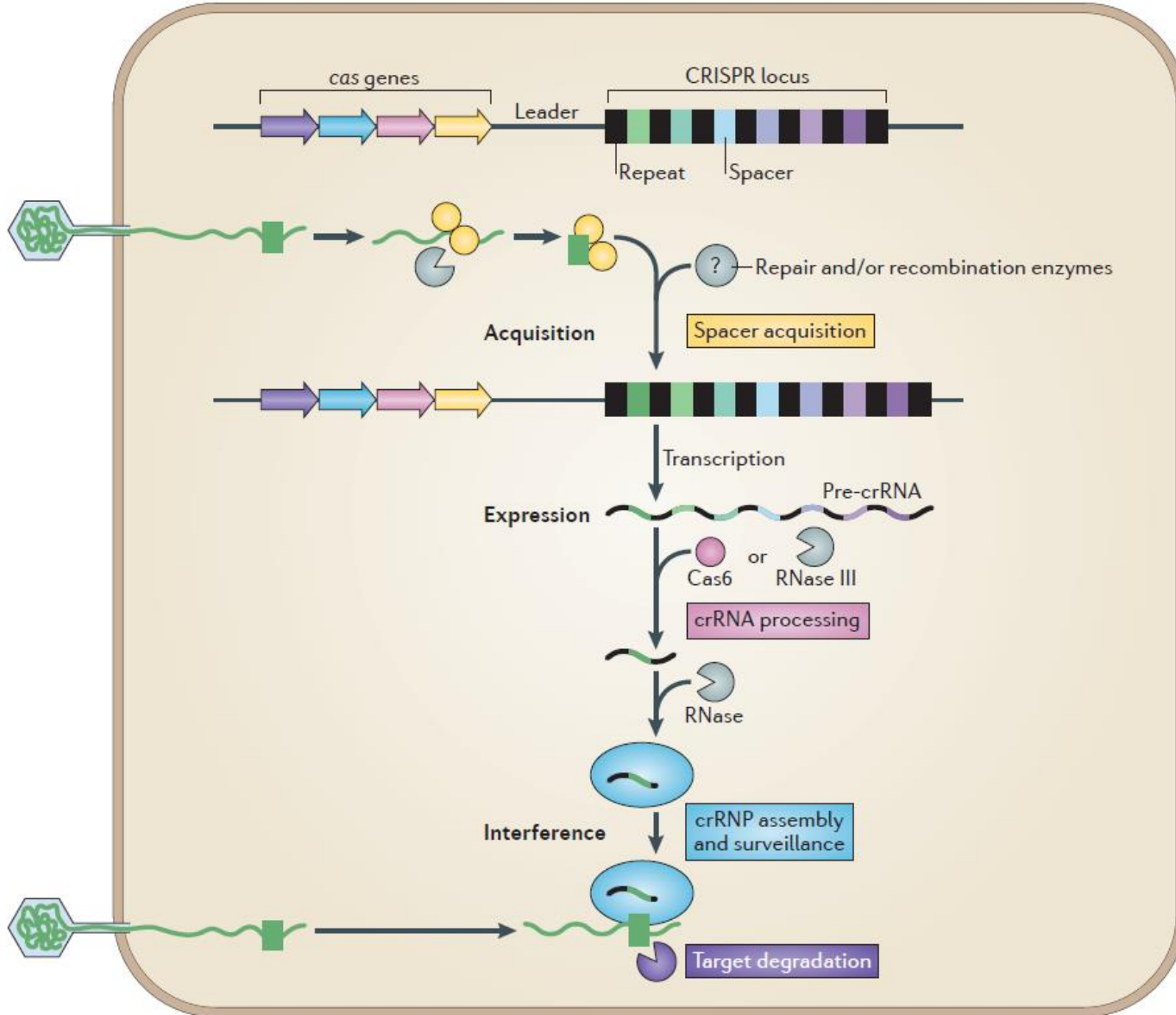
CRISPR array = memory of adaptive immune system

# CRISPR-Cas mechanism – steps 2 & 3



Spacer acquisition

# CRISPR-Cas mechanism – steps 2 & 3

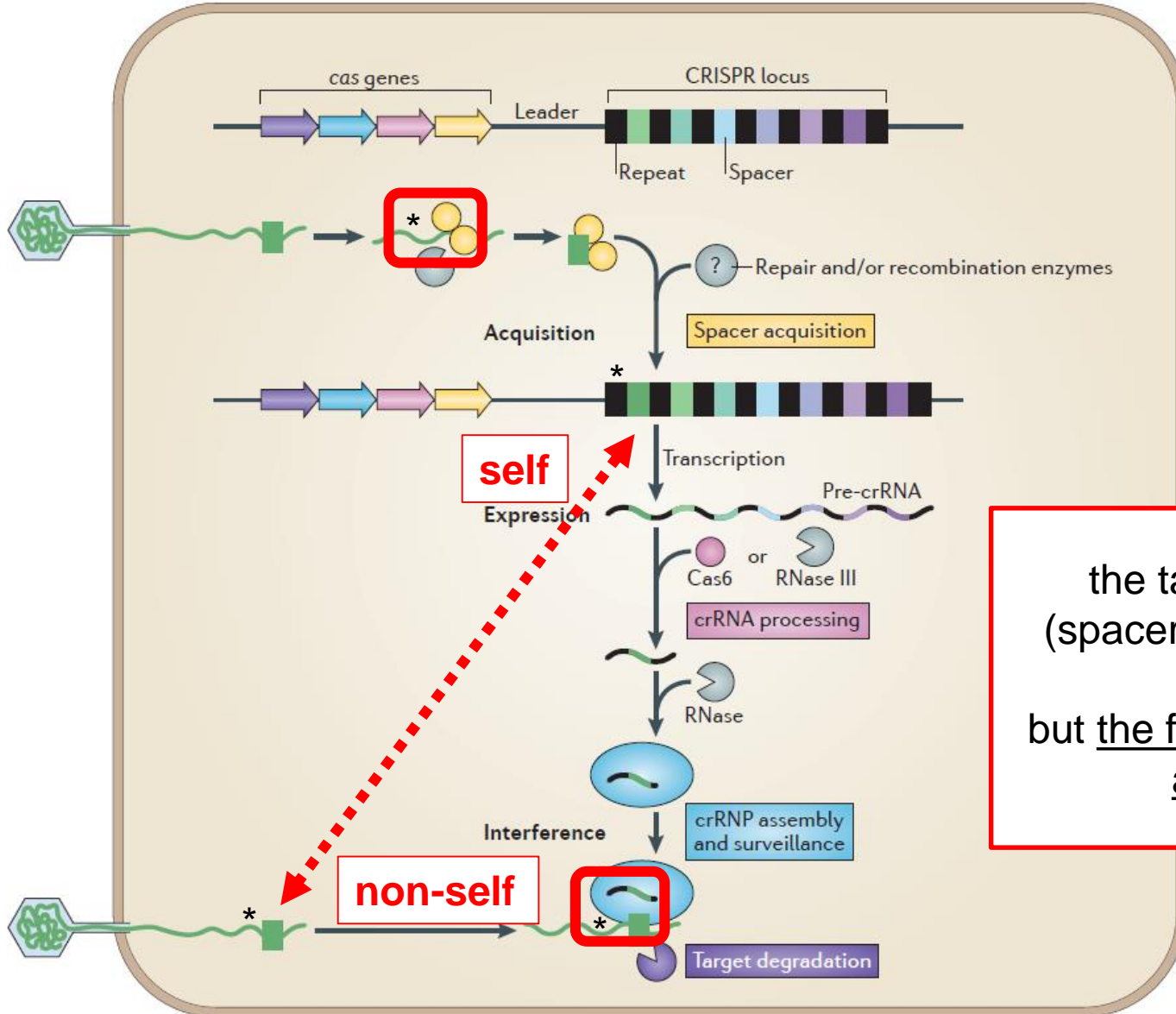


Spacer acquisition

Guide expression

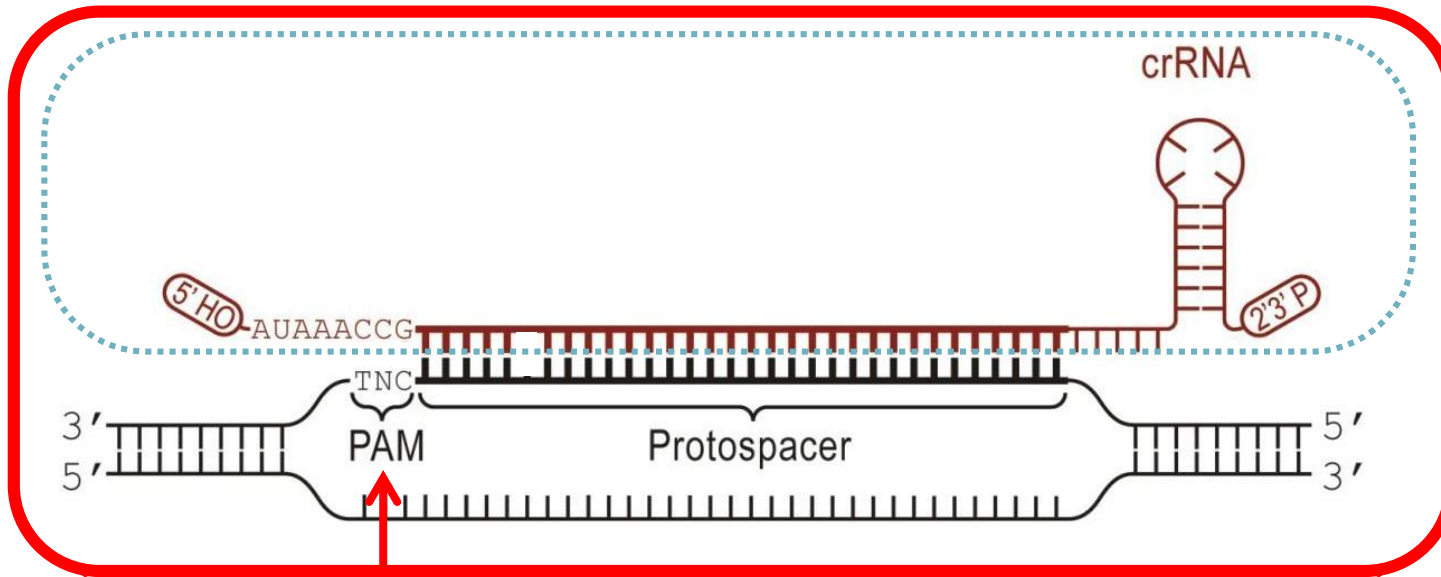
Target interference

# CRISPR-Cas – self / non-self discrimination

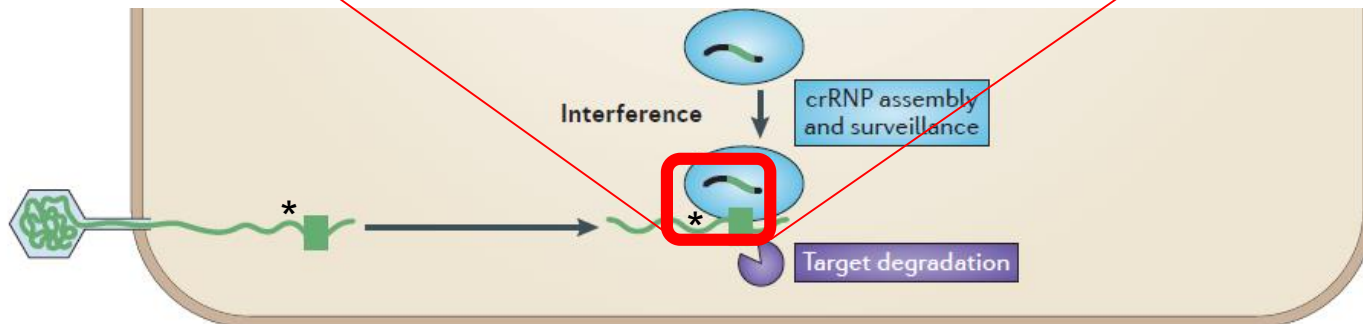


the target sequences (spacers & proto-spacers) are similar, but the flanking sequences\* are different

# CRISPR-Cas – self / non-self discrimination

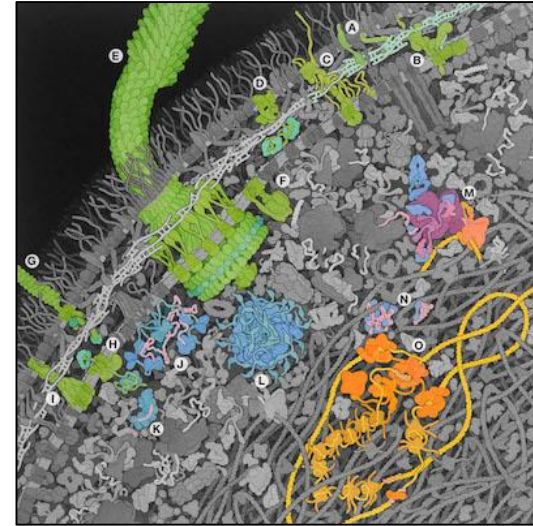
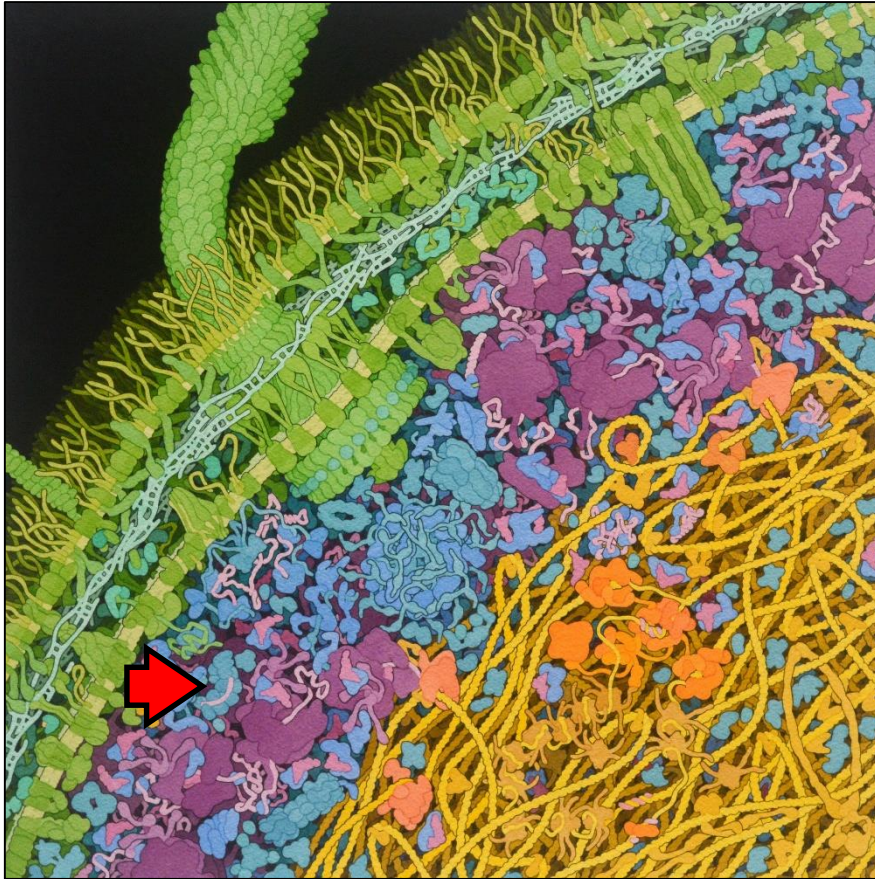


Protospacer Adjacent Motif (PAM)





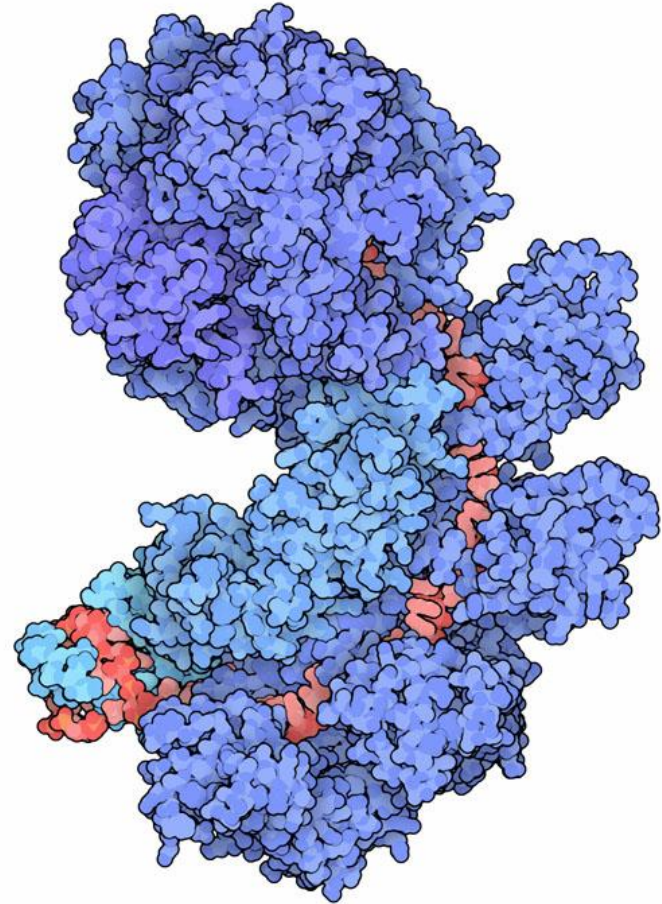
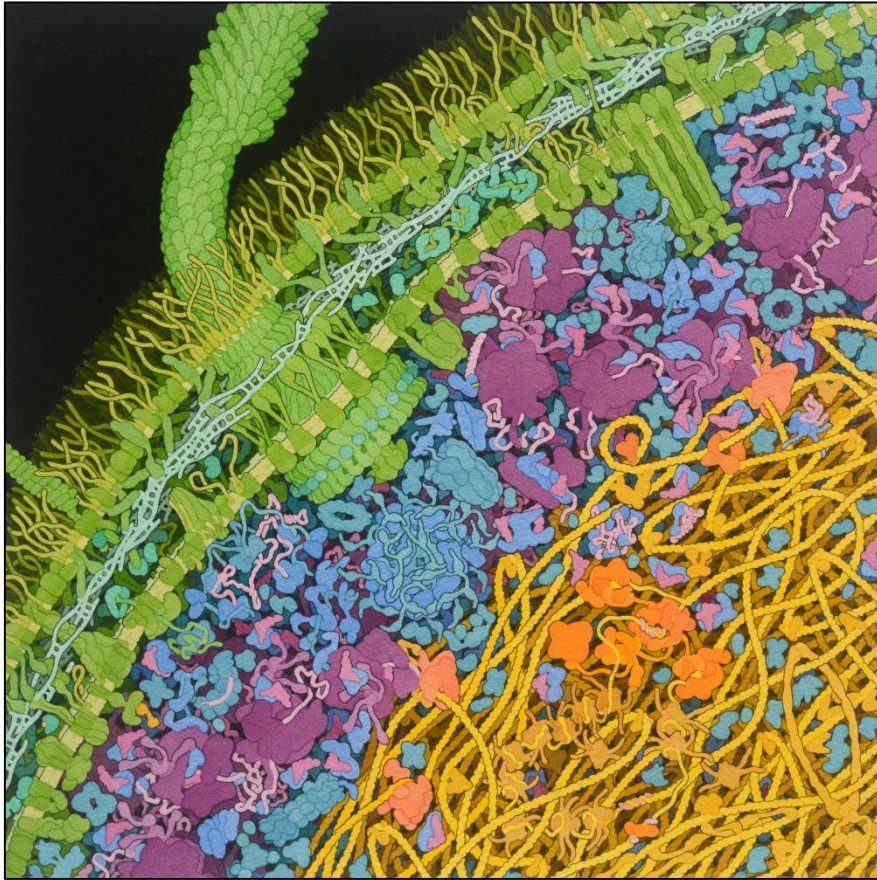
# Cascade in *E. coli* cell



- A. Lipoprotein, the most abundant protein in the cell, and OmpA. Two structural proteins that connect the outer membrane and the peptidoglycan network.
- B. A complex of proteins that synthesize the peptidoglycan network, including penicillin-binding proteins that are the target of penicillin.
- C. The Lpt system that transports lipopolysaccharides from the inner membrane (where they are synthesized) to the outer membrane.
- D. The Bam system that chaperones the folding of outer membrane proteins.
- E. Flagellum and flagellar motor.
- F. ATP synthase.
- G. Fimbria and fimbrial usher. One subunit is being transported to the usher by a chaperone.
- H. Secretory translocon, being assisted by chaperones.
- I. AcrAB/TolC multi-drug efflux pump.
- J. Degradosome, a complex of proteins for degrading obsolete RNA.
- K. Cascade and CRISPR.
- L. Pyruvate dehydrogenase complex.
- M. Expressome. The flexible arms on the ribosome are L12 subunits, which help coordinate the binding of tRNA and elongation factors.
- N. Two sRNA with their associated proteins.
- O. Replisome including several DNA polymerases and single-strand DNA-binding proteins protecting the single-stranded intermediates formed during the process of DNA replication.



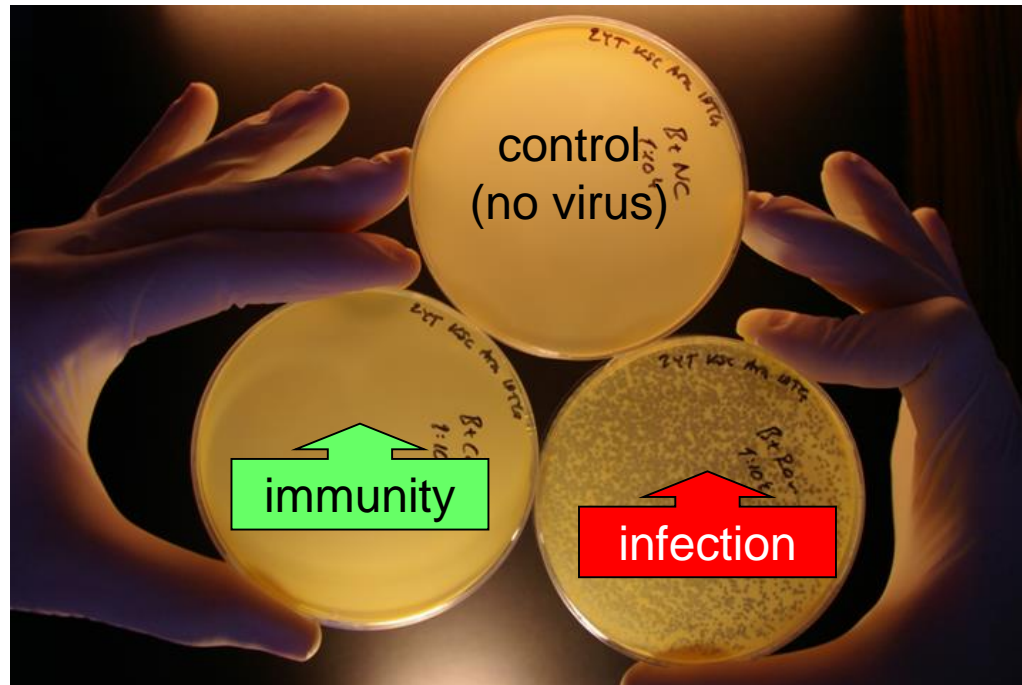
# Cascade in *E. coli* cell





# crRNA-guided DNA interference

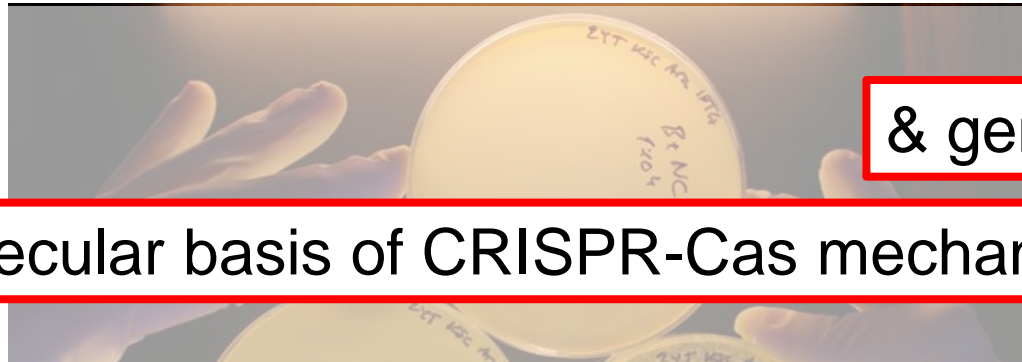
*E. coli* BL21  
design  
CRISPR-Cas





# crRNA-guided DNA interference

*E. coli* BL21  
design  
CRISPR-Cas



& genome editing

## Molecular basis of CRISPR-Cas mechanism

- Specific maturation of CRISPR-derived RNA guides
- Binding of crRNA guides by Cas effector complex
- Anti-viral defense by crRNA-guided DNA interference
- Transplantation of CRISPR-Cas system to another host
- CRISPR design allows for specific (multiplex) DNA targeting

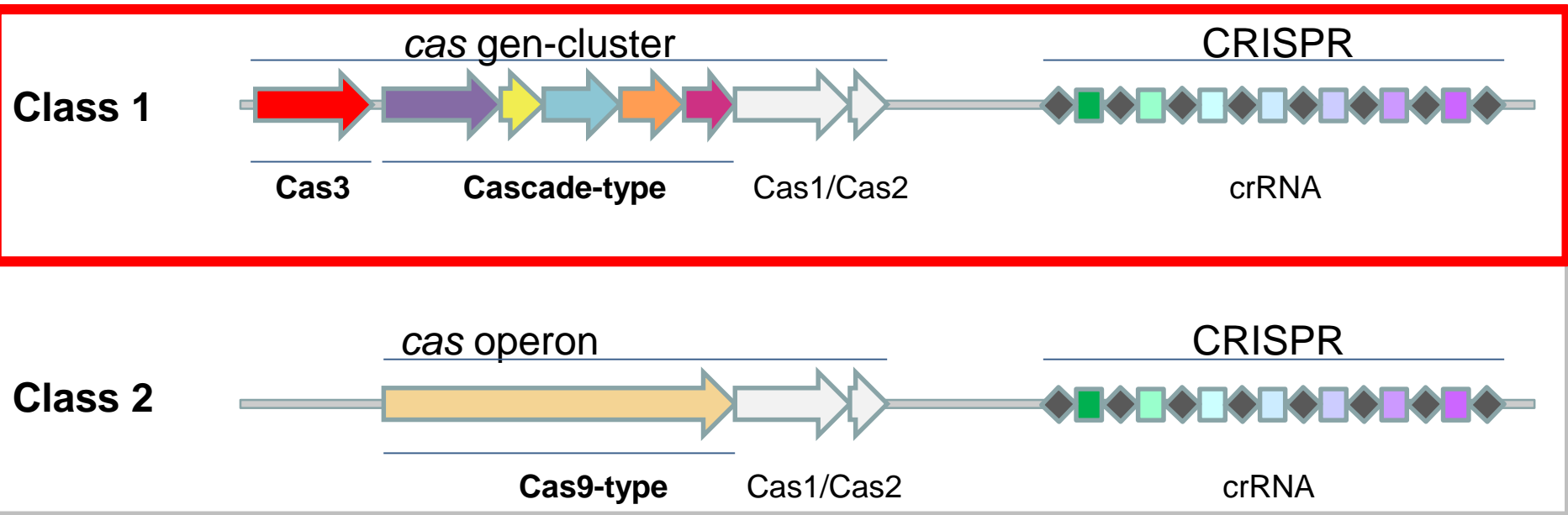
# outline

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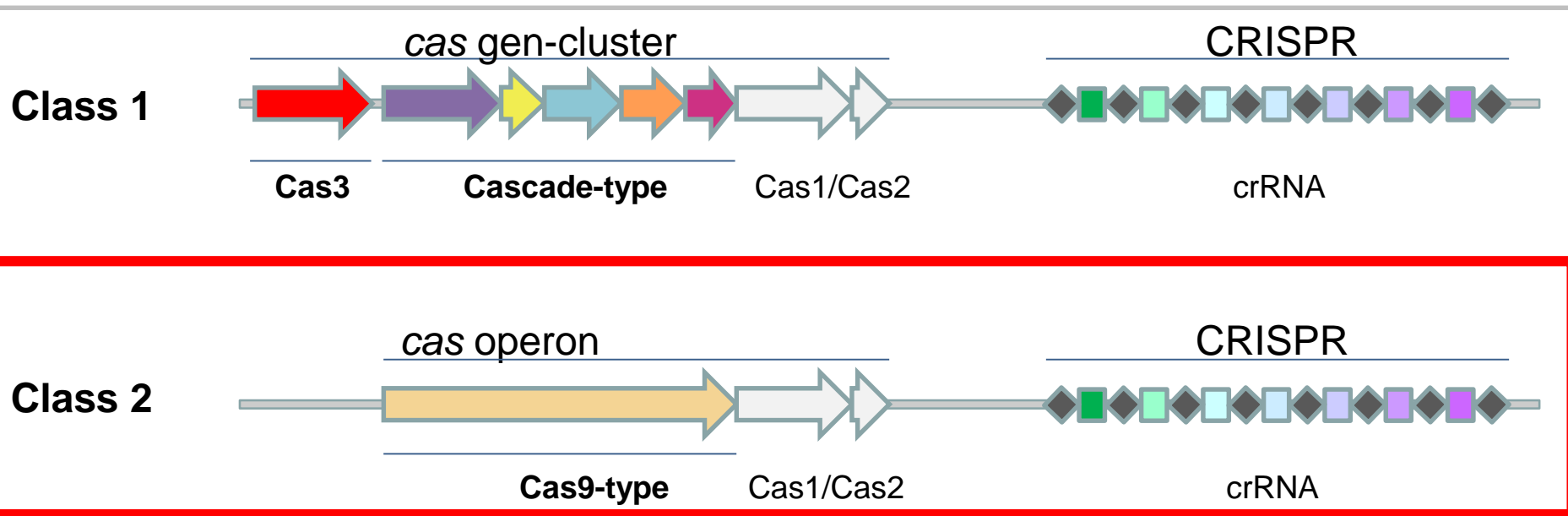


- History - *discovery & diversity*
- Biology - *anti-virus defence mechanism*
- ➔ Applications - *genome editing & gene therapy*

# CRISPR-Cas – 2 classes



# CRISPR-Cas – 2 classes

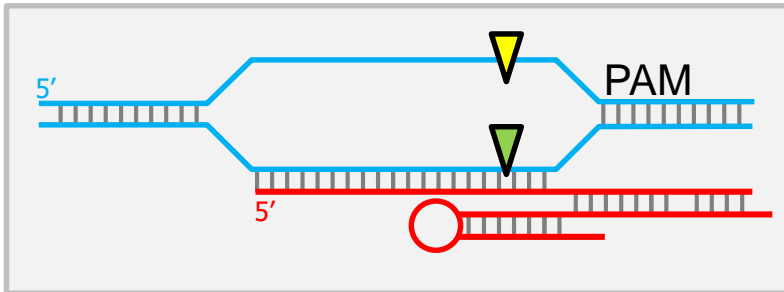




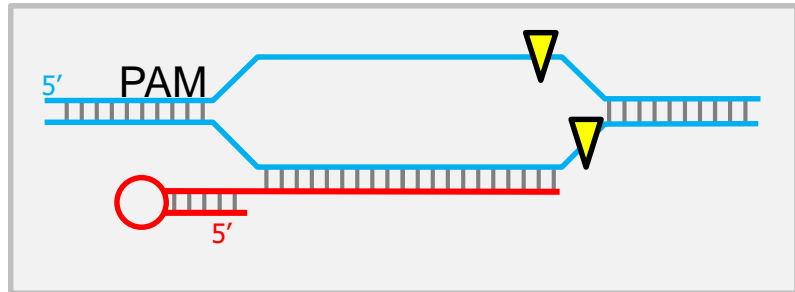
# Class 2 – Cas9



CRISPR-Cas9



CRISPR-Cas12

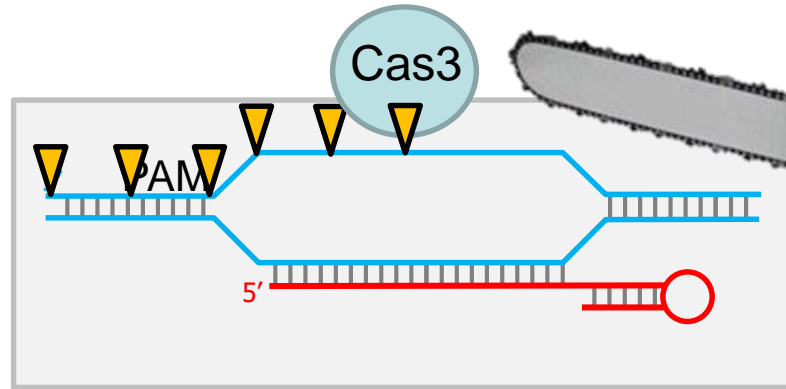


class 2 Cas nucleases are multi-functional

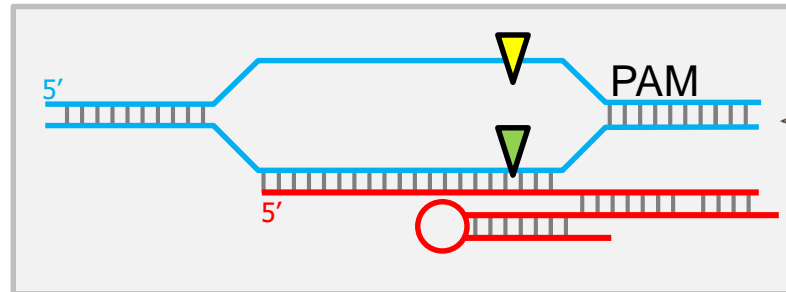
# Cascade - Cas9 - Cas12



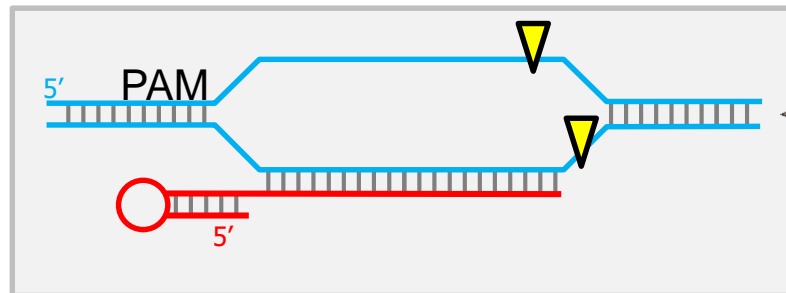
Type I  
Cascade



Type II  
Cas9



Type V  
Cas12

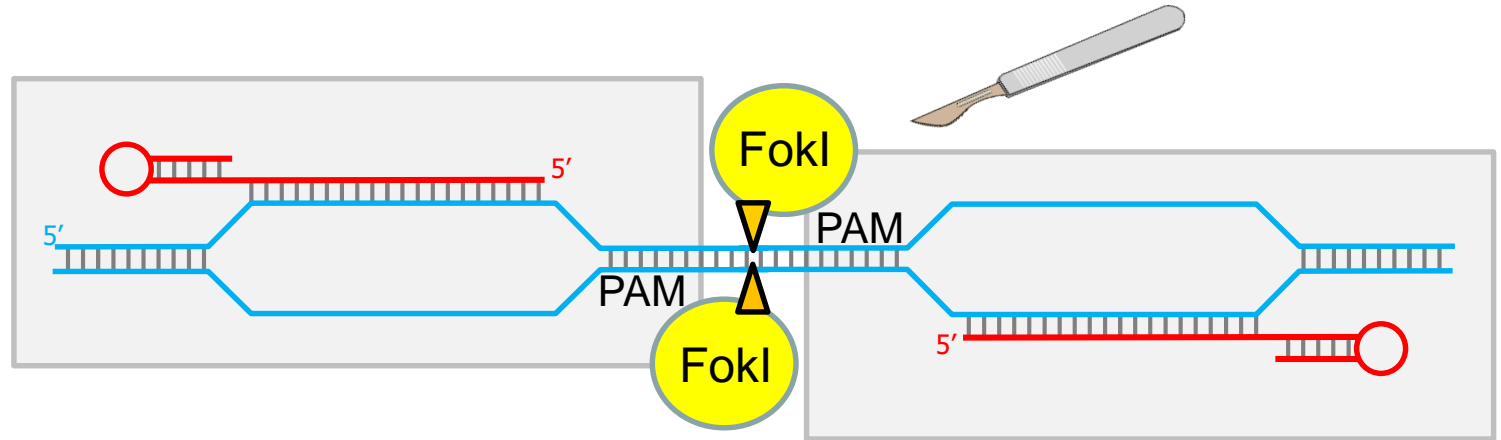




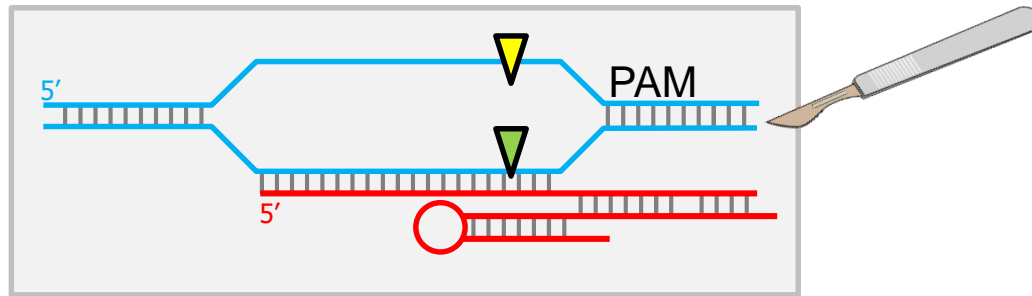
# Cascade - Cas9 - Cas12



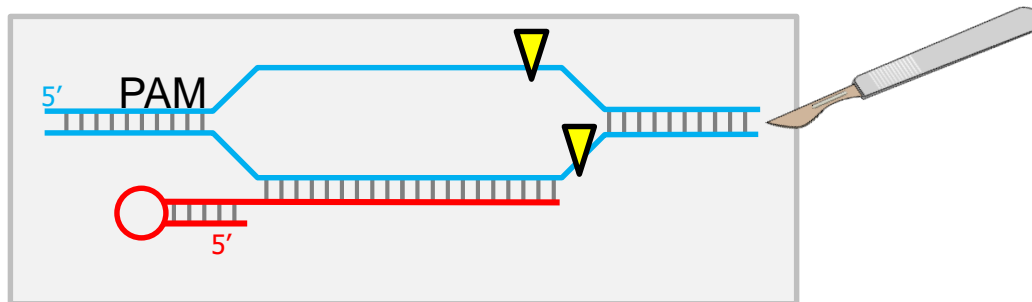
Type I  
Cascade



Type II  
Cas9



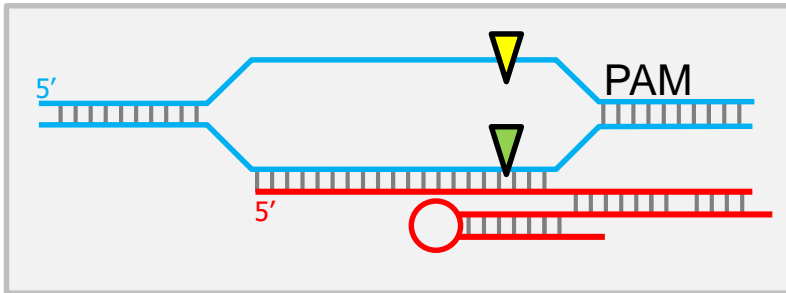
Type V  
Cas12



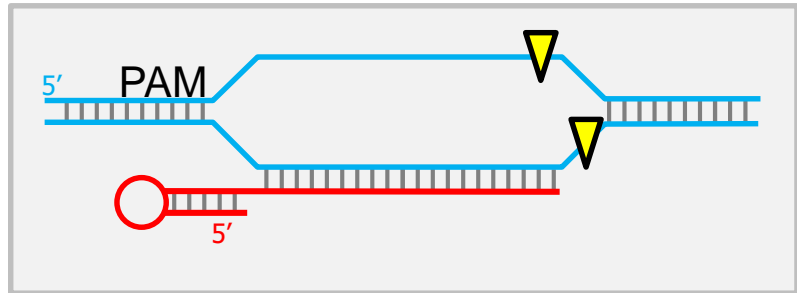
# CRISPR-Cas – DNA editing



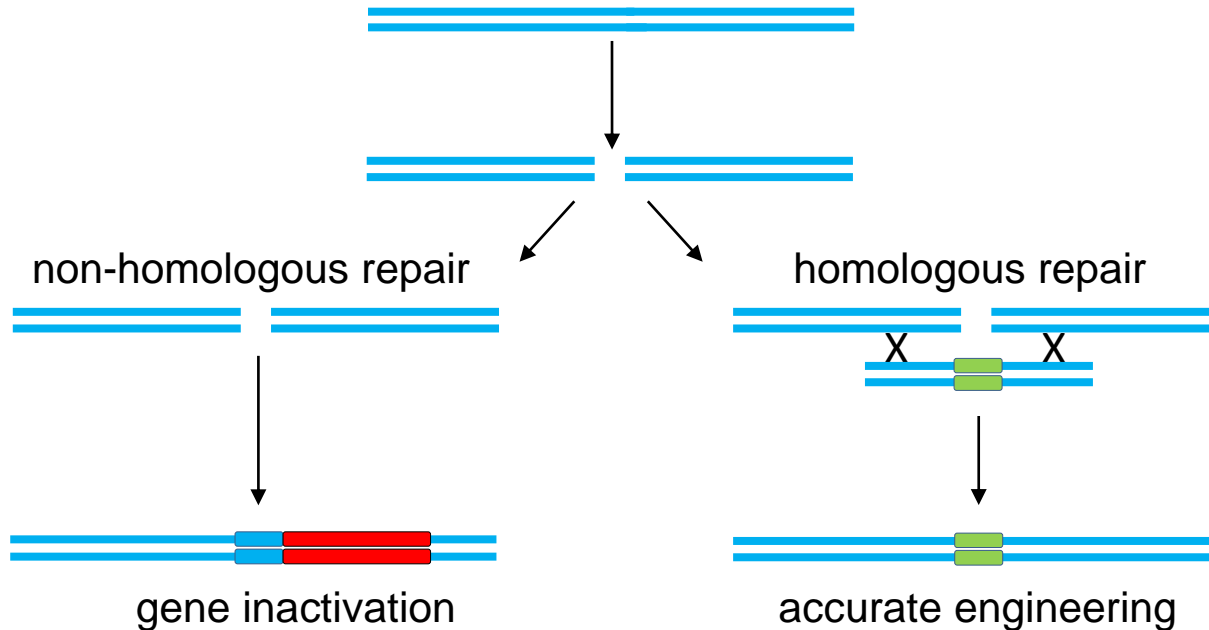
CRISPR-Cas9



CRISPR-Cas12



specific cleavage by Cas9 / Cas12



# Conclusions

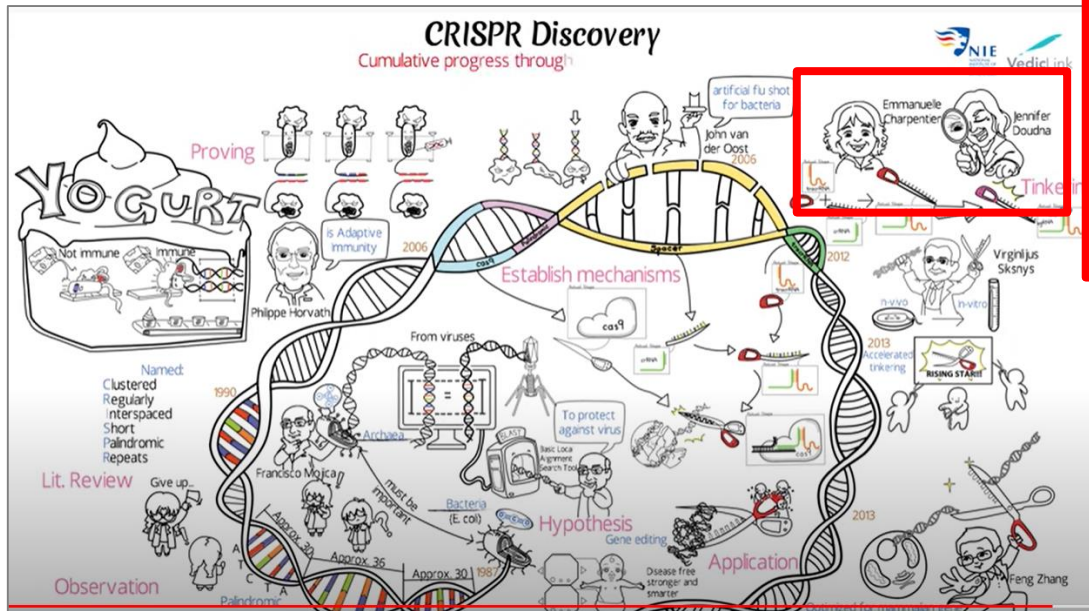


- CRISPR-Cas is an adaptive defense system in bacteria and archaea
- CRISPR RNAs guide nucleases to complementary **DNA\*** (or RNA) target sequences
- CRISPR-Cas systems are highly diverse (2 classes, 6 types, >20 subtypes)
- Cas nucleases (**Cas9, Cas12a, Cascade-FokI**)\* led to genome editing revolution
- Genome engineering in **Biotechnology**: bacteria, yeast, algae & plants
- **Clinical trials** of natural & synthetic Cas nucleases to cure human genetic diseases

# Conclusions



- CRISPR discoveries – youtube : <https://www.youtube.com/watch?v=RKh2mi3tsmc>



Scientific Background on the Nobel Prize in Chemistry 2020

A TOOL FOR GENOME EDITING

# Collaborations



## ***Wageningen***

Wen Wu  
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Jurre Steens  
Thomas Swartjes  
Raymond Staals  
Rob Joosten  
Richard v. Kranenburg  
Sjoerd Creutzburg\*  
Ioannis Mouggiakos\*  
Prarthana Mohanraju\*  
Yifan Zhu\*  
Daan Swarts\*  
Matthijs Jore\*  
Magnus Lundgren\*  
Edze Westra\*  
Stan Brouns\*



## ***Boston***

Feng Zhang

## ***Berkeley***

Jennifer Doudna

## ***Bethesda***

Eugene Koonin

## ***Bozeman***

Blake Wiedenheft

## ***Rotterdam***

Joyce Lebbink

## ***Utrecht***

Niels Geijsen



Netherlands Organisation for Scientific Research

