



UNIVERSITÀ
DEGLI STUDI DI TRIESTE



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Widespread RNA-based *cas* regulation monitors crRNA abundance and anti-CRISPR proteins

Liu, Chao et al. Cell Host & Microbe, 2023, Volume 31, Issue 9, 1481 - 1493.e6

Presented by Letizia Barbieri and Nora Costella



OUTLINE

- **INTRODUCTION**
- STUDY AIM
- RESULTS
- CONCLUSION

Prokaryotes VS Mobile Genetic Elements

Coevolution between **prokaryotes** and their **predators** has led to sophisticated **defence** mechanisms

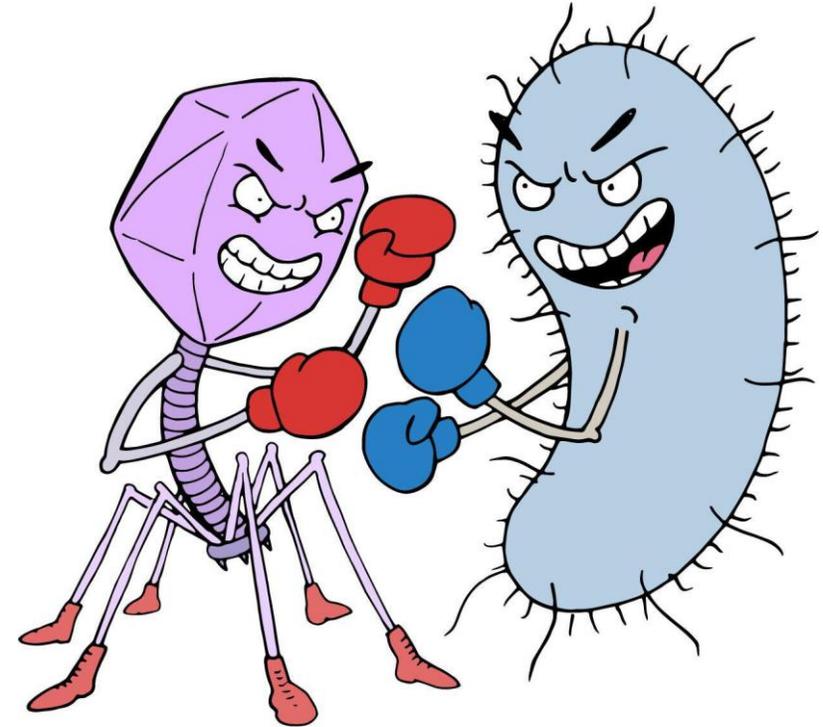
Prokaryotes employ:

Innate defence

- Preventing phage adsorption
- Restriction of invading DNA
- Abortive infection

Adaptive defence

- **CRISPR-Cas system**



Stores **memory** of the past infections and, upon reinfection, uses **RNA-guided nucleases** to silence phage or other mobile genetic elements (MGEs)

CRISPR-Cas main actors

- **CRISPR array** - archive of previous infections

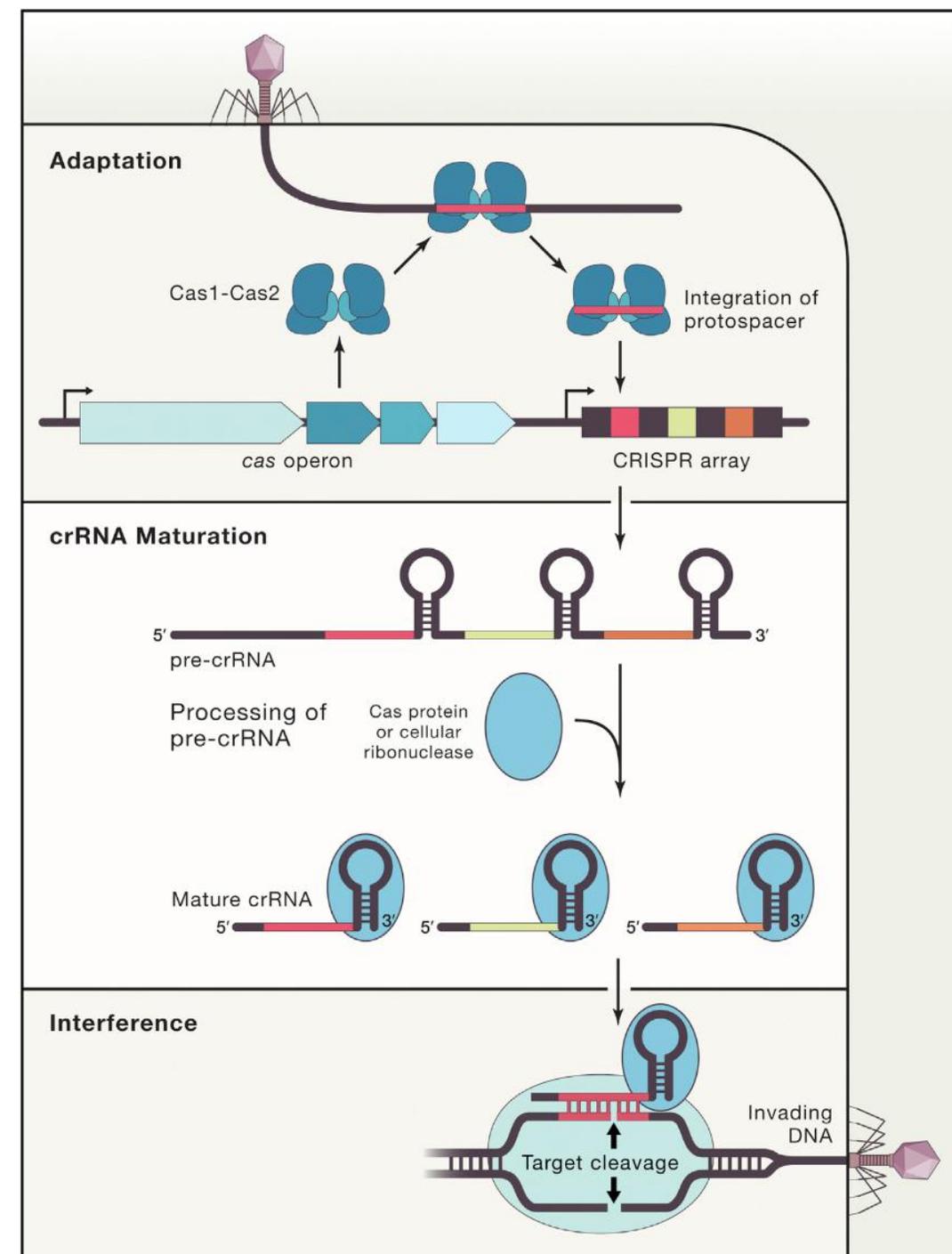
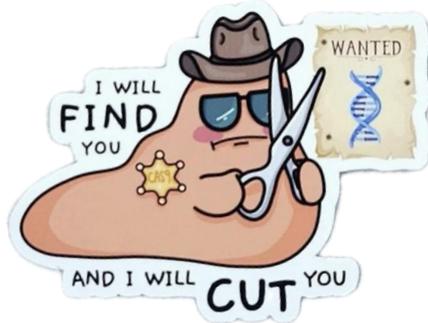
Genomic locus:

- **Repeats:** identical sequences recognized by Cas
- **Spacers:** derived from integration of **protospacers** (= DNA fragments from past infections)

Transcribed as **pre-crRNA** → processed to **crRNA**

- **Cas** - nuclease proteins with active role in:

1. **Adaptation** (= spacer acquisition)
2. **crRNA biogenesis**
3. **Interference** → Cas use crRNA as guide to degrade foreign DNA



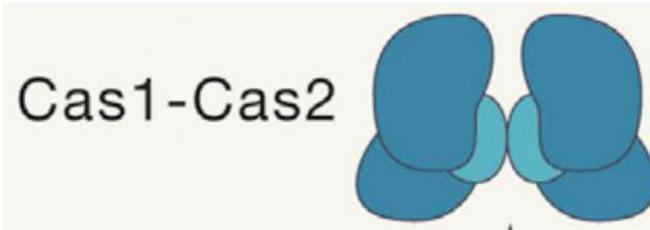
CRISPR-Cas systems Classification

	Class 1 multi-protein Cas complexes for interference			Class 2 Single Cas for interference		
Type	Type I	Type III	Type IV	Type II	Type V	Type VI
Effector	Cascade complex + Cas3 ✂	Csm (III-A) or Cmr (III-B) + Cas10 ✂	?	Cas9 (HNH and RuvC domains) ✂	Cas12 (only RuvC domain) ✂	Cas13 (HEPN domain) ✂
Guide	crRNA	crRNA	?	crRNA + tracrRNA	crRNA	crRNA
Target recognition	through PAM	5' repeat portion of crRNA unbound	?	through PAM	T-rich PAM	RNAs complementarity to crRNA
Cutting	ssDNA break intermediate first, then dsDNA break	DNA being transcribed and RNAs	?	blunt dsDNA break	dsDNA breaks (with overhangs) distal from PAM	ssRNA + collateral host ssRNAs

1. Adaptation – memory creation

Incorporation of protospacers from invading nucleic acids into the host's CRISPR array, generating spacers

Machinery (for almost all CRISPR-Cas systems in *E.Coli*)

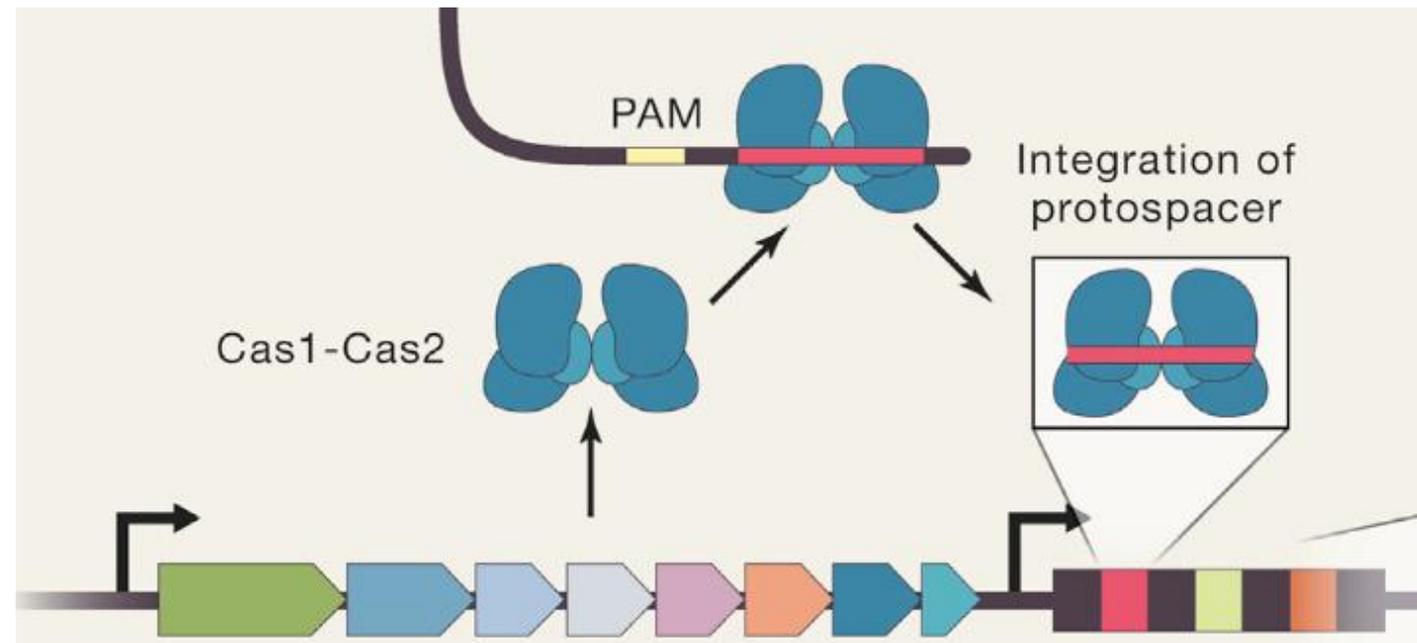


- Catalytic integrase
- Structural role

Adaptation steps:

- **Detection** of foreign genetic elements
- **Selection & processing** of protospacers for integration – e.g. type I & II recognize, and then remove, protospacers with PAM complementary to a Cas1 subunit
- **Integration** of spacers - preferentially at the leader end of the CRISPR array

Challenge: integrate **ONLY** invader genome to avoid lethal autoimmunity



2. crRNA biogenesis - memory retrieval

CRISPR array is transcribed in **long pre-crRNA**, his maturation leads to **crRNA** having:

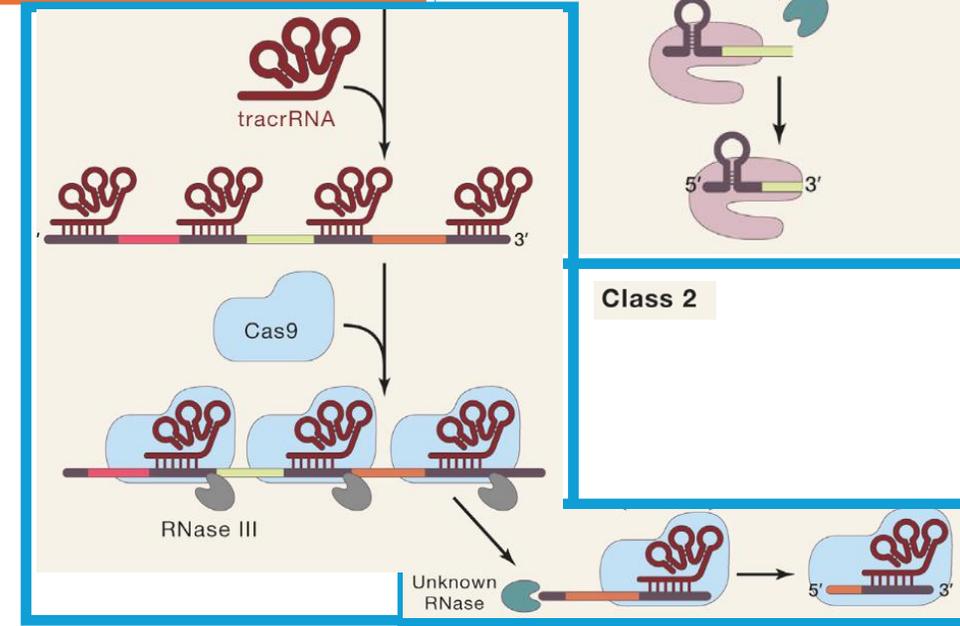
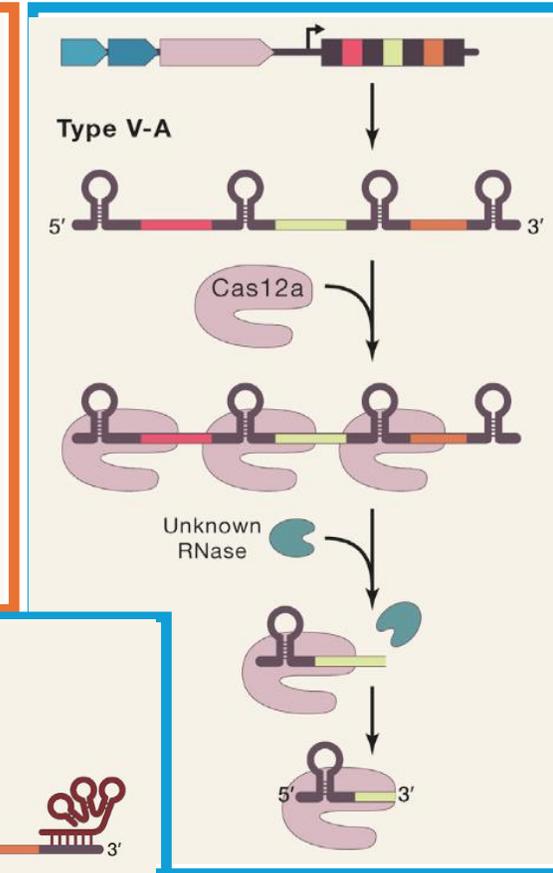
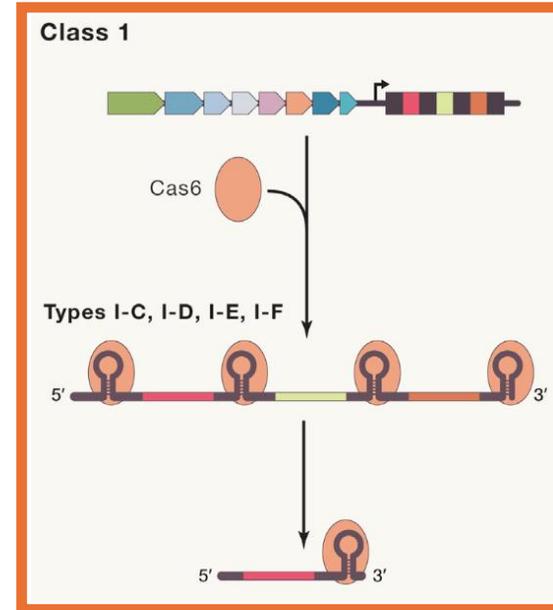
- **Repeats** → for Cas recognition
- **Spacer** → for target binding

→ Class 1 crRNA maturation

Cas6 recognizes and cuts hairpin formed by palindromic sequences in repeats portion

→ Class 2 crRNA maturation

- **Type II**: needs duplex **pre-crRNA:tracrRNA** and **RNase II**
- **Type V & VI**: effector proteins are sufficient



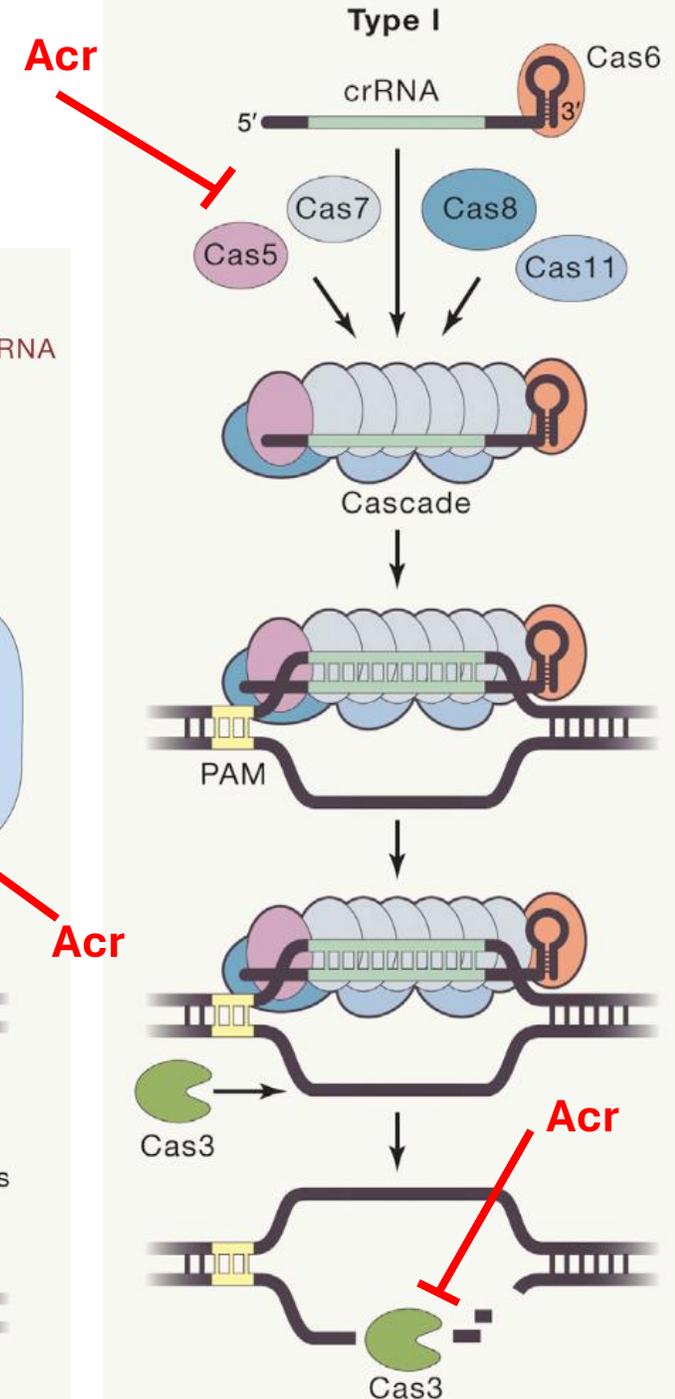
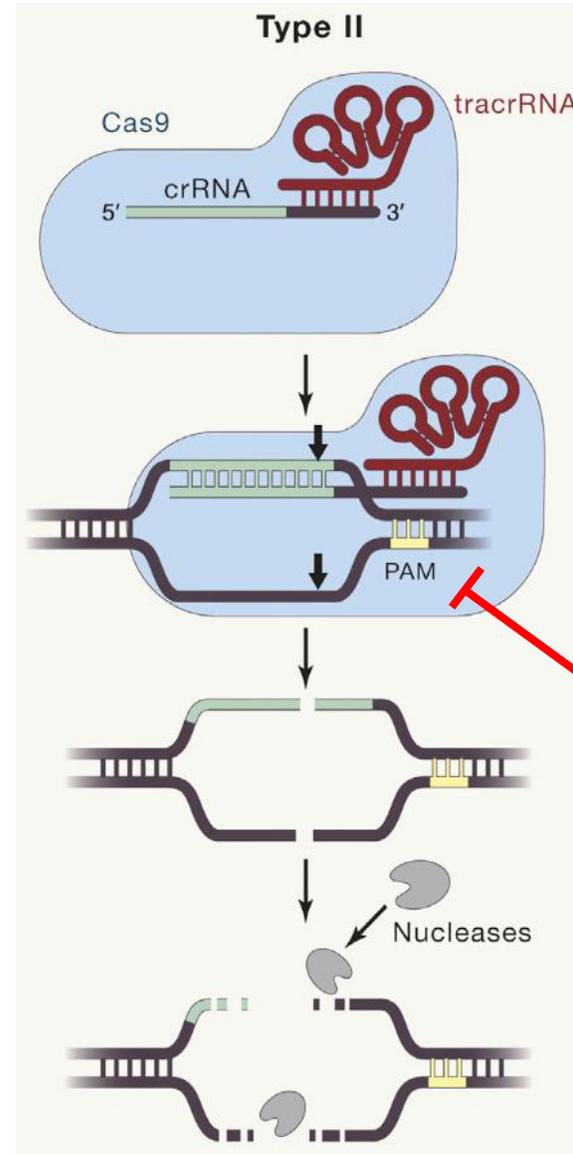
3. Interference - defence activation

crRNA guides sequence-specifically the effector machinery to cleave invading nucleic acids

- **Crucial challenge** → prevent autoimmunity: cut only invading DNA and not CRISPR array
- **Solution** → effectors recognize PAM, which are ABSENT in CRISPR array

Phages defend themselves from CRISPR-Cas system

Acr (anti-CRISPR) proteins are a strategy employed by some phages to circumvent CRISPR-Cas immunity



CRISPR-Cas comes with **fitness costs**

Limited adaptation

Acquisition of **beneficial MGE**-encoded traits is restricted by CRISPR-Cas systems

Autoimmunity

Adaptation machinery **can't distinguish self- vs non-self DNA**

Metabolic burden

Maintaining these complex systems is costly

How to **balance** these costs with the benefits of CRISPR-Cas systems?

Regulatory networks

for **context-dependent induction** of CRISPR-Cas immunity

Effective antiviral defence



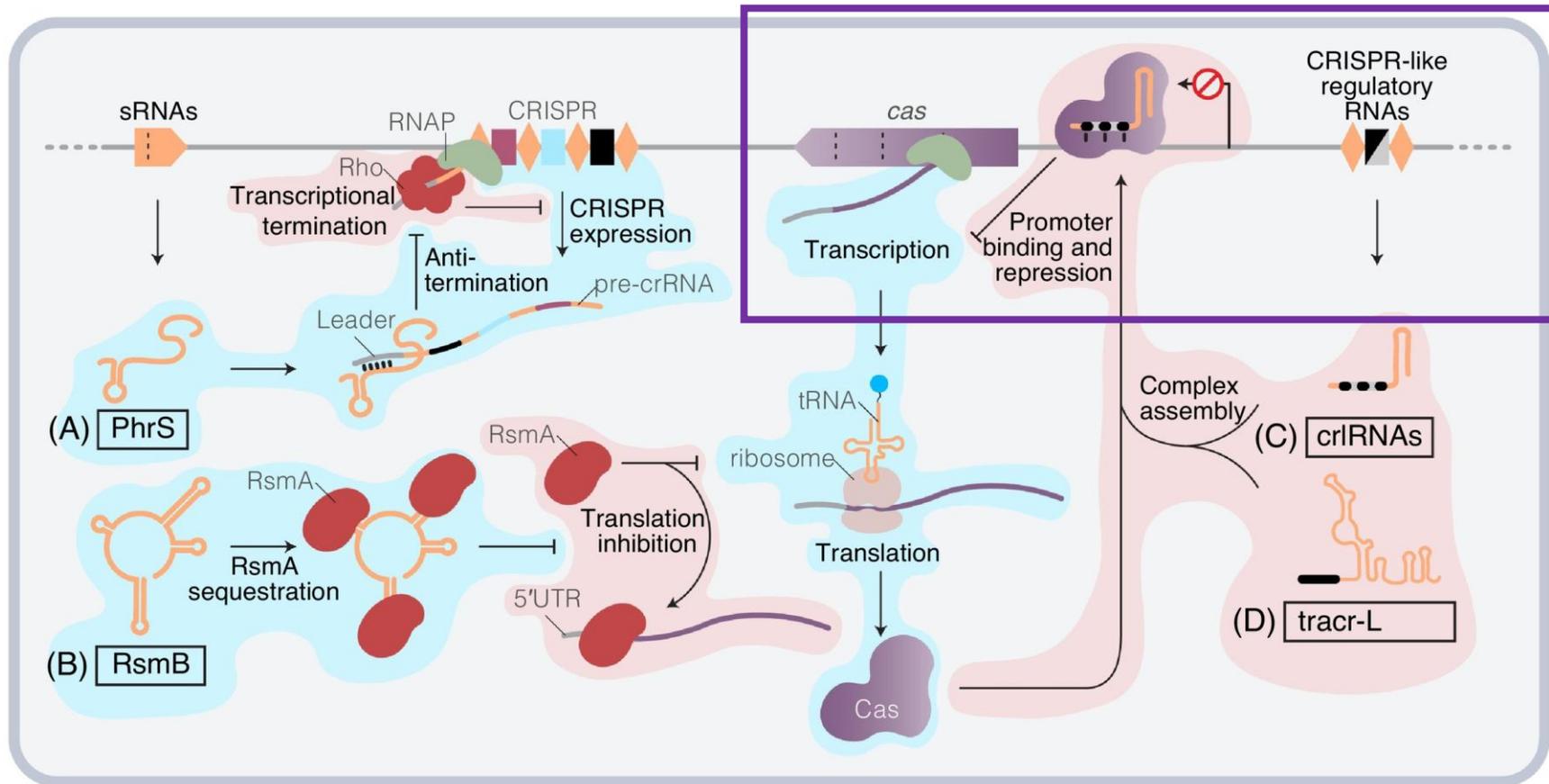
Avoid autoimmunity



Careful regulation of immunity is a universal principle of cellular life!

→ ncRNA-based regulatory networks of CRISPR-Cas immunity

Diverse families of **ncRNA regulate crRNA or Cas production** acting on the transcriptional or post-transcriptional level

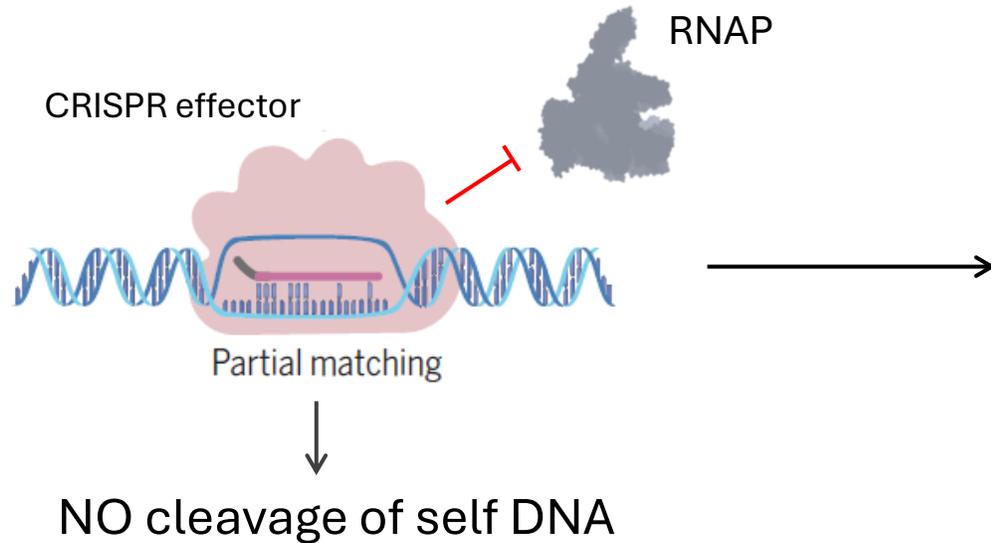


Some regulatory ncRNA act as non-canonical guides that **guide Cas to *cas* promoter** for transcriptional **autoregulation**

CRISPR effector is not just an immune effector but also a **gene regulator!**

→ CRISPR effectors as gene regulators

Transcriptional regulation is based on **PARTIAL or LIMITED spacer-protospacer complementarity**



Repression of target promoter
(protospacer)

Genes subjected to this kind of regulation:

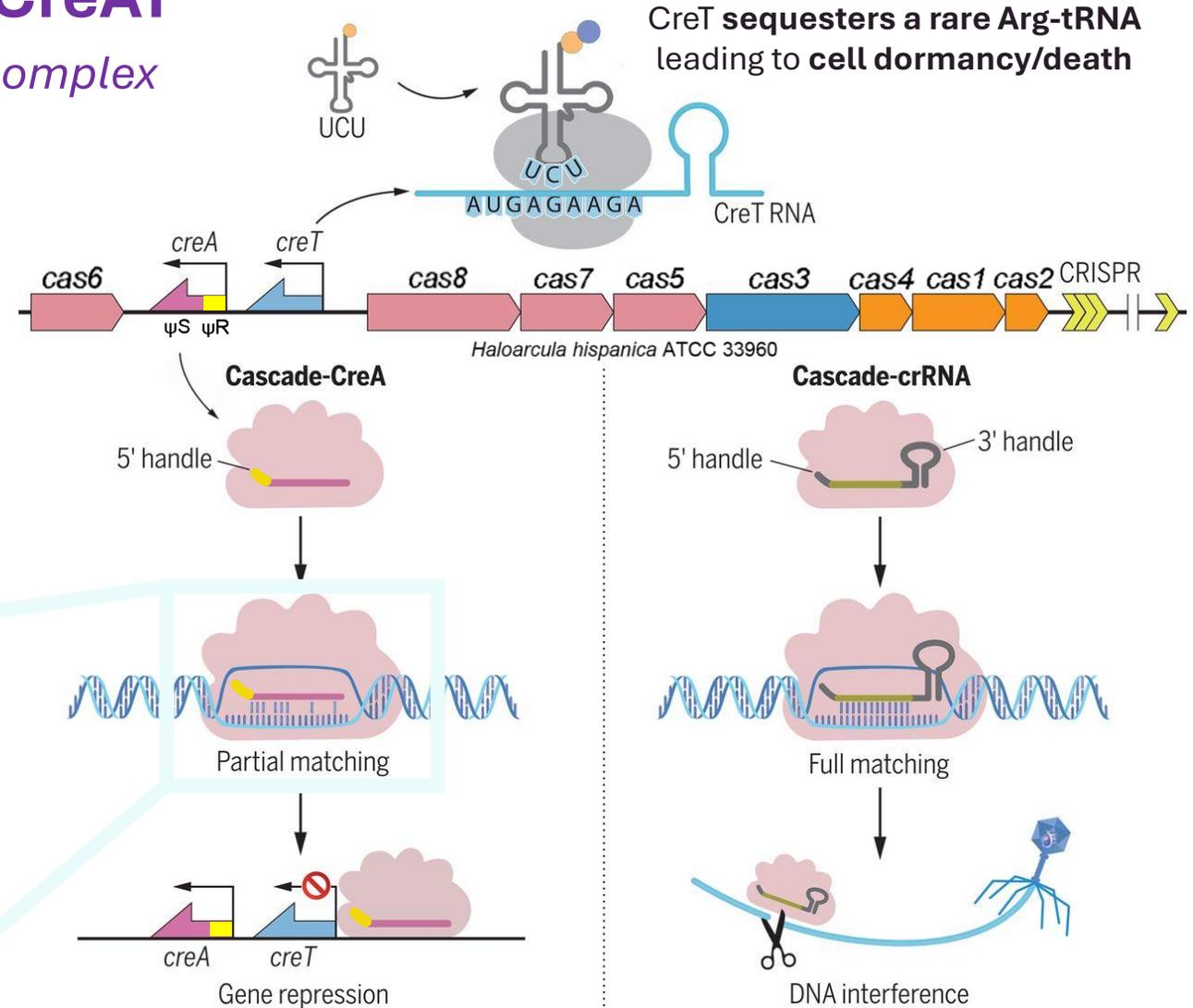
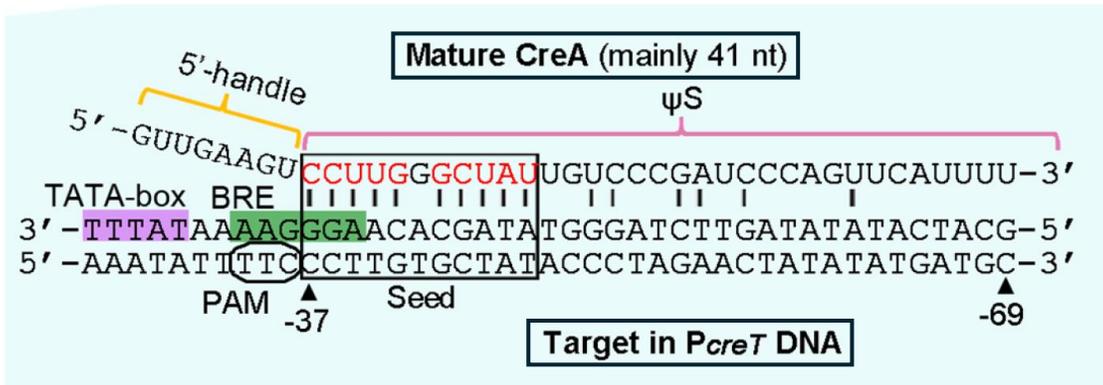
- *cas* genes (Cas autorepression)
- Virulence-related genes
- **Type VIII Toxin-Antitoxin** system
- ...

Type VIII Toxin-Antitoxin system CreAT

The regulatory role of *type I-B CRISPR* effector complex

CreA (CRISPR repeat-like antitoxin):

- crRNA-like RNA (**crRNA**) transcribed from a degenerated mini-CRISPR
- within **type I-B CRISPR-cas** loci of *Haloarcula hispanica*



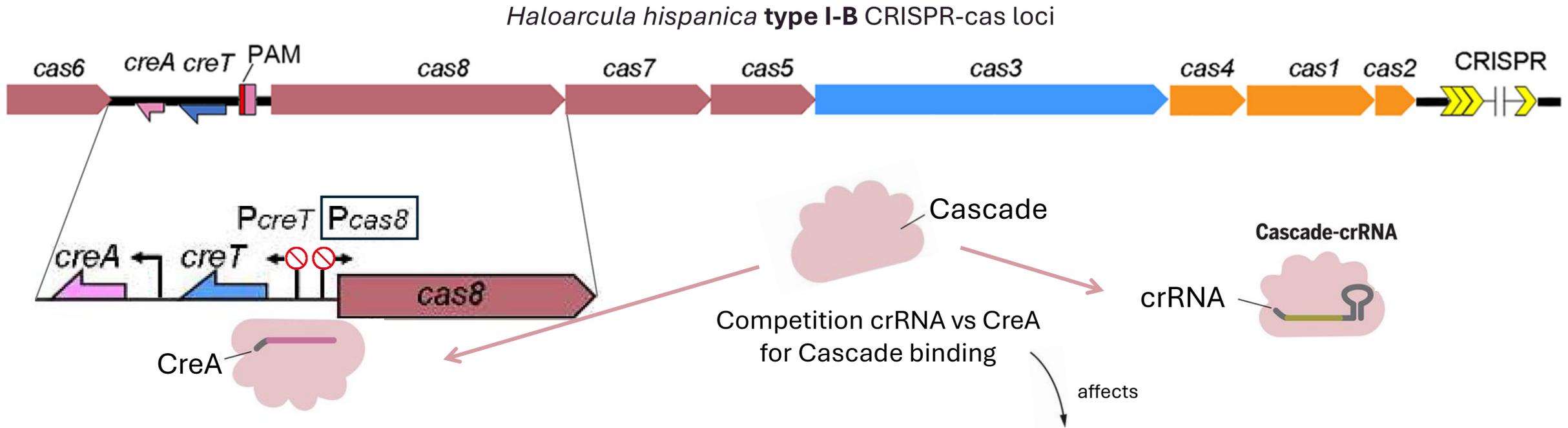
CreA antitoxin reprograms **Cascade** to transcriptionally repress the RNA **CreT** toxin

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In this study:

SPOILER ALERT



CreA target site includes a previously overlooked ***cas8* promoter**

CreA-Cascade synchronously represses *creT* and *cas*

A new **regulatory network** for **context-dependent *cas* expression** to **avoid autoimmunity**

OUTLINE

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Article

Widespread RNA-based *cas* regulation monitors crRNA abundance and anti-CRISPR proteins

Chao Liu,^{1,3,7} Rui Wang,^{1,7} Jie Li,^{2,7} Feiyue Cheng,^{1,7} Xian Shu,^{1,3,7} Huiwei Zhao,¹ Qiong Xue,¹ Haiying Yu,² Aici Wu,^{1,3} Lingyun Wang,^{1,4} Sushu Hu,¹ Yihan Zhang,^{1,5} Jun Yang,^{1,6} Hua Xiang,^{2,3,*} and Ming Li^{1,3,8,*}

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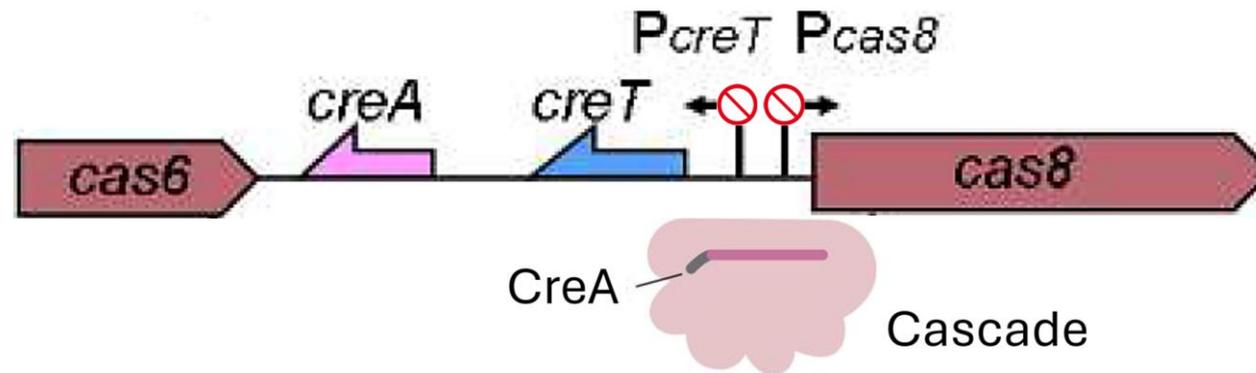
⁷These authors contributed equally

⁸Lead contact

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<https://doi.org/10.1016/j.chom.2023.08.005>

1. Demonstrating that *H. hispanica* **CreA** represses also ***cas*** transcription

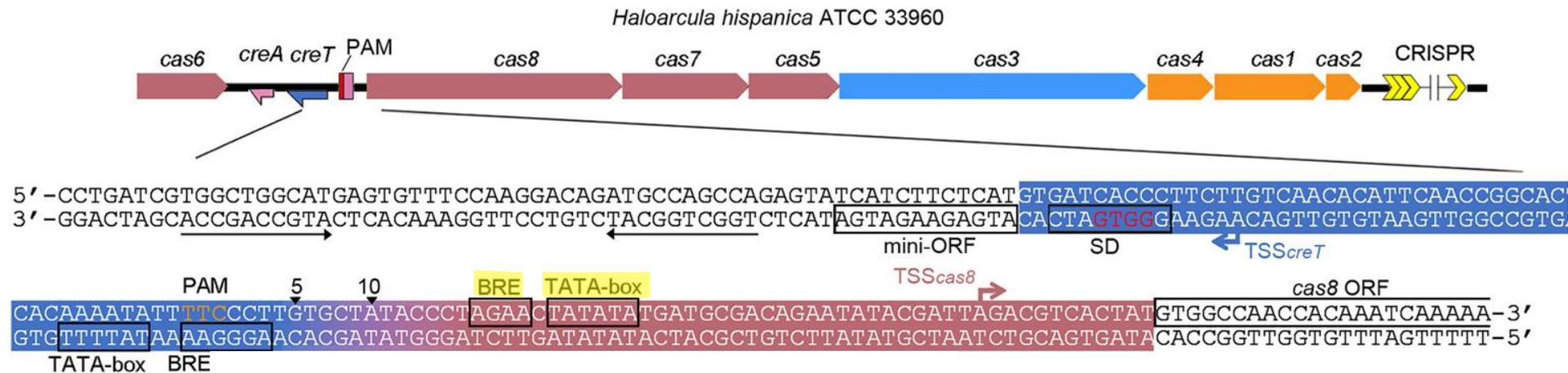


1. Demonstrating that *H. hispanica* CreA represses also cas transcription

Previous study: a *H. hispanica* ΔTA mutant showed **70-90% reduction of cas transcripts except for cas6** => An overlooked **promoter precedes cas8**



→ Reanalysis of previous RNA-seq data → **cas8 TSS identification** 12 bp upstream cas8 ORF
=> Prediction of archaeal promoter elements **BRE** and **TATA** box

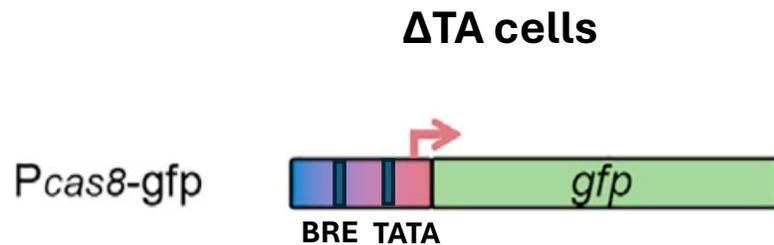


1. Demonstrating that *H. hispanica* CreA represses also *cas* transcription

Validation of *cas8* promoter

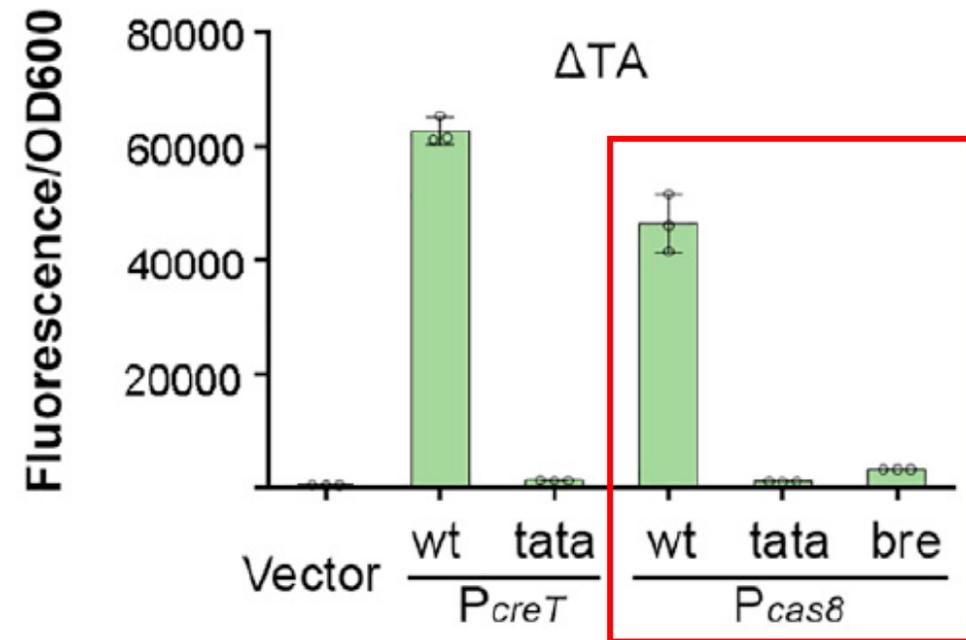
cas8 TSS identification allowed to predict archaeal promoter elements **BRE** and **TATA** box

→ **VALIDATION:**



GFP-reporter system to evaluate the activity of:

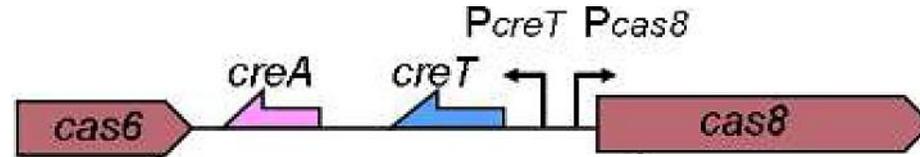
- **wt** P_{cas8}
- **TATA-mutated** P_{cas8}
- **BRE-mutated** P_{cas8}



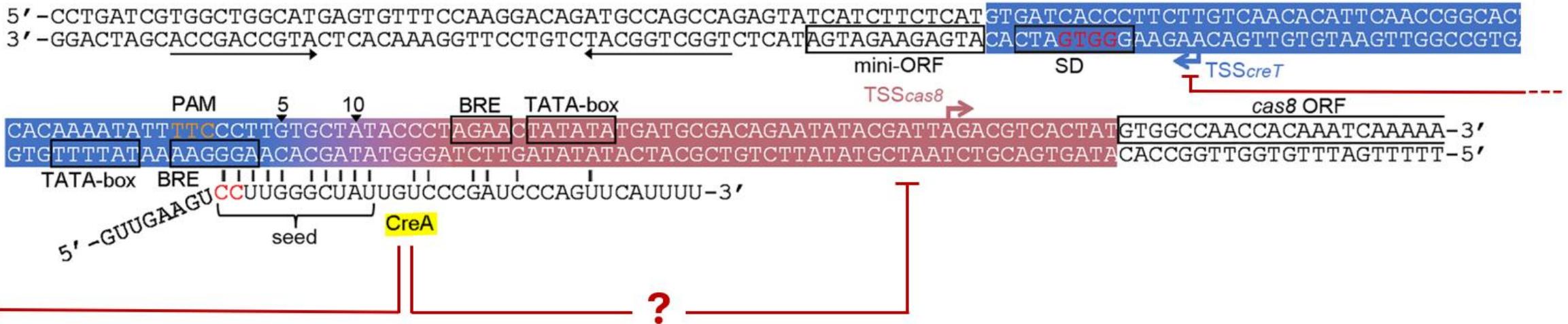
P_{cas8} activity **nullified** by mutating archaeal promoter elements sequences

1. Demonstrating that *H. hispanica* CreA represses also *cas* transcription

Validated *cas8* promoter



P_{cas8} and P_{creT} run divergently and tightly flank **CreA target site**



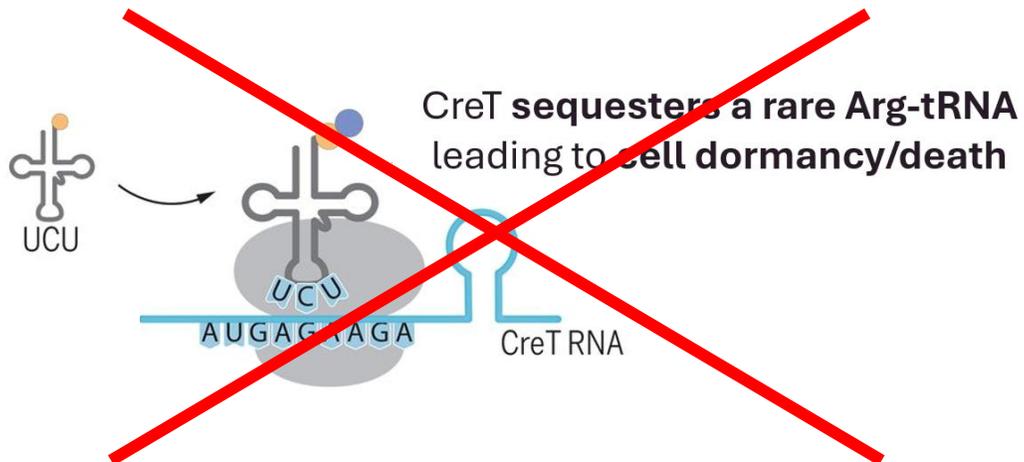
Is P_{cas8} also repressed by CreA?

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Is P_{cas8} also repressed by CreA?

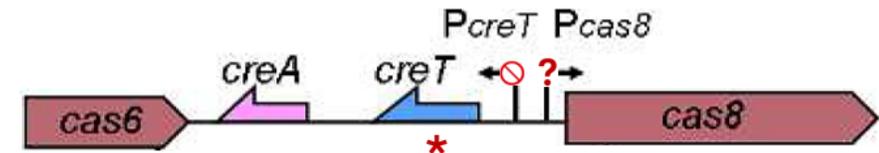
What happens if we **impair CreA binding** to its target site by **mutating CreA sequence**?

Problem: Abolishing CreA-mediated repression would lead to **CreT toxin expression!**



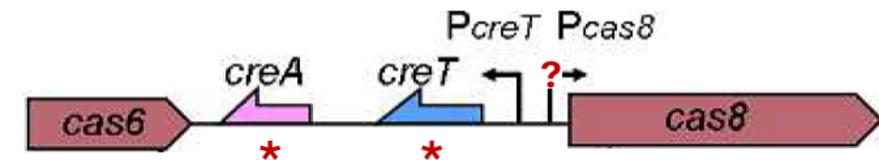
Solution: mutate **CreT** to make it **non-toxic**

Tm mutant cells - **CONTROL**



- **creT mutated** → **non-toxic** if expressed
- *creA* wt

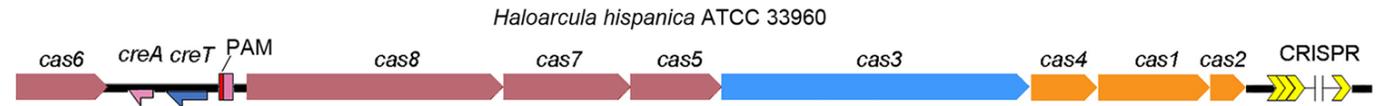
TA double mutant cells – **No CreA binding**



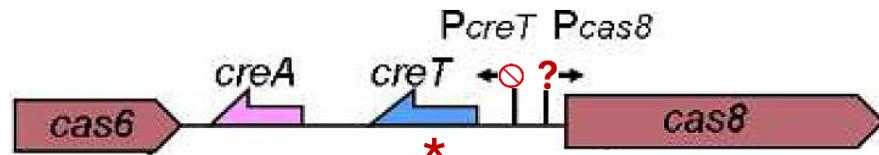
- **creT mutated** → **non-toxic** if expressed
- **creA mutated** → **impaired complementarity** to its target site => abolished repression

1. Demonstrating that *H. hispanica* CreA represses also *cas* transcription

Is P_{cas8} also repressed by CreA?

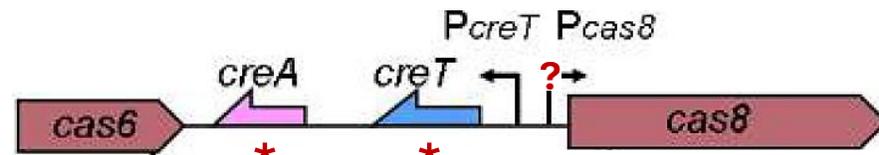


Tm mutant cells - CONTROL



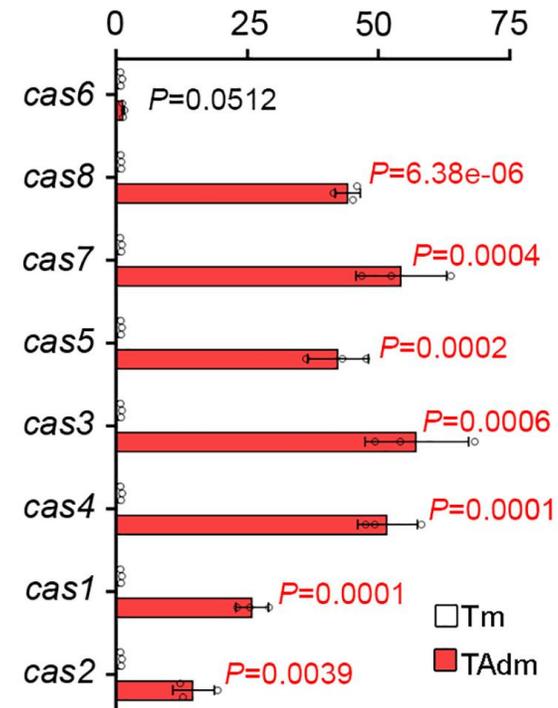
- **creT mutated** → **non-toxic** if expressed
- **creA wt**

TA double mutant cells – No CreA binding



- **creT mutated** → **non-toxic** if expressed
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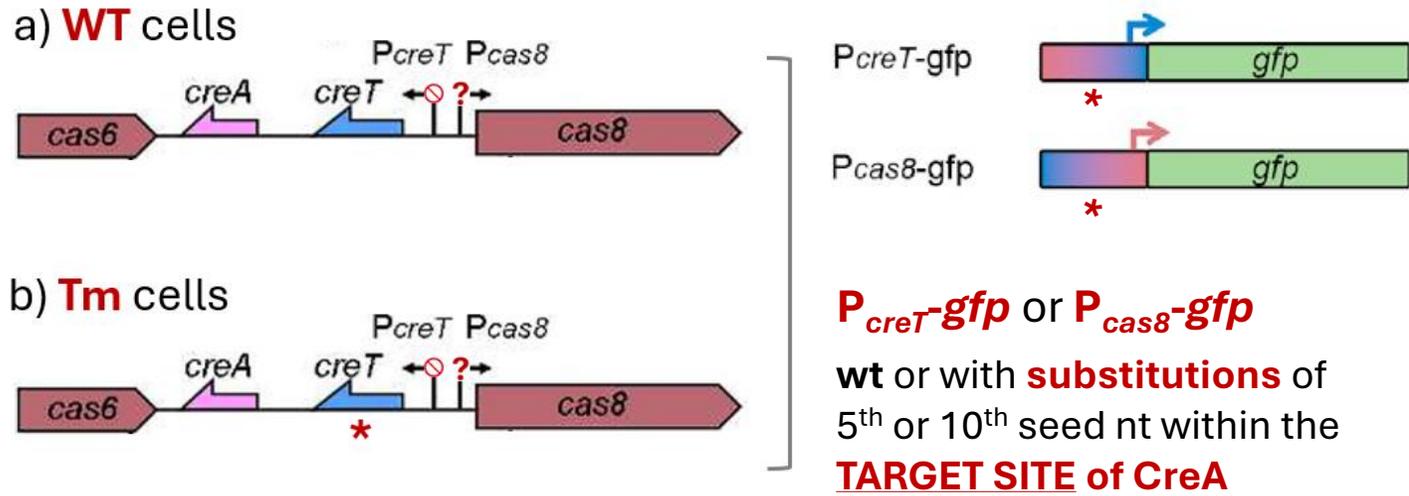
Relative RNA abundance
(Normalized $2^{-\Delta Ct}$)



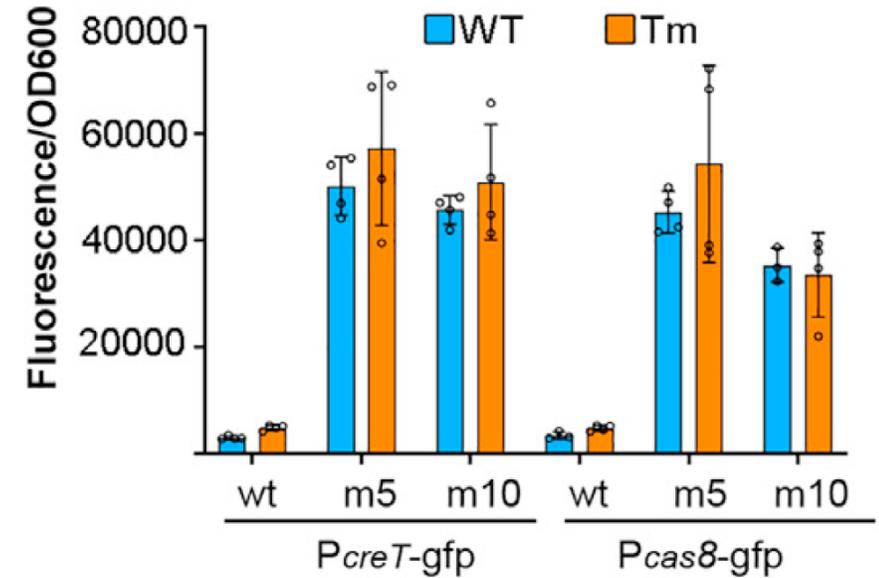
15- to 54-fold up-regulation of *cas8* and downstream *cas* in TAdm; no significant changes for *cas6*

1. Demonstrating that *H. hispanica* CreA represses also *cas* transcription

Is P_{cas8} also repressed by CreA?



GFP-reporter assay to evaluate P_{cas8} and P_{creT} activity depending on **CreA** capability to bind its target site



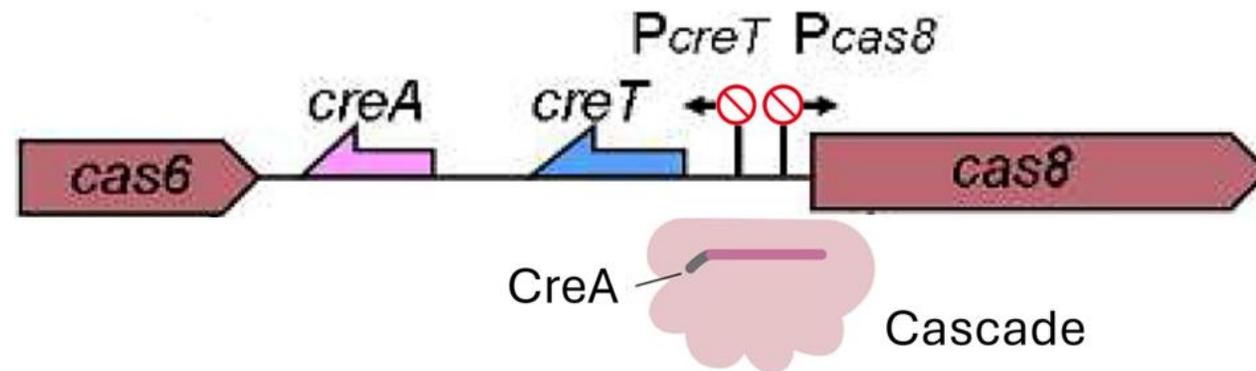
P_{cas8} and P_{creT} are **both de-repressed in m5 and m10** in WT and Tm cells when the **seed sequence is mutated**

1. Demonstrating that *H. hispanica* CreA represses also *cas* transcription

Is P_{cas8} also repressed by CreA?



Conclusion: P_{cas8} and P_{creT} are **synchronously downregulated by CreA**



2. CreA-guided cas regulation shifts from **reducing autoimmunity** to **enhancing immunity against imperfect targets**



2. CreA-guided *cas* repression reduces autoimmune risk

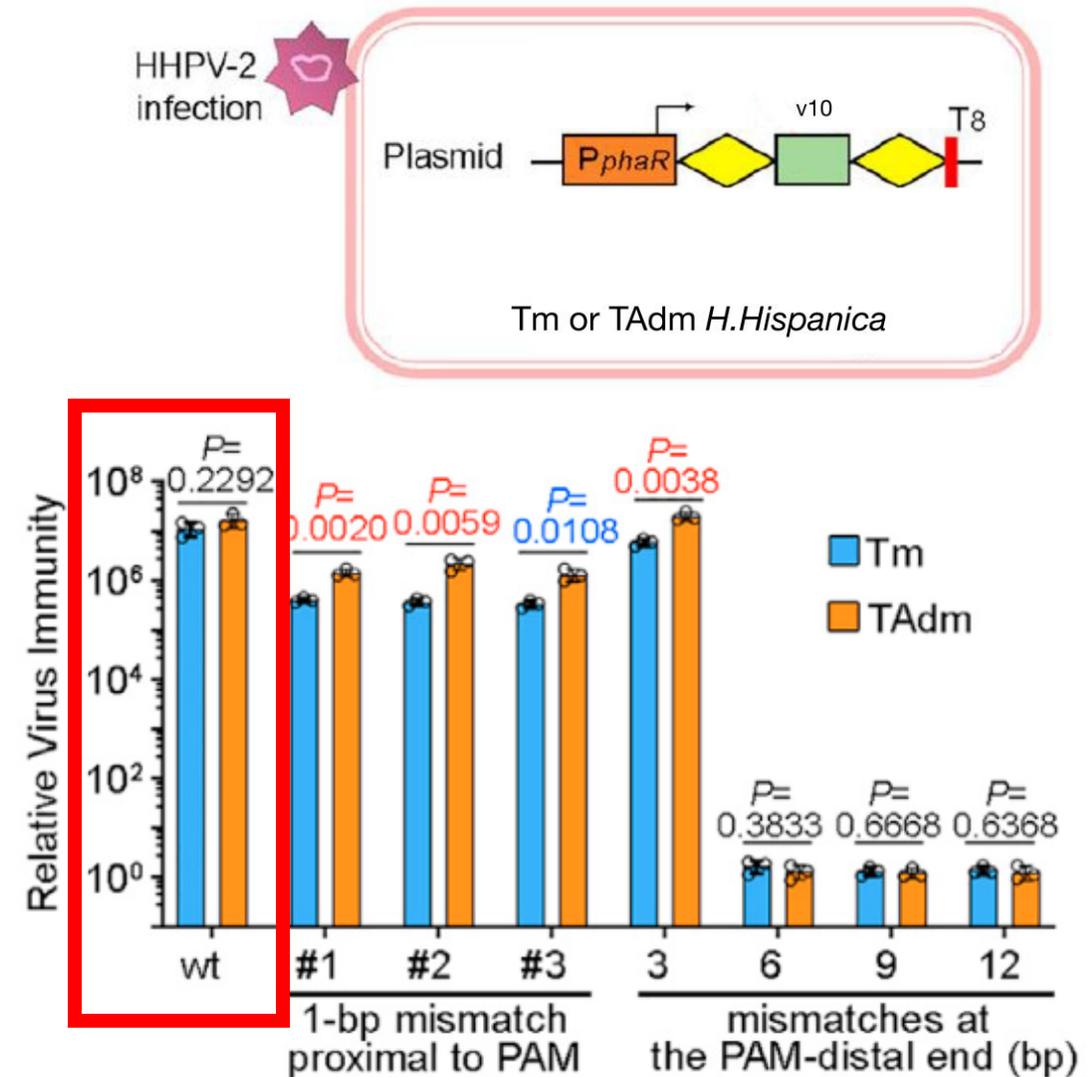
Can *cas* overexpression lead to a more proficient CRISPR immunity?

Model: Tm & TAdm + construct with **synthetic mini-CRISPR with v10 spacer** (targeting HHPV-2 virus) under the control of **P_{phaR} constitutive promoter**
→ Cells are infected with **HHPV-2**

Methods: immunity level revealed by Plaque Assay

Result: equivalent viral immunity levels in both Tm and Tadm
→ Strong response also with wild levels of Cas

BUT viruses can escape CRISPR immunity by **mutating** the protospacer, generating imperfect targets



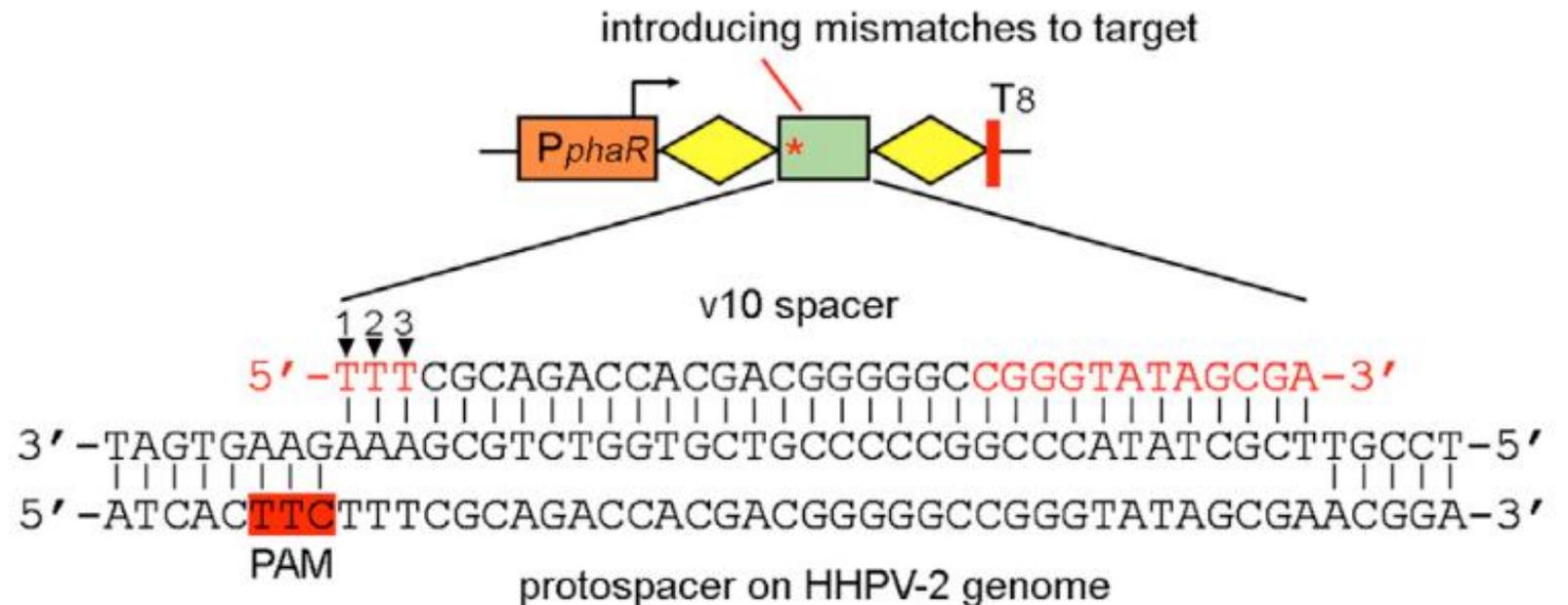
2. CreA-guided *cas* repression reduces autoimmune risk

Can *cas* overexpression lead to more efficient immunity against imperfect targets?

Model: Tm & TAdm + construct with **synthetic mini-CRISPR with v10 spacer** (targeting HHPV-2 virus) with different **mutations** under the control of P_{phaR} constitutive promoter

→ Cells are infected with **HHPV-2**

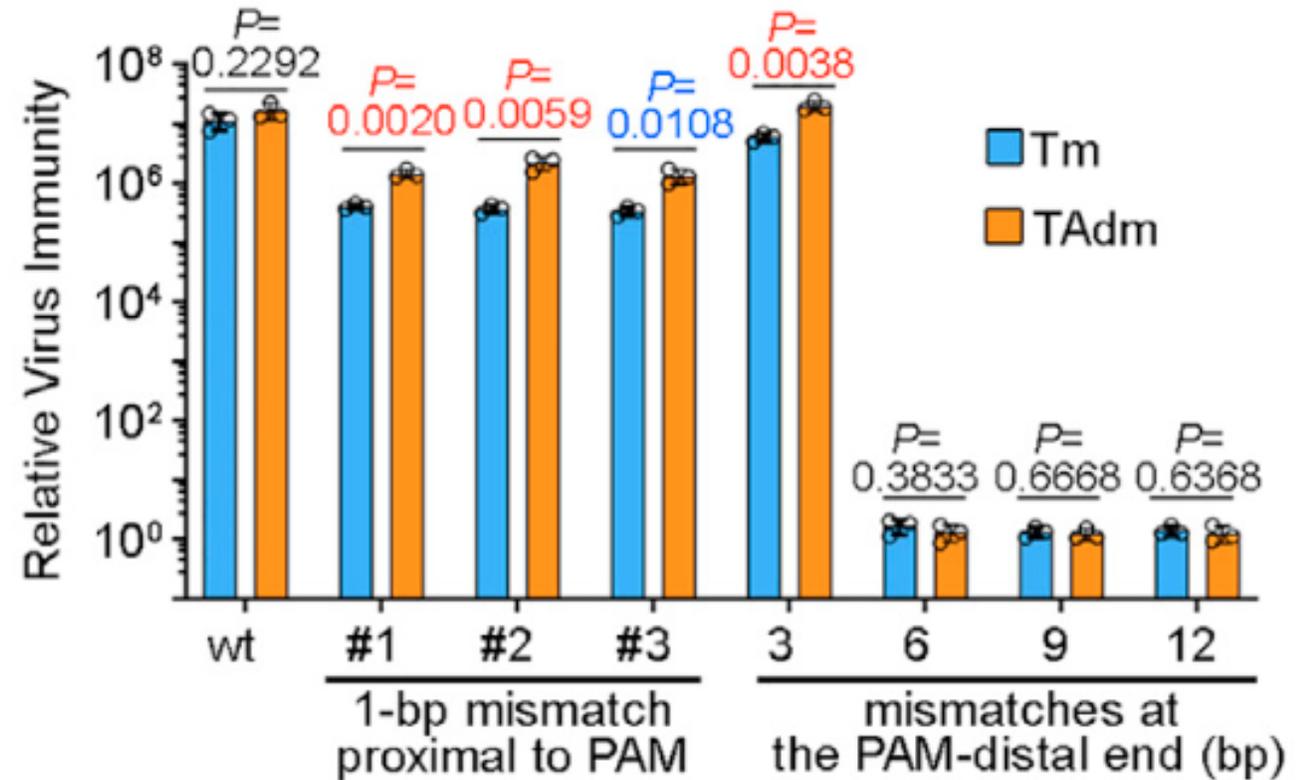
Methods: immunity level measured by Plaque Assay



2. CreA-guided *cas* repression reduces autoimmune risk

Can *cas* overexpression lead to more efficient immunity against imperfect targets?

- Single nt mutation proximal to PAM
Result: TAdm higher immunity level than Tm
- Little mismatches in the PAM-distal end
Result: TAdm higher immunity level than Tm
- Big mismatch in the PAM-distal end
Result: neither TAdm & Tm can activate CRISPR immunity



Interference step

The overexpression of *cas* leads a **more proficient** CRISPR immunity, against **some escape mutants** that interact weakly with crRNA

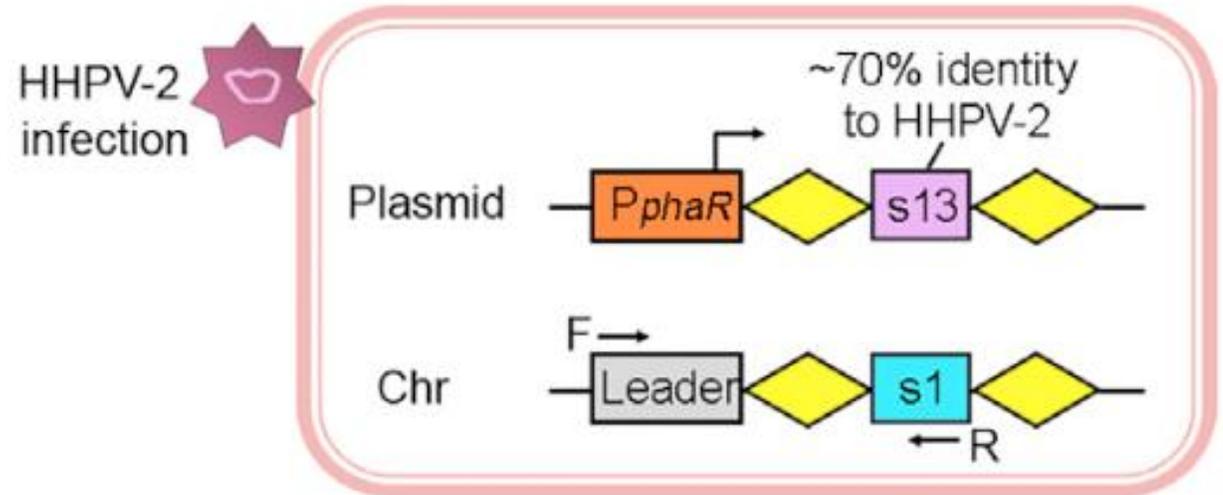
2. CreA-guided *cas* repression reduces autoimmune risk

As *cas* genes are involved in adaptation, can their up-regulation lead to more effective primed spacer acquisition?

Model: WT, Tm and TAdm + construct for overexpression of **crRNA s13**

crRNA s13 = a *H. hispanica* **spacer** that can **activate primed adaptation** (70% identity with HPPV-2)

→ Cells are infected with **HHPV-2 virus** in different quantity (multiplicity of infection = MOI)



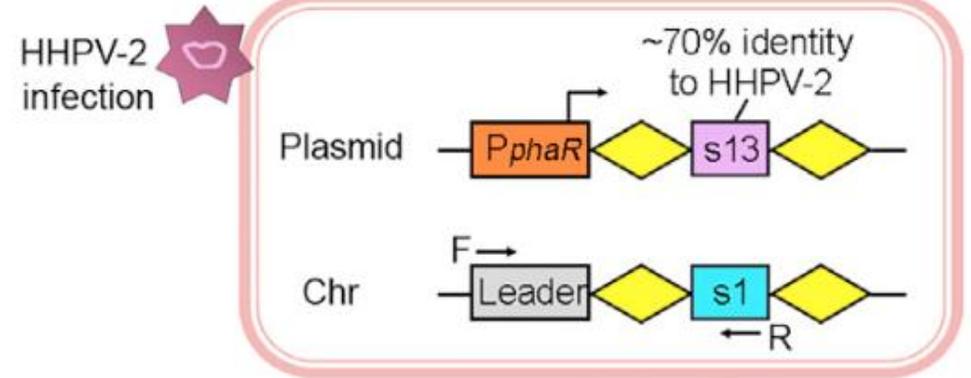
The s13-encoding plasmid makes **all cells express the same quantity of crRNA s13**
→ the results will reflect only the difference in *cas* expression

2. CreA-guided *cas* repression reduces autoimmune risk

Knowing that *cas* genes are involved in adaptation, can their up-regulation lead to a more effective primed spacer acquisition?

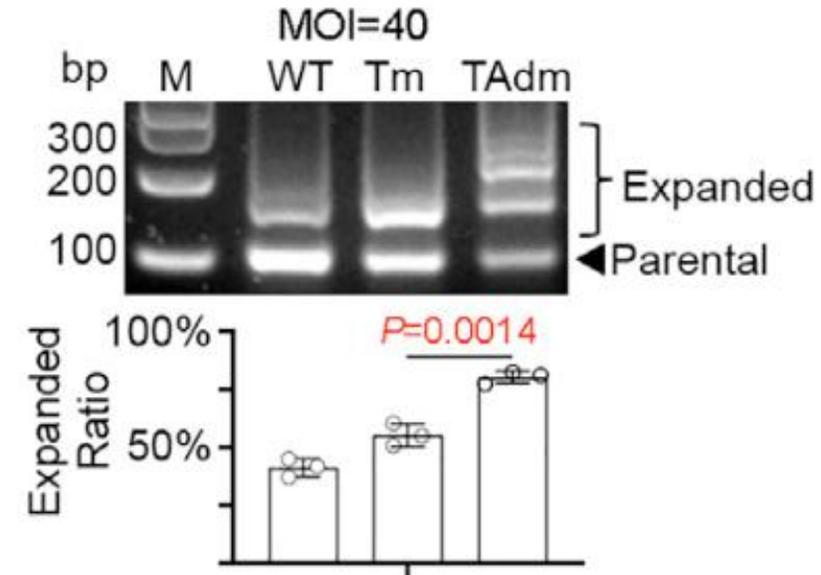
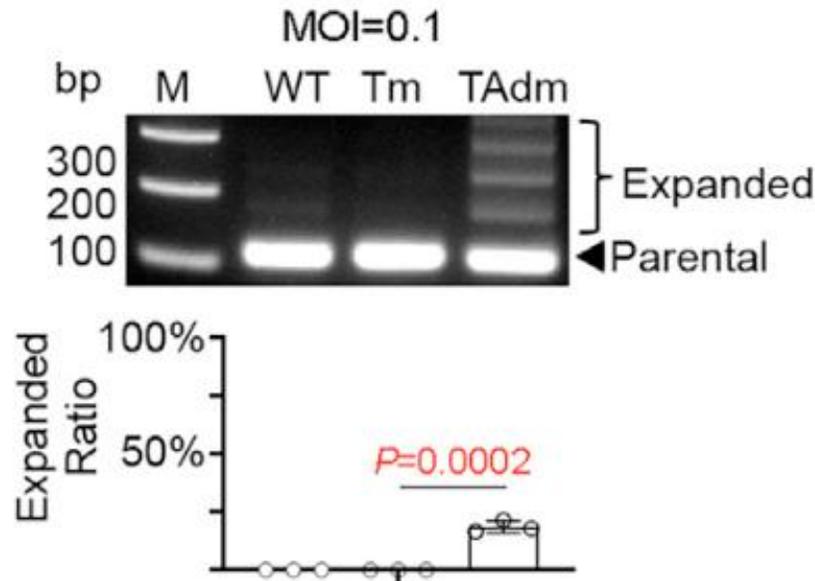
Methods: measure new spacers acquisition through PCR

Result: TAdm show **higher frequency of incorporation** of new spacers in the CRISPR array compared to WT and Tm, in both low (0.1) and high (40) MOI



Adaptation step

The overexpression of *cas* leads a **more proficient primed spacer acquisition**

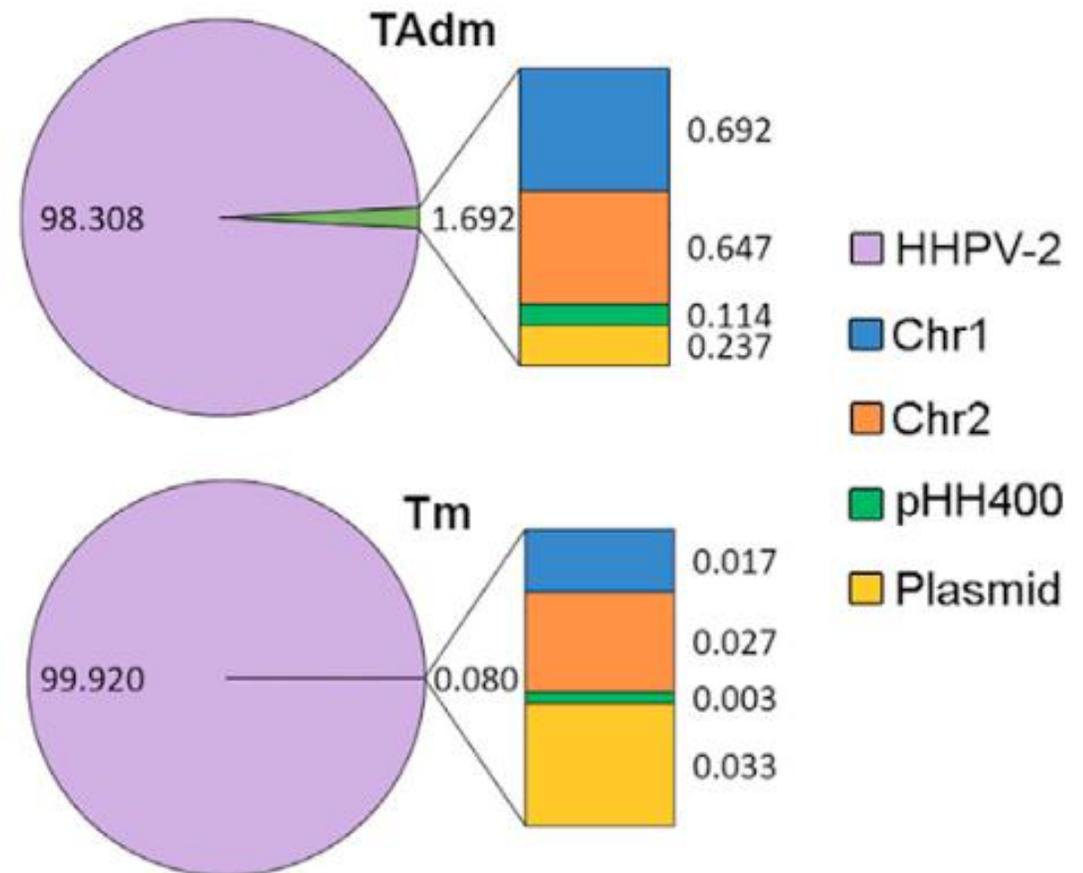


BUT

Can *cas* up-regulation lead to a higher risk of acquisition ALSO of self-derived spacers?

Result: Illumina seq data reveal → **TAdm** mistakenly acquires **self-derived spacers** at a higher frequency than Tm, leading to **autoimmunity**

TAdm 1.69% vs Tm 0.08%
self-derived spacers

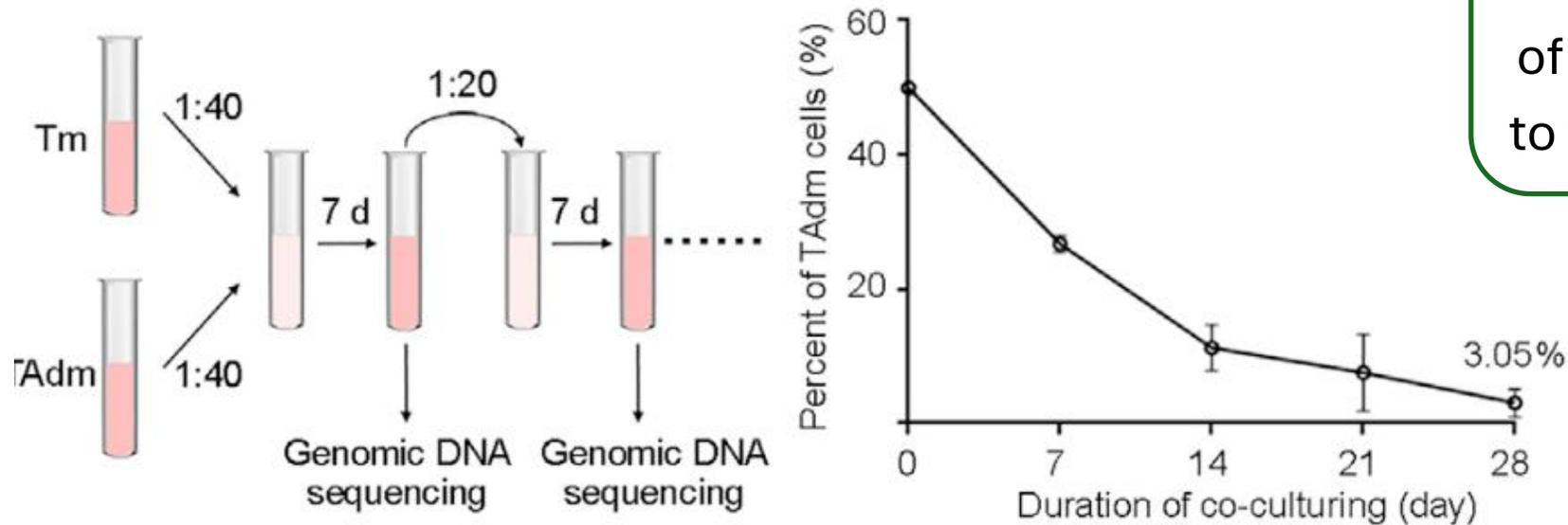


2. CreA-guided *cas* repression reduces autoimmune risk

Does the accumulation of self-targeting spacers compromise the fitness?

Model: Tm & TAdm in long-term **co-culture**

Methods: measure cell percentage by high-throughput DNA seq



The **fitness** is mainly **compromised** because of **autoimmunity** related to the *cas* overexpression

Result: Tm outcompeted (= have a better fitness) TAdm over time, with proportion of TAdm dropping to 3.05%

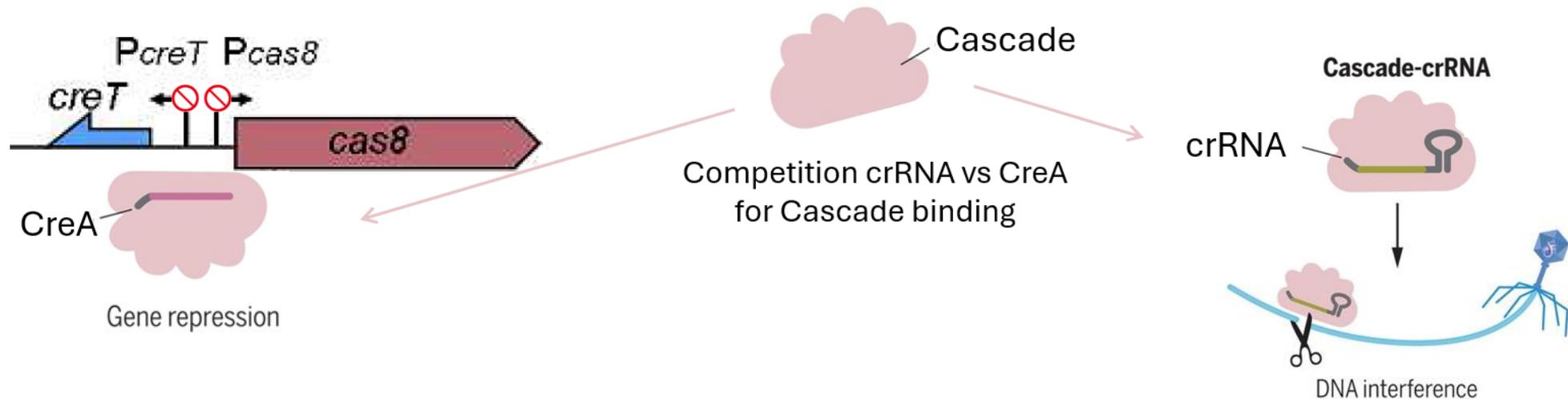
2. CreA-guided *cas* repression reduces autoimmune risk

Conclusion: the repression of *cas* genes is necessary to balance viral immunity with the **prevention of autoimmunity**

3. The entity of CreA-guided gene repression depends on crRNA levels

crRNA and CreA both bind Cascade:

- **crRNA + Cascade** = immunity effector machinery
 - **CreA + Cascade** = repression complex \rightarrow suppression of P_{cas8}
- competition?



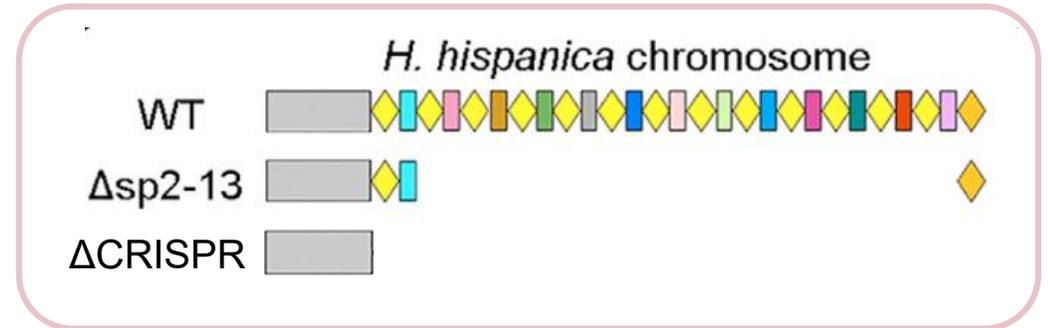
3. The entity of CreA-guided gene repression depends on crRNA levels

Does crRNA abundance affect CreA repression of P_{cas} ?

Hyp: \downarrow **crRNA** \Rightarrow \uparrow CreA-mediated *cas* repression

Model:

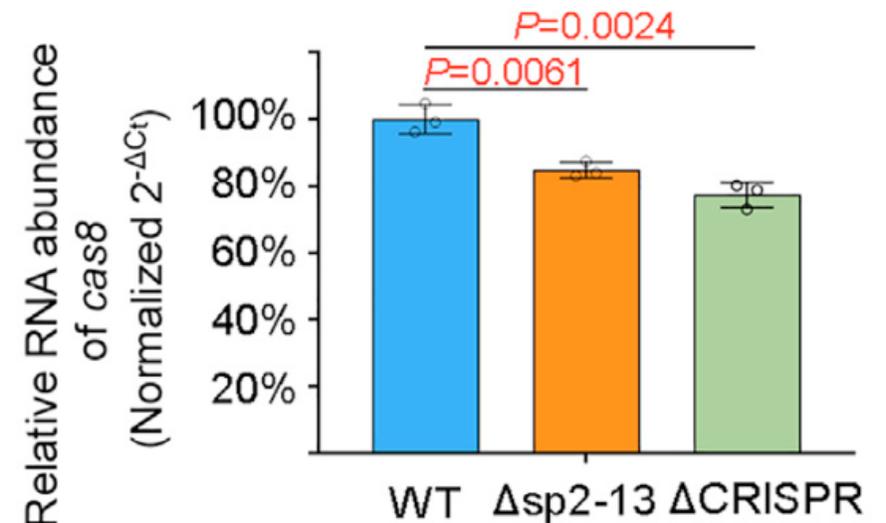
- WT cells
- **Δ CRISPR** cells
- **Δ sp2-13** (= CRISPR array with only 1 spacer) cells



Methods: quantify *cas8* transcription by RT-qPCR

Result: in **Δ CRISPR & Δ sp2-13** *cas8* expression decreases by ~20% compared to wt

(= more repression if less crRNA are present)



3. The entity of CreA-guided gene repression depends on crRNA levels

Does crRNA abundance affect CreA repression of P_{cas} ?

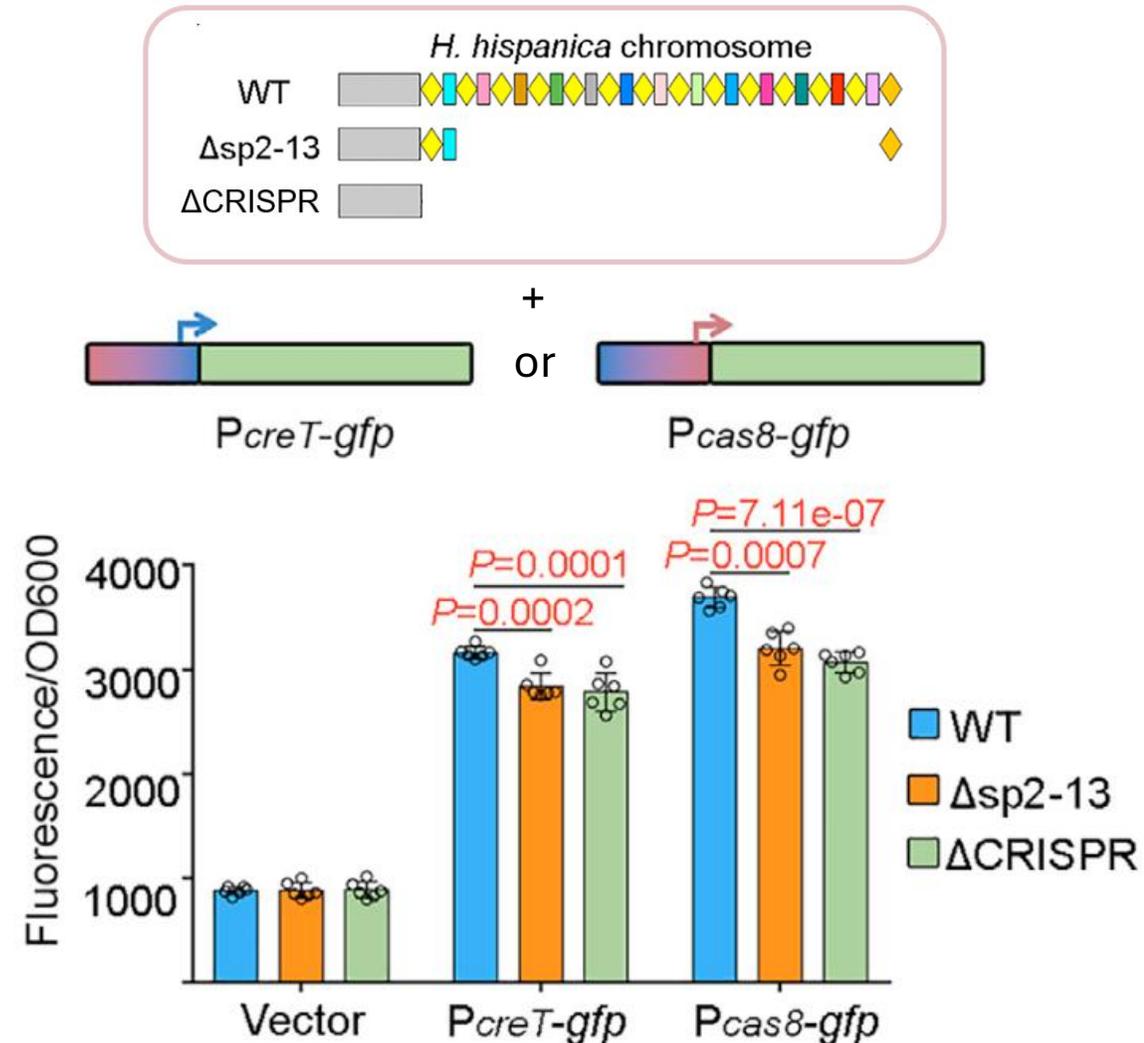
Hyp: \downarrow crRNA \Rightarrow \uparrow CreA-mediated *cas* repression

Model: WT, Δ CRISPR & Δ sp2-13 + P_{cas8} -*gfp* or P_{creT} -*gfp*

Methods: evaluate P_{cas8} or P_{creT} activity **at different crRNA levels** by *gfp*-reporter assay

Result: in Δ CRISPR & Δ sp2-13 promoters P_{cas8} and P_{creT} activity **decreases** by 10%-20%

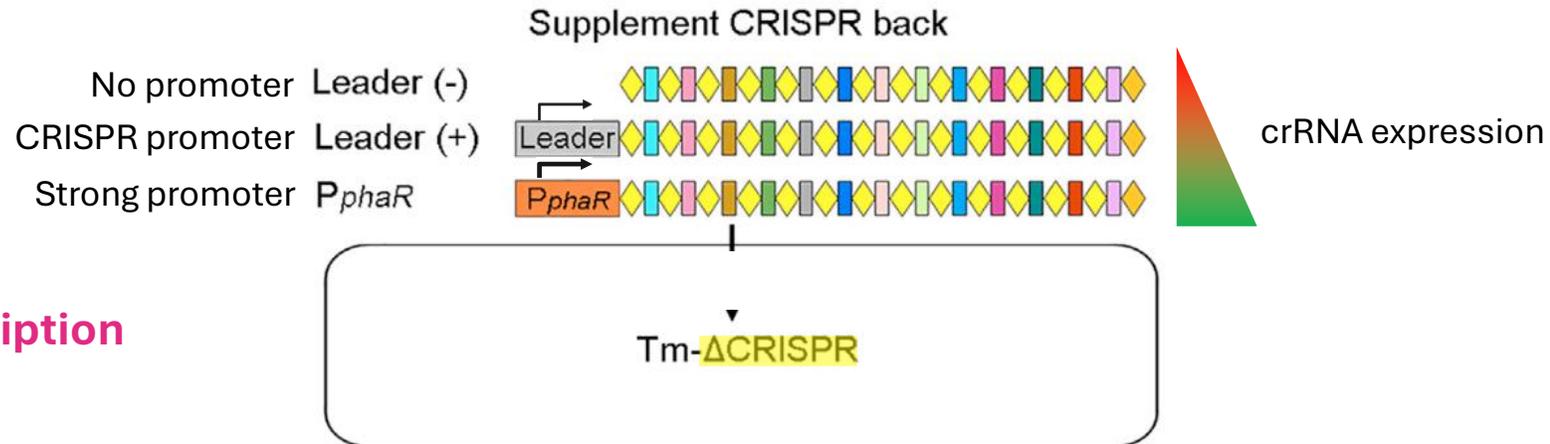
CreA tightly **suppresses** P_{cas8} & P_{creT} in case of **fewer** or **absence** of CRISPR **spacers** (crRNA)



3. The entity of CreA-guided gene repression depends on crRNA levels

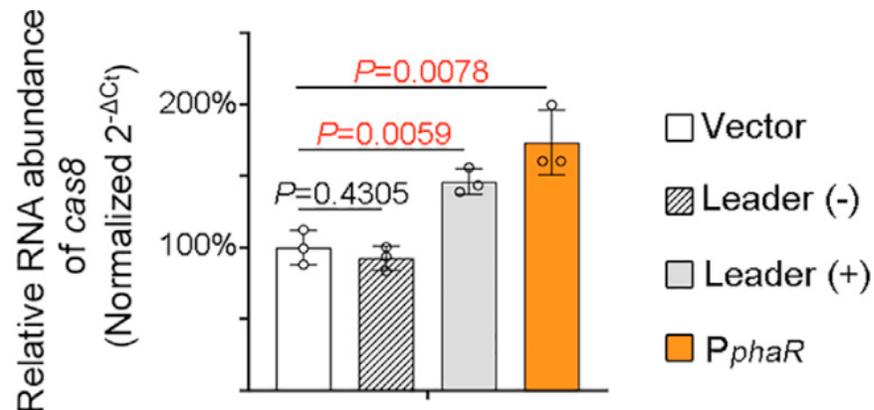
Does crRNA abundance affect CreA repression of P_{cas} ?

Hyp: \uparrow crRNA \Rightarrow \downarrow cas repression CreA mediated



Evaluate **cas8 transcription** through **RT-qPCR**

Result: crRNA expression leads to almost **2-fold increase of cas8 transcripts** in comparison with hosts having empty vector or leader-less array



CreA-guided gene repression can be alleviated by expanding the crRNA pool

3. The entity of CreA-guided gene repression depends on crRNA levels

Does the overexpression of CreA lead to tighter repression of P_{cas8} ?

Hyp: $\uparrow cas$ repression CreA mediated => $\downarrow crRNA$

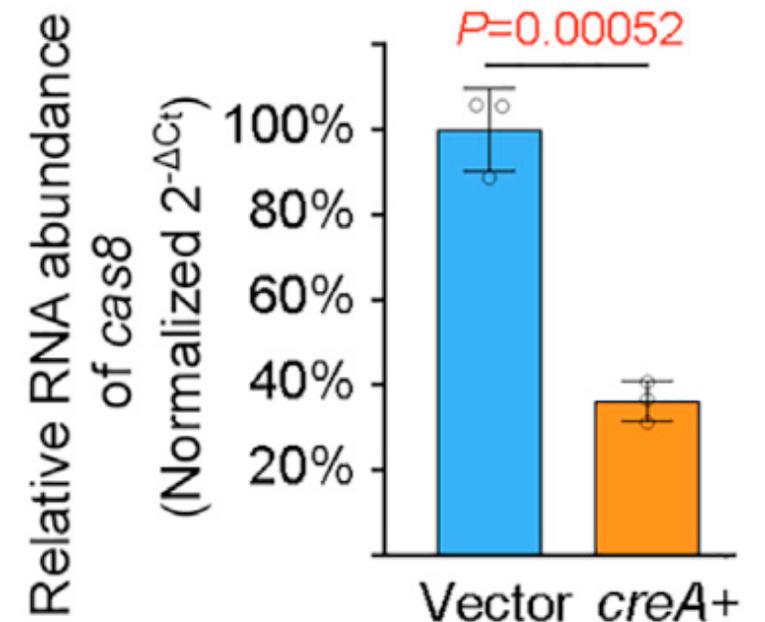
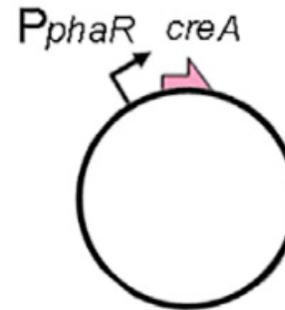
Model: WT + constructs:

- Empty vector
- **Strong promoter P_{phaR} upstream CreA** → **overexpression of CreA**

Methods: quantify **$cas8$ transcription** by RT-qPCR

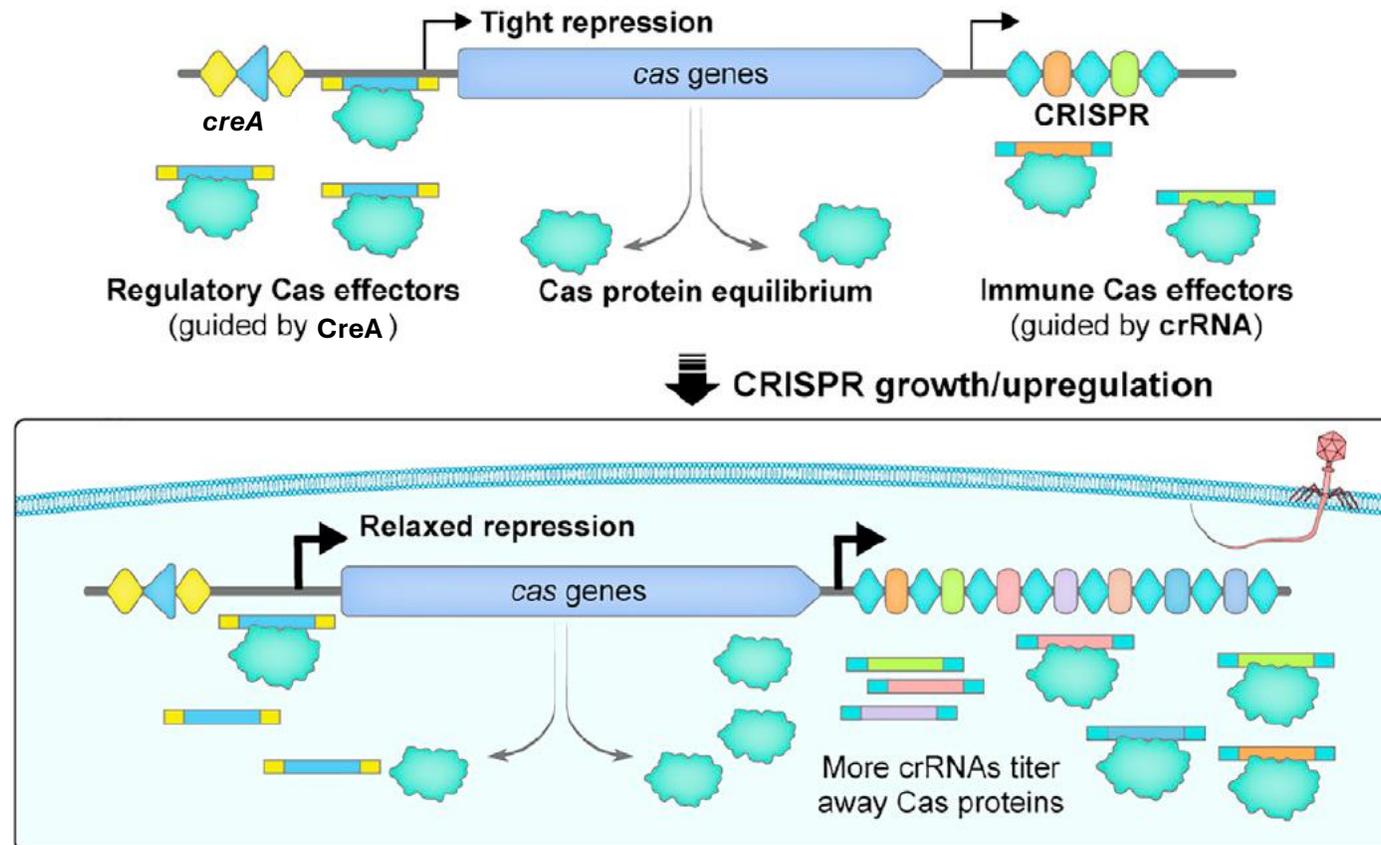
Result: relative abundance of **$cas8$ transcript** decreases by 60% in cells **overexpressing CreA**

CreA overexpression leads to an **enhanced repression of cas genes**

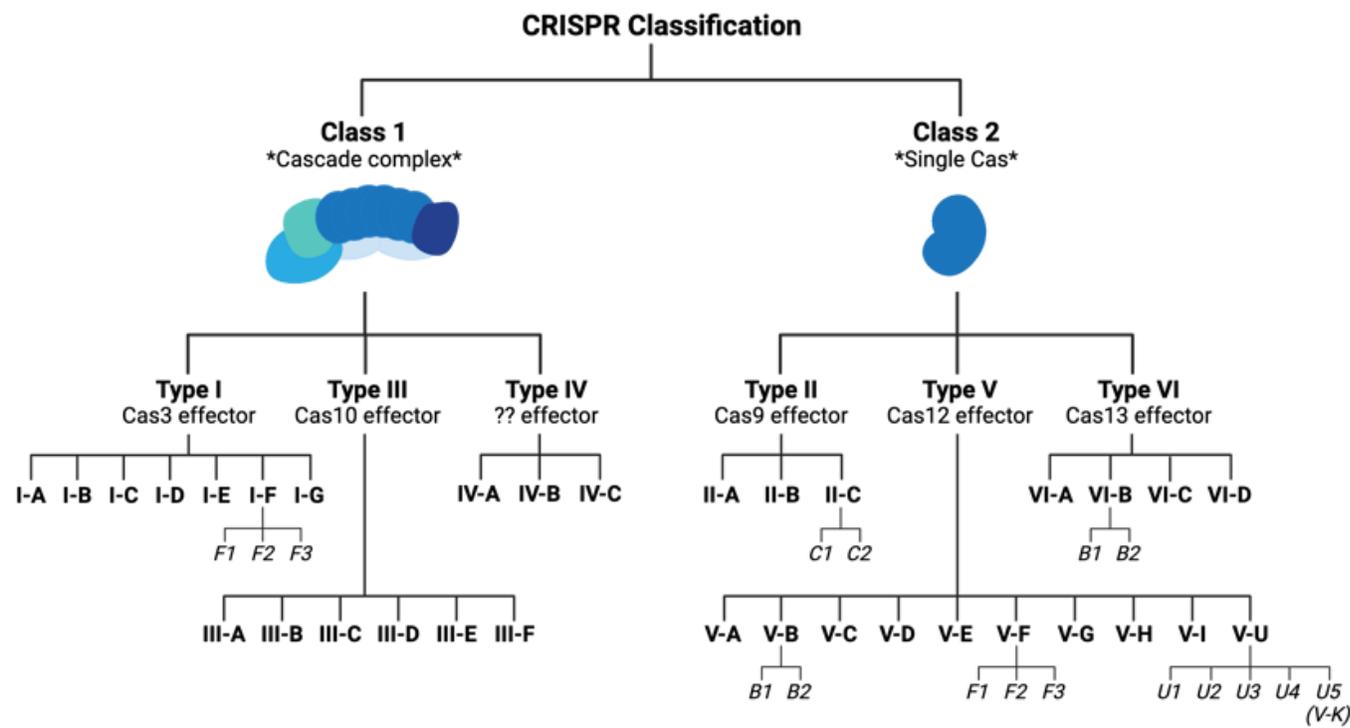


3. The entity of CreA-guided gene repression depends on crRNA levels

Conclusion: **cas autorepression** can be **fine-tuned** by altering the relative transcripts **abundance** of the defensive RNA guides **crRNA** and regulatory **CreA**



4. Inspecting the wider **distribution** of CreA in different CRISPR-Cas systems

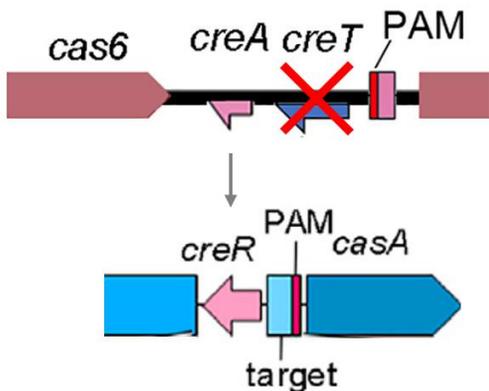


4. Inspecting the wider distribution of CreA

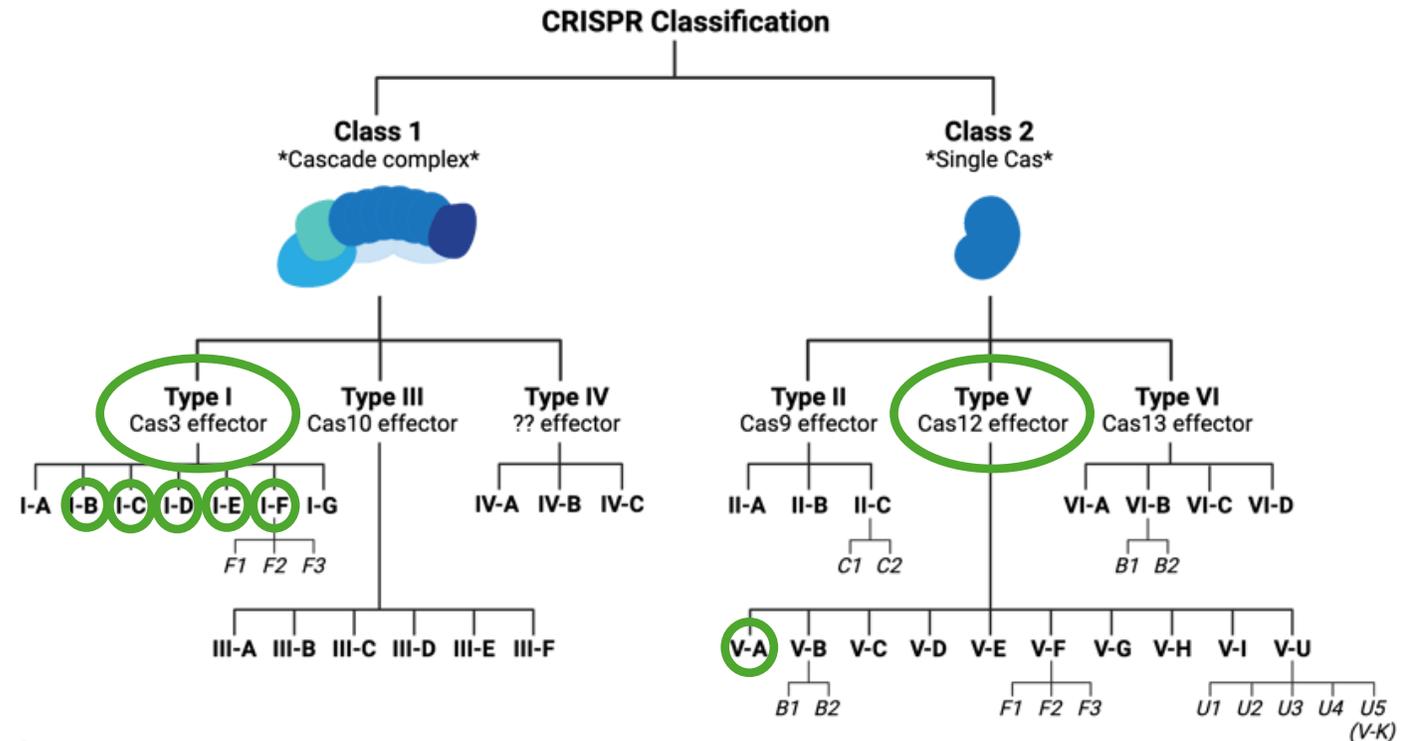
Sequence analysis of more CRISPR-Cas systems revealed **creA-like** elements also in: various **type I** systems (Cascade complex) & dozens of **type V-A** systems (Cas12a)

In most cases **CreA** targeted the **promoter of cas genes**

Many of these *creA*-like elements **DO NOT co-locate with a toxic gene!**



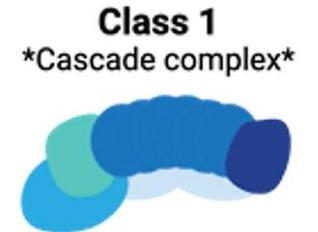
«standalone» *creA* genes will be referred as ***creR*** (*cas*-regulating RNA)



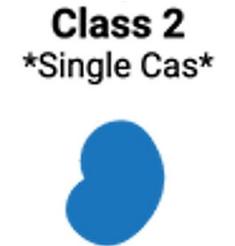
Cas autoregulation directed by **CreA** or **CreR** appears to be a **general mechanism** of both **class 1** and **class 2** CRISPR systems

Experimental validation of **CreR** regulatory role in directing the **autorepression circuit**:

5. CreR in **class 1** type **I-E** CRISPR system



6. CreR in **class 2** type **V-A** CRISPR system



Conclusion: Cas **autoregulation** directed by **CreA** or **CreR** is a **general mechanism** of both **class 1** and **class 2** CRISPR systems and **monitors crRNAs** abundance in the cell

7. Phage **Acr** proteins can **relieve** or **subvert** CreR-guided **Cas** autorepression

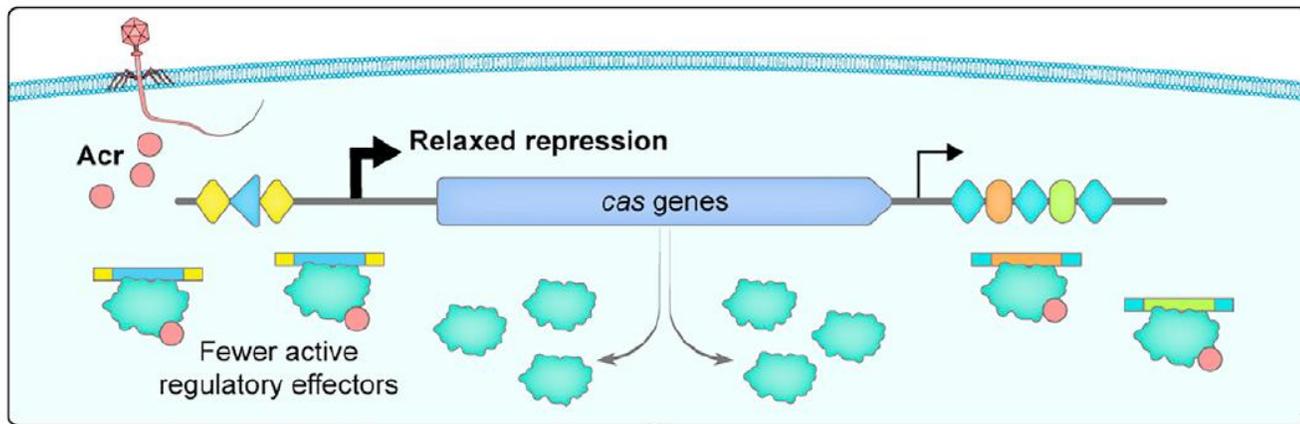
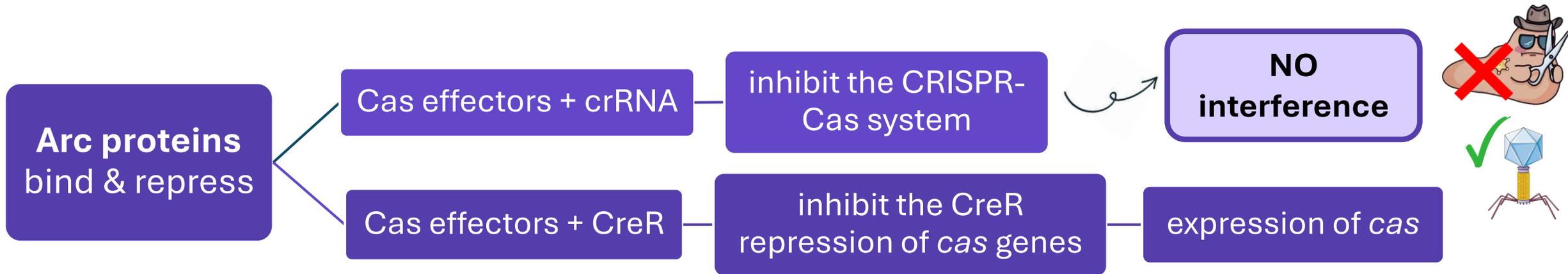
cas autoregulation circuit can surveil phage attack

Anti-Anti-CRISPR effect:

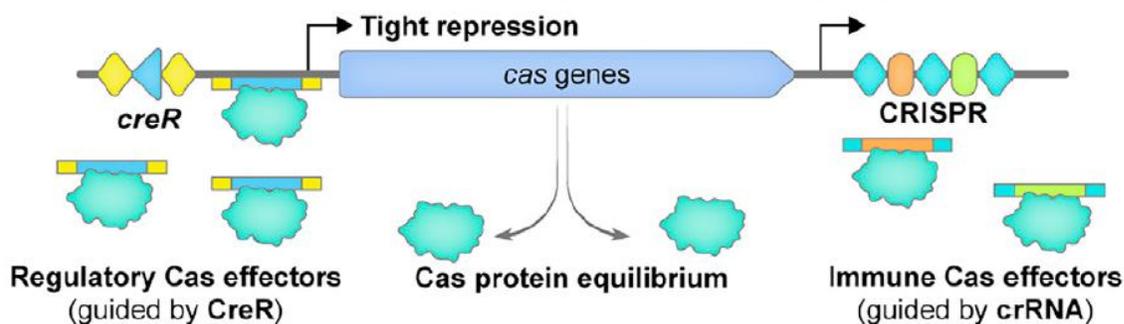
Arc proteins can **relieve** or **subvert** the **CreR** repression leading to the increase of **new Cas** production for phages defeat



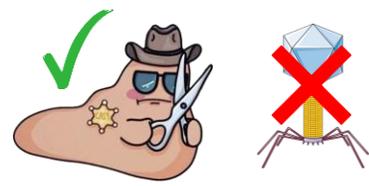
7. Phage Acr proteins can relieve or subvert CreR-guided Cas autorepression



Anti-CRISPR (Acr) attack



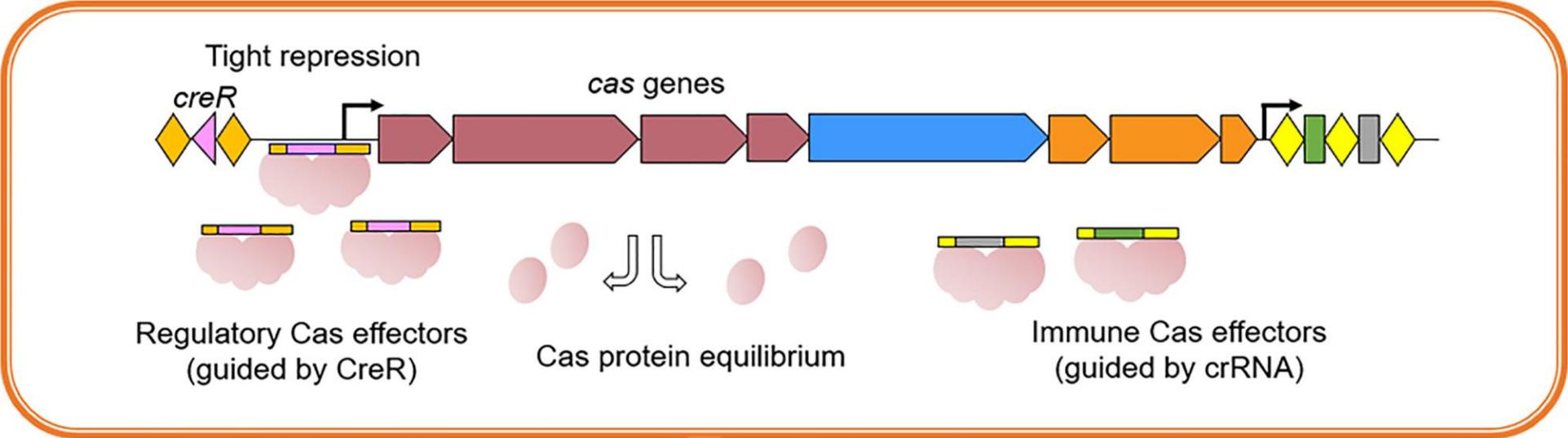
Enhanced CRISPR-Cas immune response



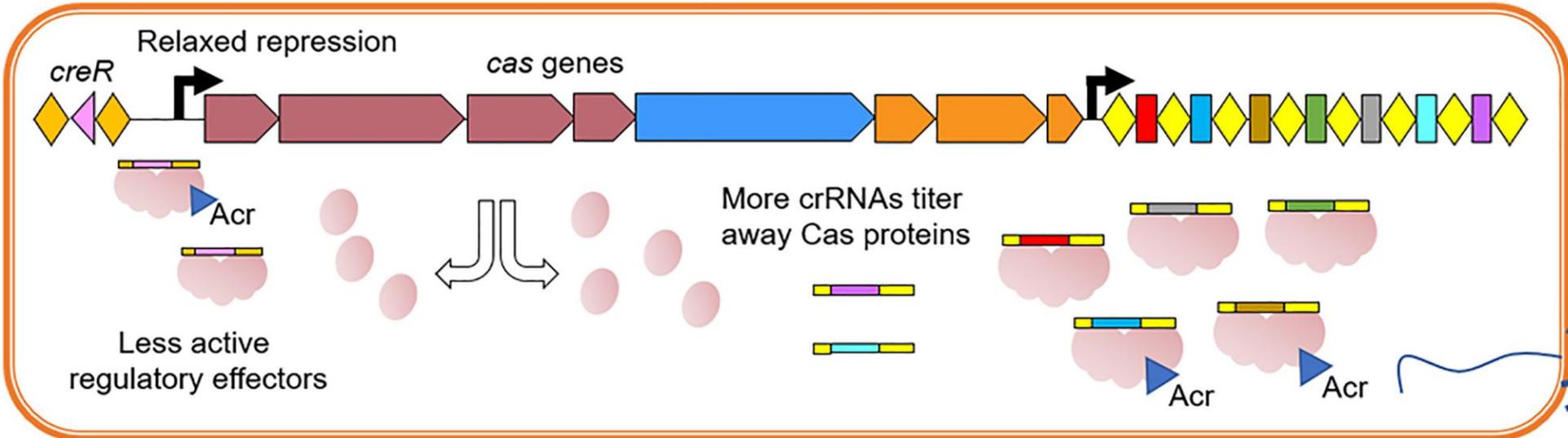
OUTLINE

- INTRODUCTION
- STUDY AIM
- RESULTS
- **CONCLUSION**

Model of CreA or CreR-guided Cas autorepression



CRISPR growth/upregulation or Acr expression



Cas autoregulation circuit directed by crlRNA CreA or CreR:

Balances benefits and downsides of CRISPR-Cas



Might represent a distinct **anti-anti-CRISPR** strategy



With CreT, might have **promoted persistence of CRISPR-Cas** in prokaryotes



Exploring ncRNA in bacterial immunity and phages counterattack

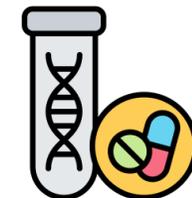


Revealing molecular parallels to **conserved eukaryotic antiviral pathways**



Developing novel **biotechnologies**

- **Control** strategies for CRISPR-Cas biotechnologies
- Improving **antimicrobial** phage therapy efficacy



References

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- Levy A, Goren MG, Yosef I, Auster O, Manor M, Amitai G, Edgar R, Qimron U, Sorek R. CRISPR adaptation biases explain preference for acquisition of foreign DNA. *Nature*. 2015 Apr 23;520(7548):505-510. doi: 10.1038/nature14302. Epub 2015 Apr 13. PMID: 25874675; PMCID: PMC4561520.

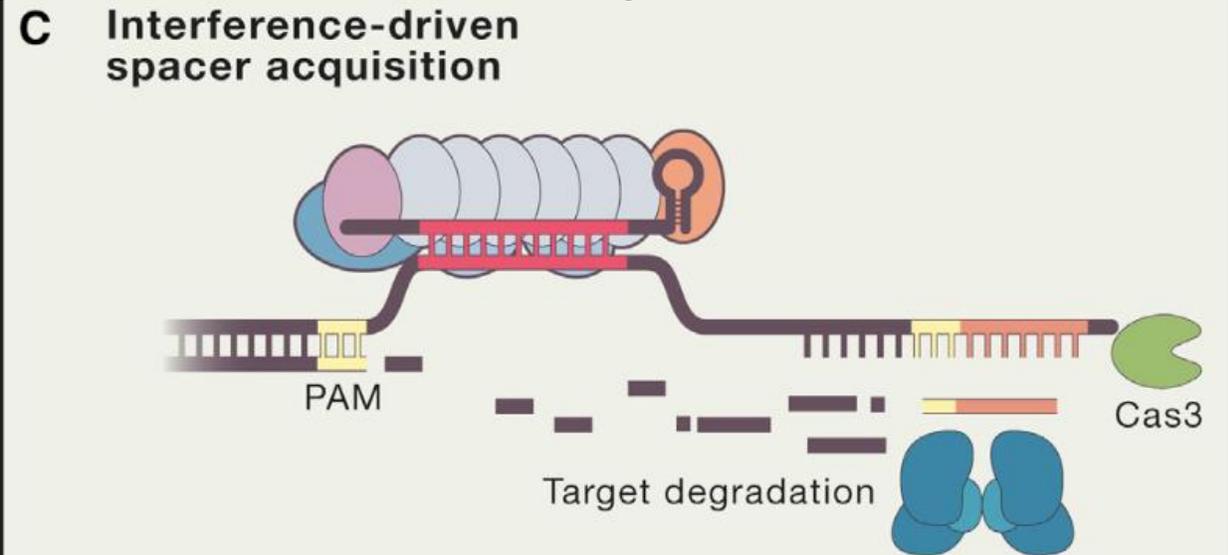
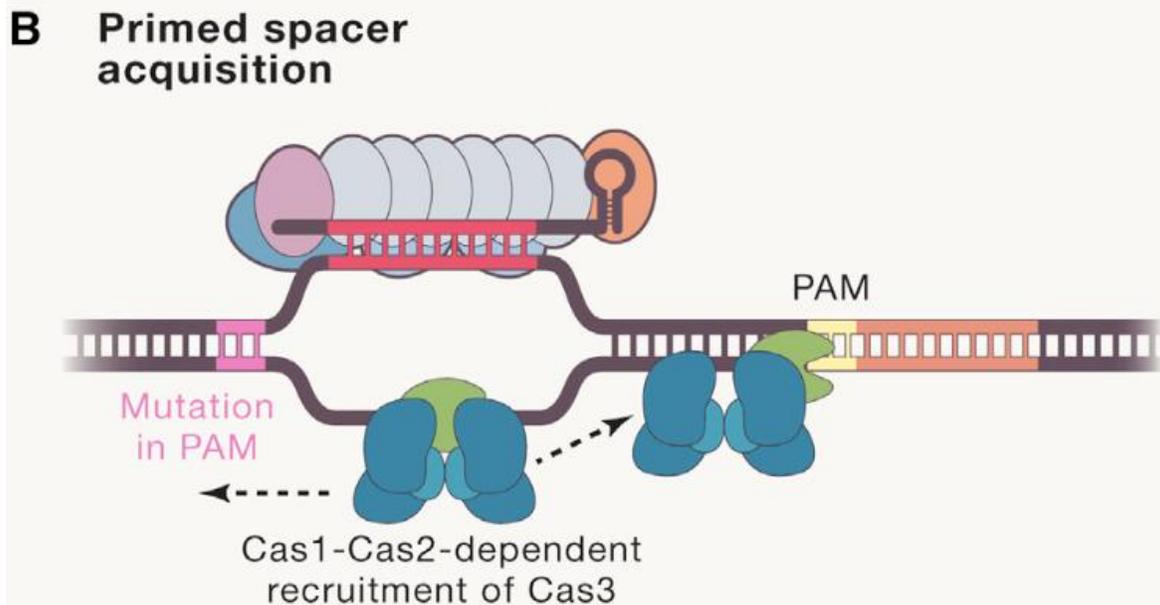
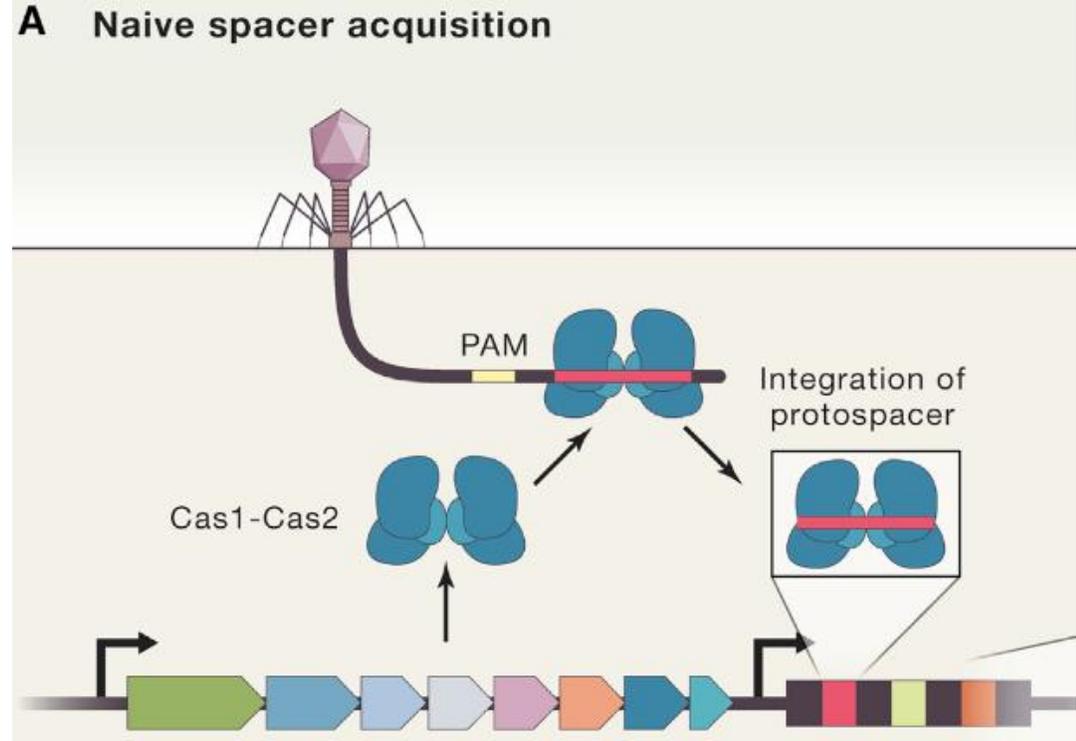


**Complementary
slides**

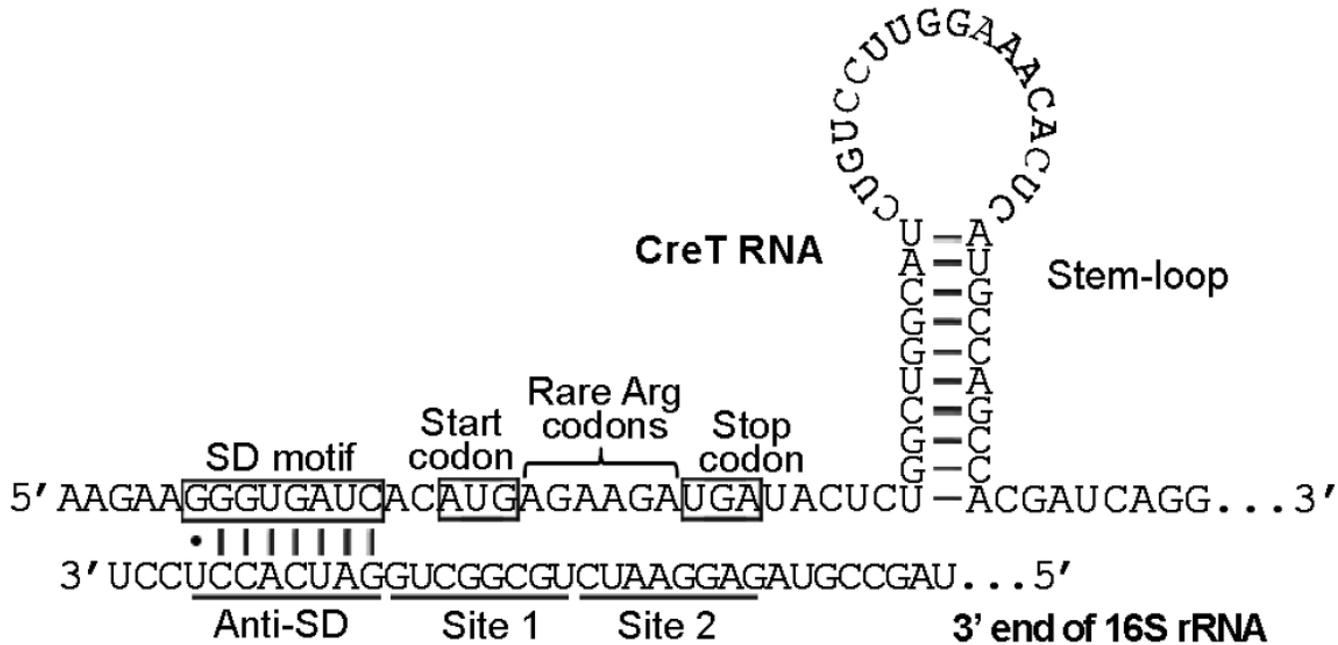
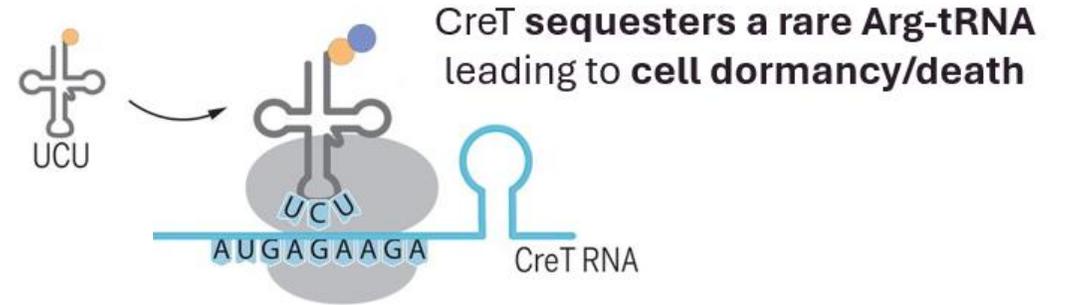
1. Adaptation

Methods spacer acquisition:

- A. Naïve:** from **previously unfronted** MEGs
- B. Primed:** from **pre-existing spacers partially complementary** to previously encountered MEGs
- C. Interference-driven:** from **interference degraded** MEGs



CreT toxic RNA

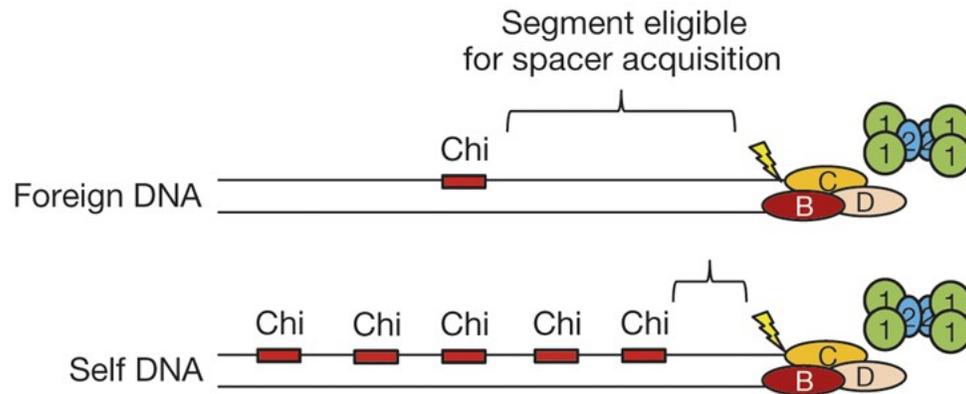
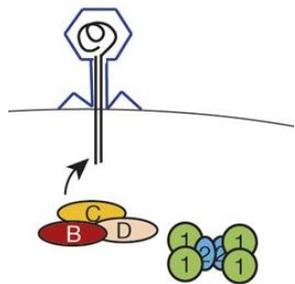
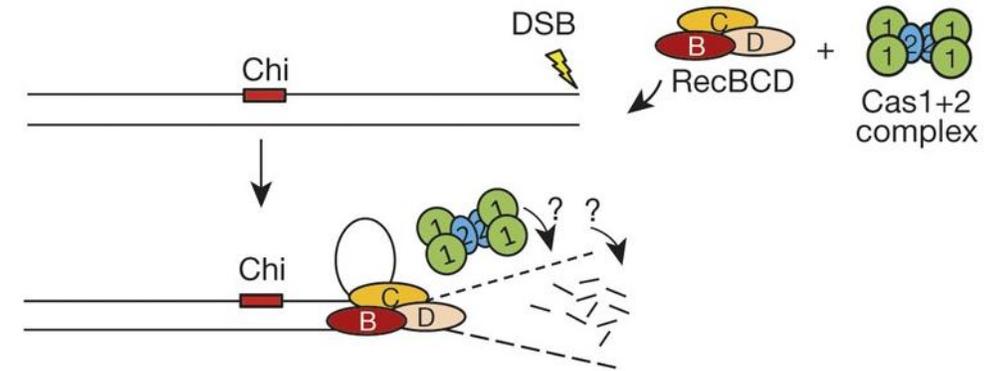


Adaptation machinery lacks intrinsic ability to distinguish self- and non-self DNA → autoimmunity risk

E. coli:

DSB/linear DNA => RecBCD indiscriminately unwinds/**degrades** DNA until the nearest **Chi** site

Generated DNA fragments = **substrates for spacer acquisition** by Cas1–Cas2



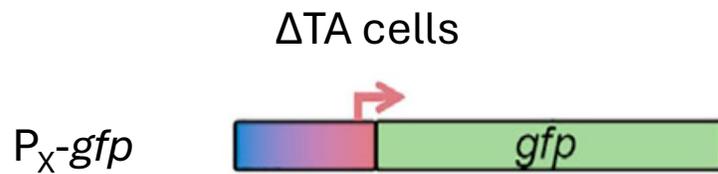
↓ **Chi sites density on foreign DNA** = more extensive RecBCD processing
=> **more** substrate for spacer acquisition

↑ **Chi sites density on chromosome** = less RecBCD processing
=> **reduced** self-derived spacer acquisition

A matter of «preferential acquisition» rather than «recognition» of non-self DNA

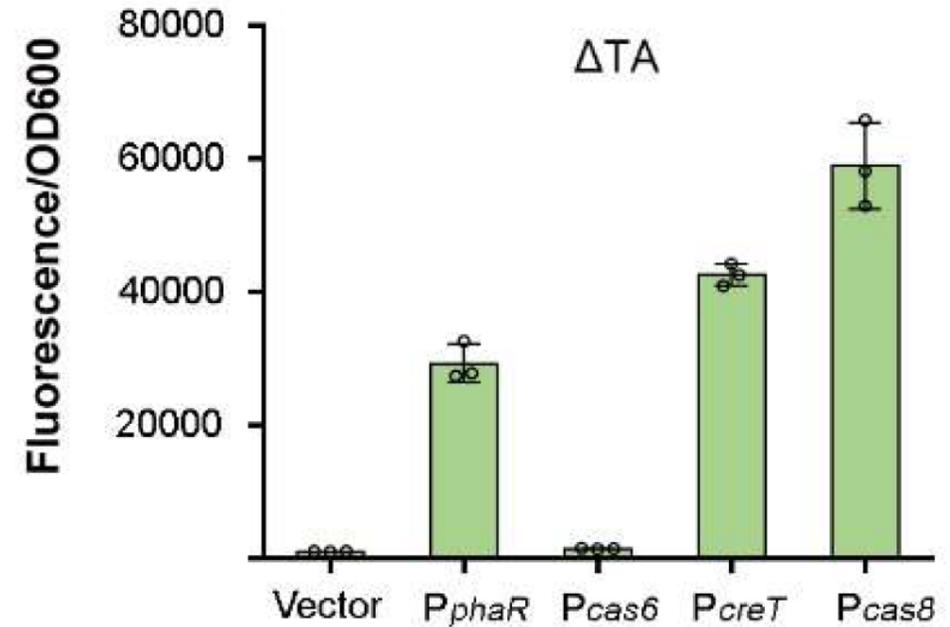
1. Demonstrating that *H. hispanica* CreA represses also *cas* transcription

Validation and characterization of *cas8* promoter



GFP-reporter system to **compare** the activity of:

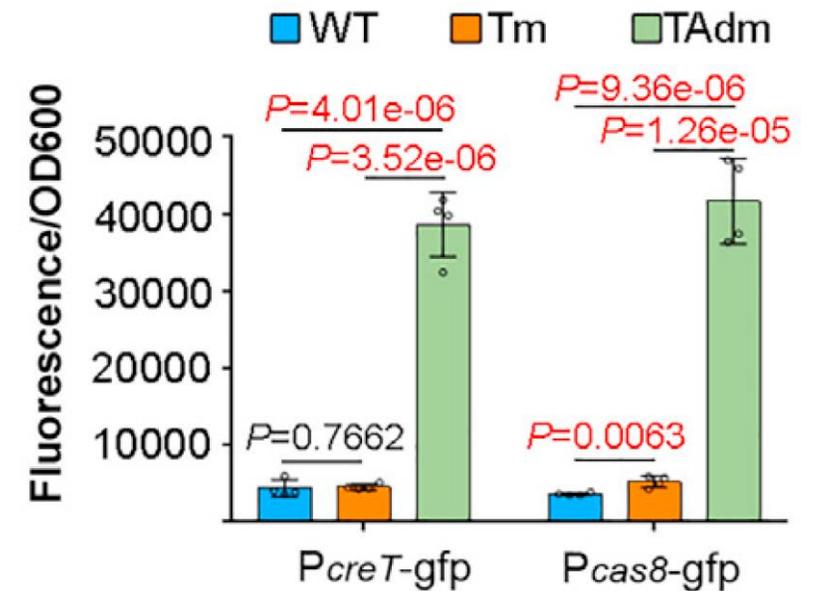
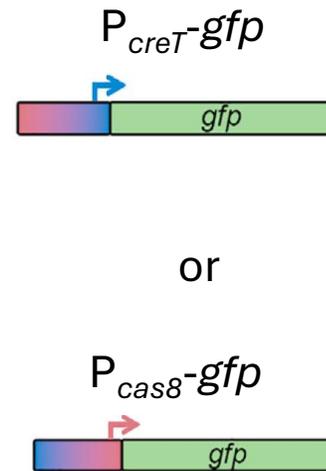
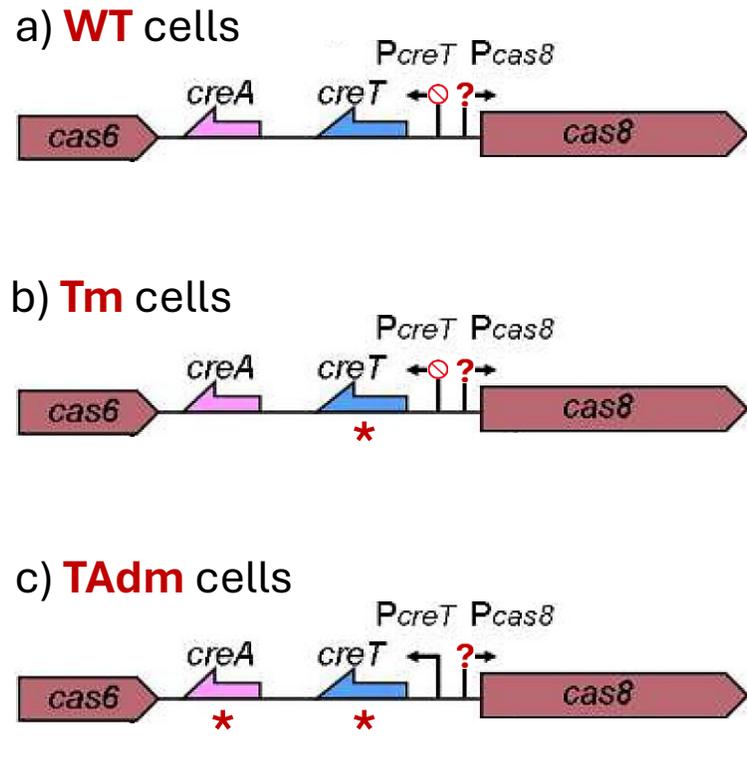
- P_{cas8}
- P_{cas6}
- P_{creT}
- P_{phaR} (strong constitutive promoter)



P_{cas8} exhibits **135 times** the activity of P_{cas6} and outperforms P_{creT} and P_{phaR} by 1.4 and 2.0 times

1. Demonstrating that *H. hispanica* CreA represses also *cas* transcription

Is P_{cas8} also repressed by CreA?



P_{cas8} and P_{creT} are **both >10-fold more active in TAdm** compared to WT and Tm

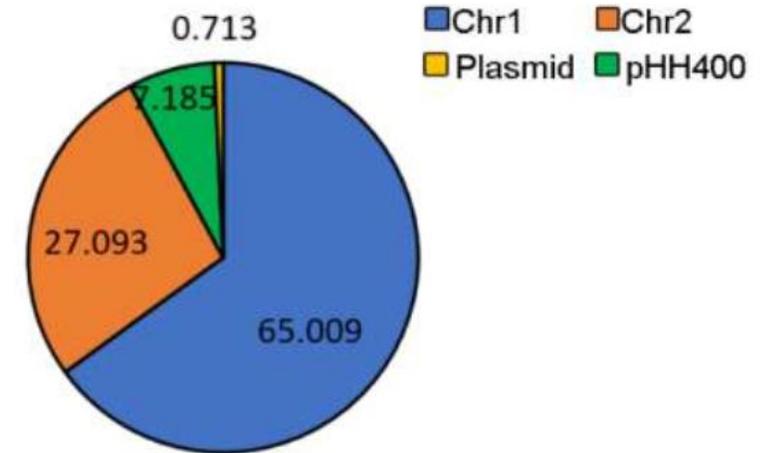
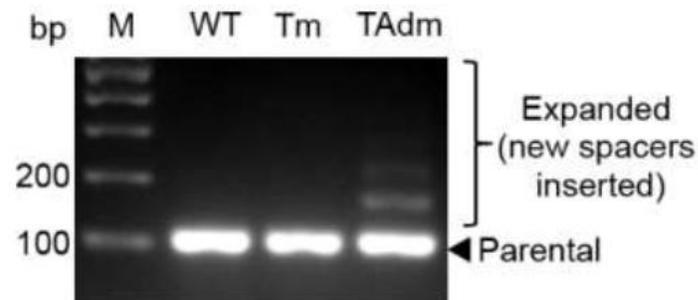
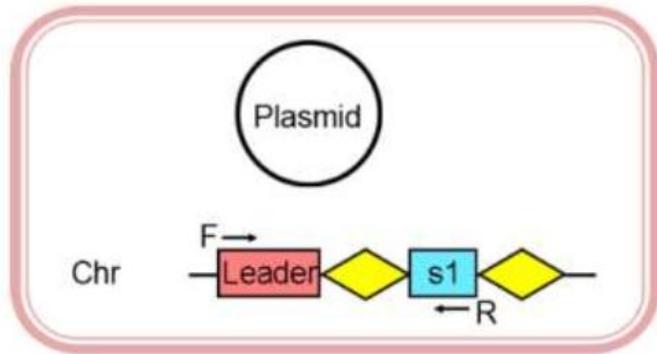
GFP-reporter assay to evaluate P_{cas8} and P_{creT} activity depending on **CreA capability to bind its target site**

2. CreA-guided *cas* repression reduces autoimmune risk

Are self-derived spacers gained through primed or naïve acquisition?

Model: WT, Tm & TAdm + empty vector

Methods: measure new spacers acquisition through PCR



Result: Without virus infection TAdm show **inefficient** acquisition of **endogenous** spacers

Maybe the **13 pre-existing spacers** in the **CRISPR array** partially match the **endogenous DNA**

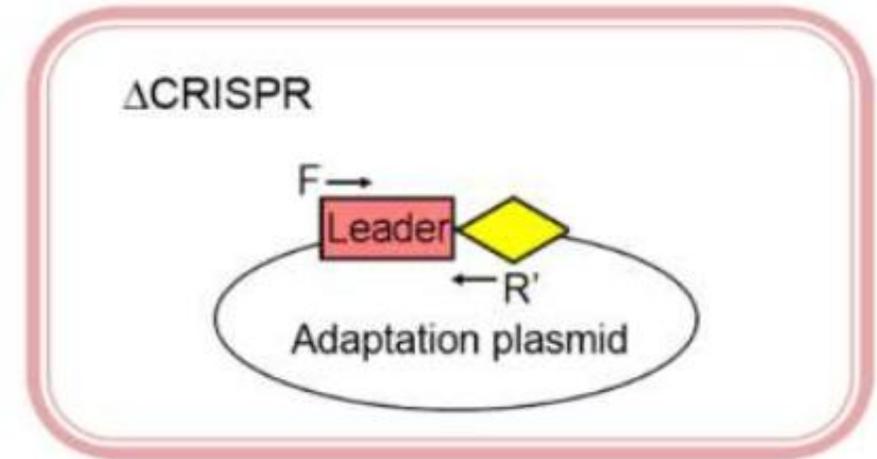
2. CreA-guided *cas* repression reduces autoimmune risk

Are self-driven spacers gained through primed or naïve acquisition?

Model: WT, Tm- Δ CRISPR & TAdm- Δ CRISPR

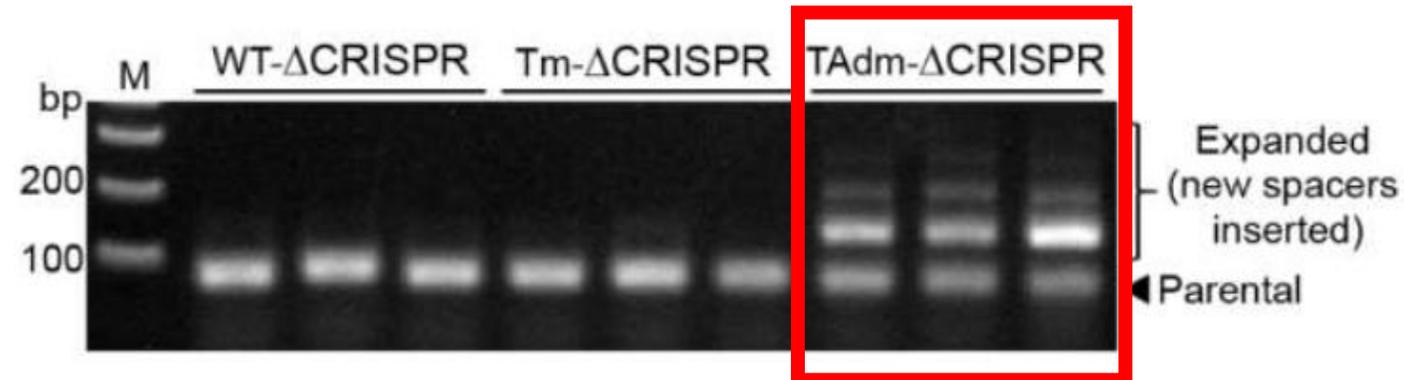
= no possibility of primed acquisition of new spacers

Methods: measure new spacers acquisition through PCR



Result: TAdm- Δ CRISPR has **new spacers without infection** through naïve acquisition

cas up-regulation increases the adaptation frequency but with a **higher risk of self-targeting** through **naïve spacer acquisition**



2. CreA-guided *cas* repression reduces autoimmune risk

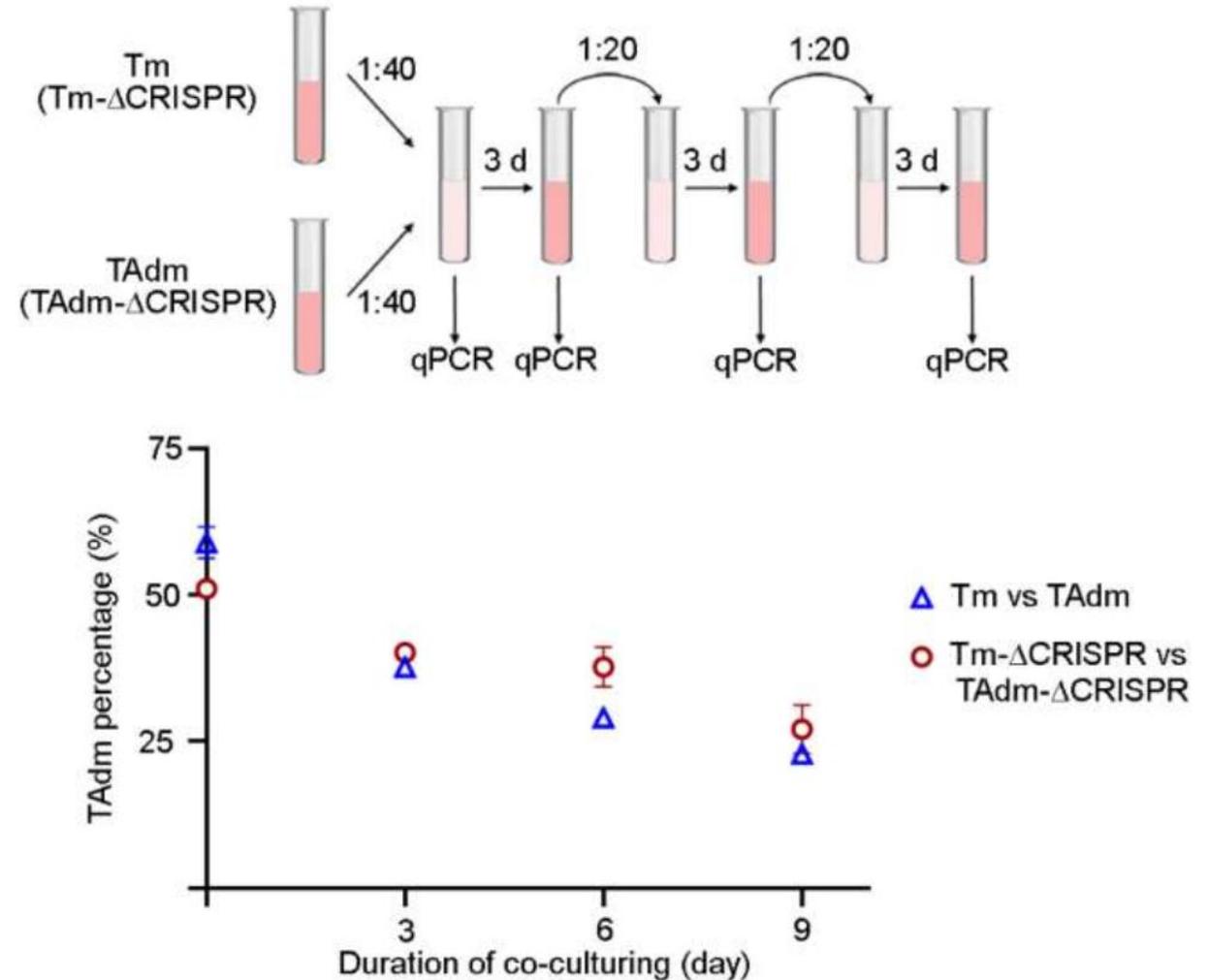
Does the accumulation of self-targeting spacers compromise the fitness?

Model: Tm- Δ CRISPR & TAdm- Δ CRISPR in long-term co-culture

Methods: measure cell percentage by qPCR

Result: Tm- Δ CRISPR outcompeted TAdm- Δ CRISPR over time, but with a **slower decrease**

The fitness is **mainly** compromised because of **autoimmunity**, but also **other metabolic disadvantages** related to the *cas* overexpression



3. CreA-guided gene repression monitors crRNA levels

CONFIRMED

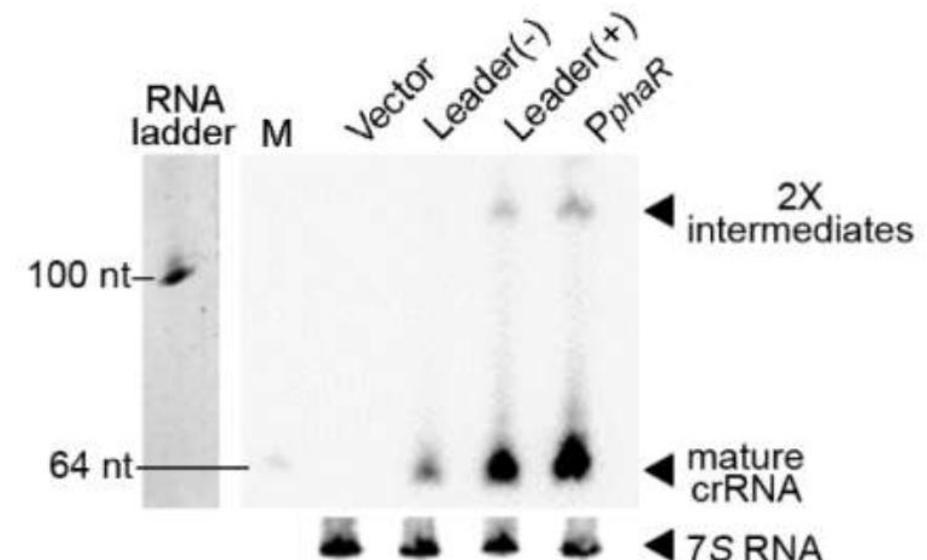
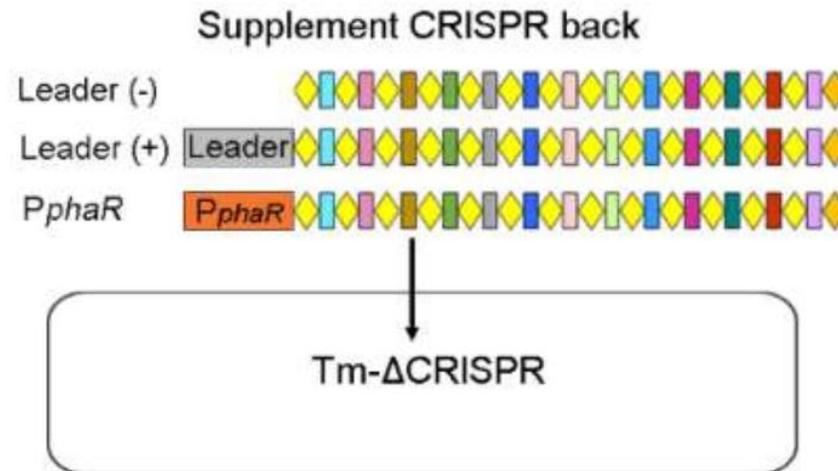
Does the shown result relay on CRISPR-array expression that leads crRNA transcript increase?

Model: Tm- Δ CRISPR + same plasmids

Methods: measure crRNA transcription by Northern Blot assay

Result: CRISPR-expressing cells DO produce significantly more crRNAs than the controls

CreA-guided gene repression can be alleviated by expanding the crRNA pool



5. Delving into the role of CreR in class 1 type I-E CRISPR system

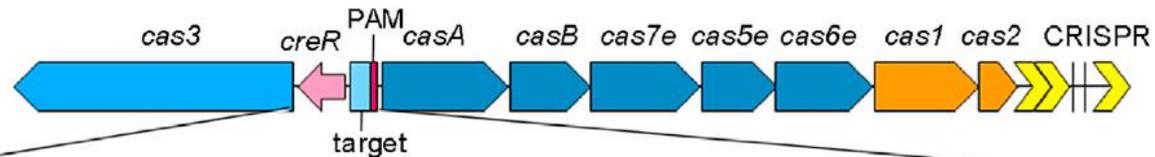
Testing cas repression effect of *Salmonella enterica* CreR (SeCreR) in cas-expressing *E. coli* MG1655 Δ hns strain

	3' handle	5' handle
<i>S. ente</i> Ψ R1	TACGTTCCCTAAAATTAT	GGGATAAACCG
<i>S. ente</i> Ψ R2	GCGTTCCTCCCGCGCCAGCG	GGGATAAACCTA
<i>S. ente</i> R	GTGTTCCCTCCCGCGCCAGCG	GGGATAAACCG
<i>E. coli</i> R	GAGTTCCTCCCGCGCCAGCG	GGGATAAACCG

High sequence similarity of the **CRISPR repeat** between *S. enterica* ATCC 51960 and *E. coli* MG1655

=> *E. coli* Cas probably recognises *S. enterica* CreR

Salmonella enterica
I-E CRISPR-Cas



```

CATCTTTTTCTGTAGCAGGGGGATTACAATGAAATCGATAATAAATTGGGATACGGGTAATTGGCCAGTTTTTTGTCTGGATTAGGTTATCCCCGCTG
GTA]GAAAAAGGACATCGTCCCCCTAATGTTACTTTAGCTATTATTTAACCCCTATGCCATTAAACCGGTCAAAAAACAGACCTAATCCAATAGGGGCGAC
cas3 start
GCGCGGGGAACGCTGCGCTATTAAGACTGTATCTACGGTTTTATCCATAATTTTAGGGAAACGTATTAAAGATAACAGATAACTATAAAAAATACATAGTC
CGCGCCCTTGGCACGCGATAATTCTGACATAGATGCCAAATAGGGTATTAAAATCCCTTGCATAATTTCTATTGTCTATTGATATTTTTTATGTATCAG
 $\Psi$ R2           $\Psi$ S           $\Psi$ R1
TGA]TTTAAGACTGTATCTACTTTTTATTTATATTTTCGTCTTCGAGAAAATGTAGTATAACTCCGATGACAGTATTTAAGAGATACCTCAGGACGGAACC
ACTAAATTCTGACATAGATGAA]AAATAAATATAAAAAGCAGAAGCTCTTTTACATCATATTGAGGCTACTGTCATAAATTCCTATGGAGTCCTGCCTTGG
casA start
TACCTATAGGTAGGTAATATTCCAAATTGGCTTAATAAATAGCCCTGCAGGAGTAAAGGTATG
ATGGATATCCATCCATTATAAGGTTTAACCGAATTATTTATCGGGACGTCCTCATTTCATAC
    
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7. Phage Acr proteins can relieve or subvert CreR-guided Cas autorepression

Phages have developed a range of Acr proteins to protect themselves against CRISPR-Cas

How does Cas autorepression respond to the action of Acr proteins?

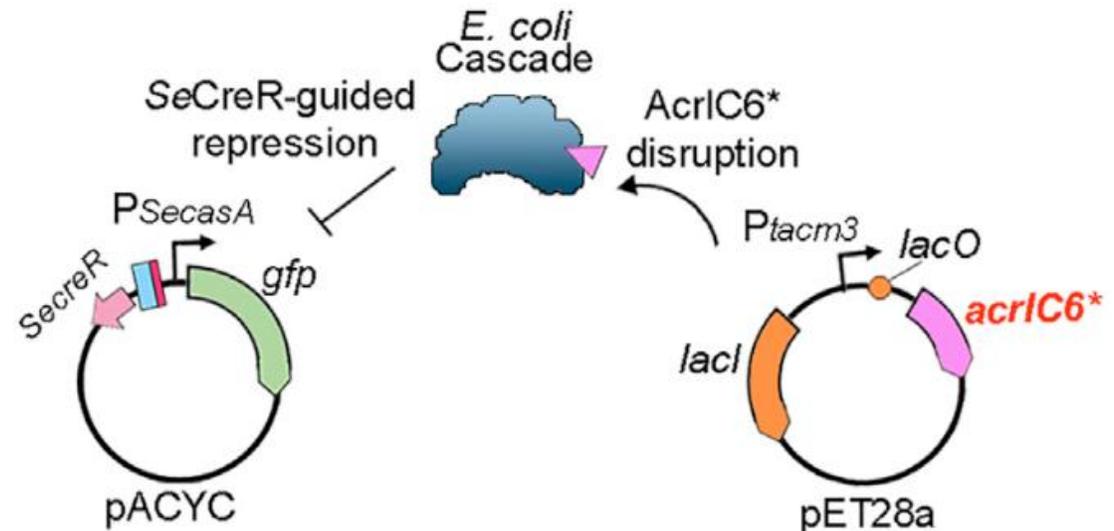
Looking at AcrIc6* targeting cas system I-E in *E. coli*

What's the effect of AcrIc6* on P_{secasA} known for being repressed by Cascade-CreR in *S. enterica*?

Model: Δhns *E. coli* cells + constructs:

- IPTG inducible promoter *tacm3* upstream *acrIc6**
- *S. enterica creR* & P_{secasA} -*gfp*

→ Treated with increasing dosage of **IPTG**



7. Phage Acr proteins can relieve or subvert CreR-guided Cas autorepression

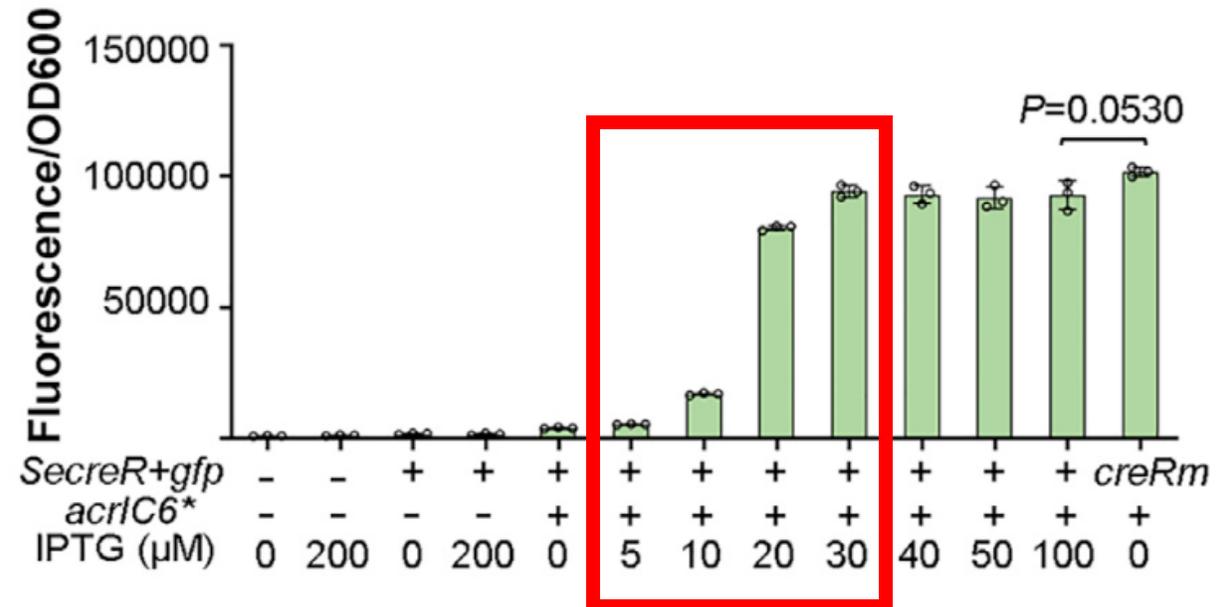
What's the effect of ArcIC6* on P_{secasA} known for being repressed by Cascade-CreR in *S. enterica*?

Methods: measure P_{secasA} activation detecting fluorescence by ***gfp*-reporter assay**

Result: induction of *arcIC6** with **5-30 μ M IPTG increases fluorescence** by a factor of 1.37-24.51, but **higher dosage does not result in further increases**

Even **low levels of ArcIC6*** can **relieve** or **subvert** the **repression effect** on P_{secasA}

→ I-E Cas autorepression circuit can respond to Acr, stimulating late mass production of new Cas effectors



7. Phage Acr proteins can relieve or subvert CreR-guided Cas autorepression

Studying ArcVA1-5 targeting cas system V-A in *M. bovoculi*



Conclusion: analogously to what demonstrated for I-E Cas autorepression circuit by AcrIC6*, **autoregulation circuit of MbCas12a responds to vary AcrVA proteins**