

# **“Rewriting Code of the Life”**

*G.B.N. Chainy*  
*30/8/19*  
*UU RC*

## ***Chinese Scientist Claims to Use Crispr to Make First Genetically Edited Babies***

The researcher, He Jiankui, offered no evidence or data to back up his assertions. If true, some fear the feat could open the door to “designer babies.” **2018**

Corn and other important crops can now be gene edited by pollen carrying CRISPR

By [Jon Cohen](#) Mar. 4, 2019 , 11:00 AM



CRISPR-Cas9 Gene Editing in Lizards through Microinjection of Unfertilized Oocytes



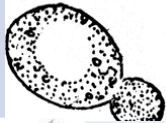


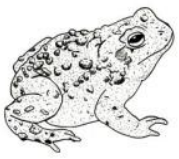


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# WHAT IS CRISPR- Cas9 ?

(Clustered Regularly Interspaced Short Palindromic Repeats/CRISPR-associated  
nuclease 9)

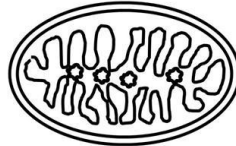
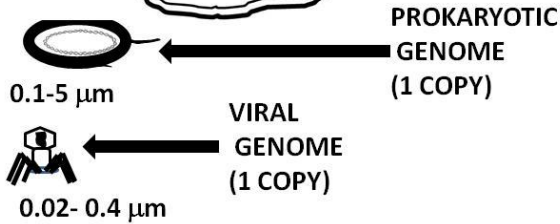
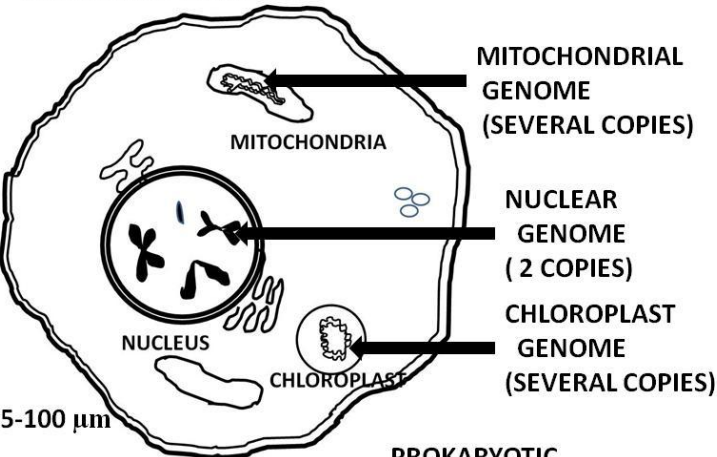
# What is a genome?

A genome is an organism's complete set of DNA, including all of its genes. Each genome contains all of the information needed to build and maintain that organism.

SPECIES	CHROMOSOME NUMBER	C VALUE DNA (pg)/COPY GENOME	GENOME SIZE(Mbp)
	1		
	1	0.017	4.24
	14	0.17	12.1
	24	0.87	430
	42	15.46	16000
	22	6	
	40	3.3	2800
	46	3.5	3200

# EUKARYOTIC GENOME

1 base pair(bp) = 660 Dalton  
 1000 bp = 1 Kilobase ( $10^3$ bp)  
 1000 kb = 1 Megabase ( $10^6$ bp)  
 1000 mb = 1 Gigabase ( $10^9$ bp)

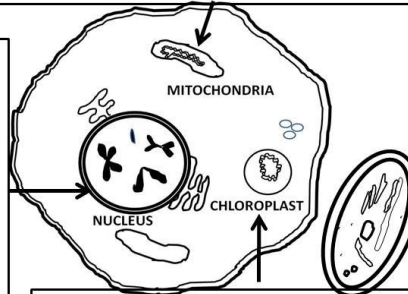


**MITOCHONDRIAL GENOME:**

- CIRCULAR, DUPLEX, MULTIPLE COPIES
- > 100 COPIES IN DIFFERENTIATING EMBRYONIC CELLS.
- SIZE 16000-20000 bp IN ANIMALS.
- SIZE 200,000 TO 250,000 IN PLANTS.

**NUCLEAR GENOME:**

- LARGE SIZE, DUPLEX, LINEAR, COMPLEX ORGANIZATION.
- MORE THAN ONE DNA MOLECULE.
- MOSTLY DIPLOID STATE
- PRESENCE OF NON CODING SEQUENCE.
- PRESENCE OF REPETITIVE DNA.
- SIZE  $10^6$  TO  $10^{11}$  bp



**CHLOROPLAST GENOME**

- CIRCULAR, DUPLEX, MULTIPLE COPIES.
- > 100 COPIES IN DIFFERENTIATING EMBRYONIC CELLS.
- SIZE 120000-160000 bp.

**A TYPICAL PROKARYOTIC GENOME**

SMALL SIZE ( $10^5$ - $10^6$  bp)

DUPLEX

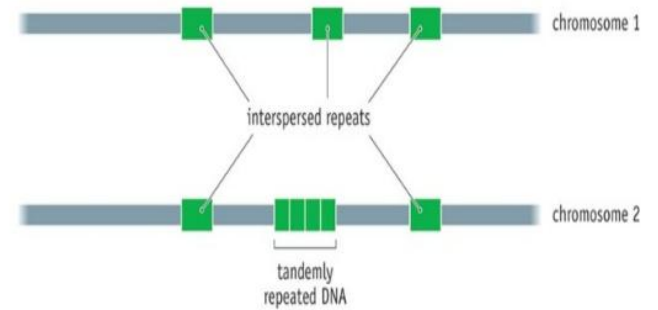
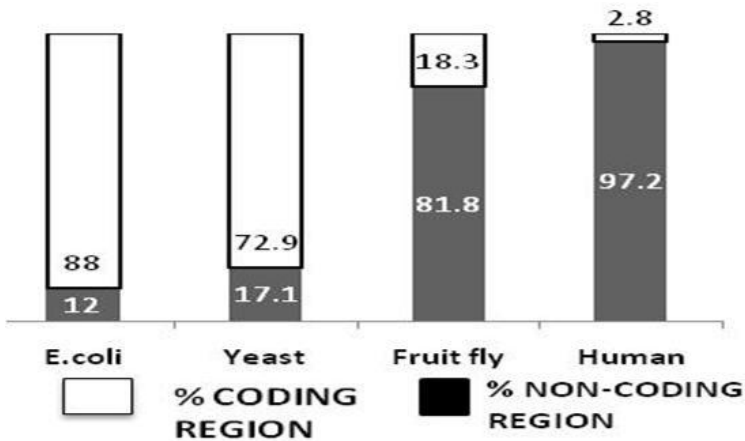
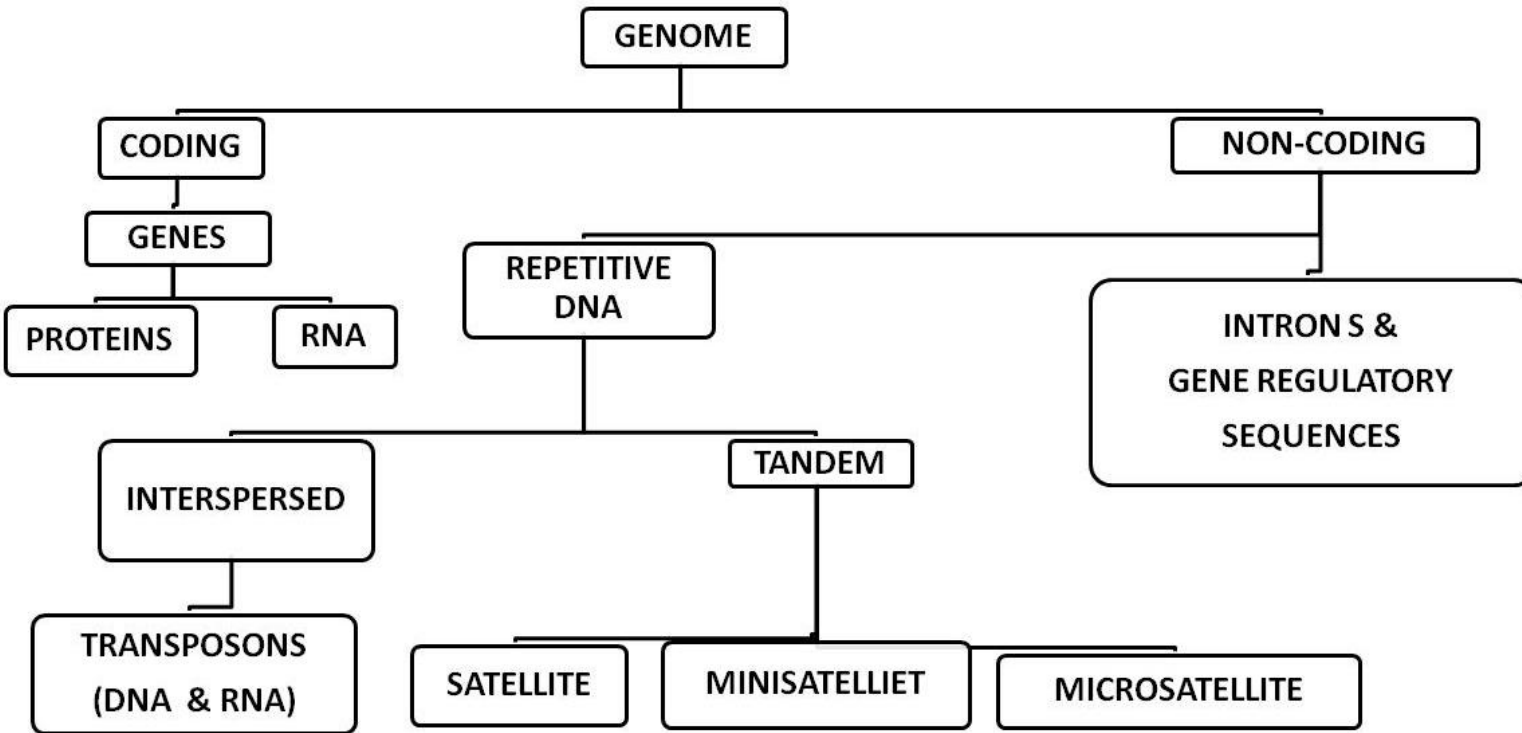
PRESENT FREELY IN CYTOPLASM

SINGLE COPY, COMPACT

NO REPETITIVE SEQUENCE

SINGLE ORIGIN OF REPLICATION

# SCHEMATIC PRESENTATION OF GENOME ORGANIZATION



CTAGAGATAGATAGATAGATAGATAGATAGATAGATACTAGAC

# **DISCOVERY OF CRISPR-Cas9**





- The park is about 3,700 hectares..
- The pink lake is 1,400 hectares
- the green one is 700 hectares.
- Both are connected to the sea by canals.

- The strange pink-purple colour of the Torrevieja lagoon is caused by pigments of the **Halobacterium** bacteria and by an alga called **Dunaliella salina**, which is responsible for the bright red colour of the lake.
- The Artemia Salina brine shrimp, which lives in the lake, is also red because it feeds on the bacteria.

### **1993 - 2005 — Francisco Mojica, University of Alicante, Spain**

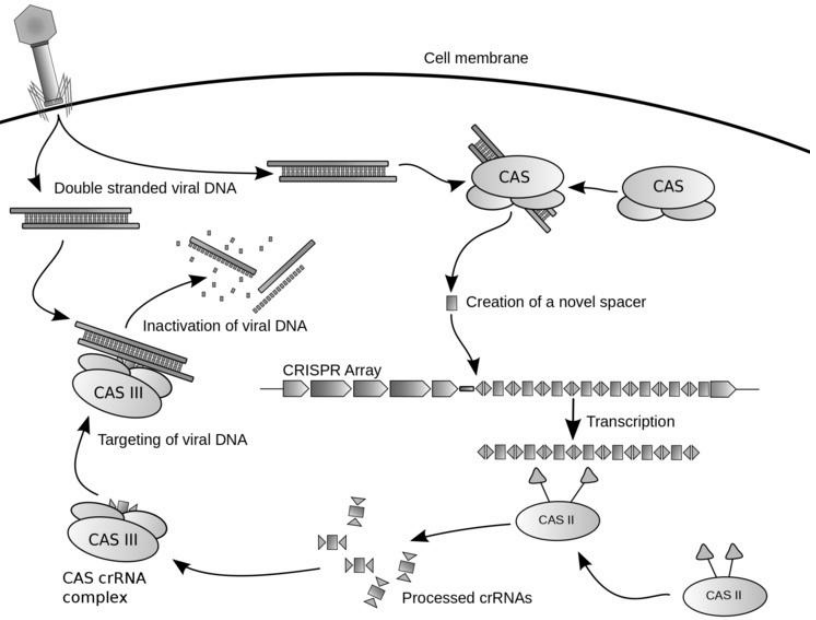
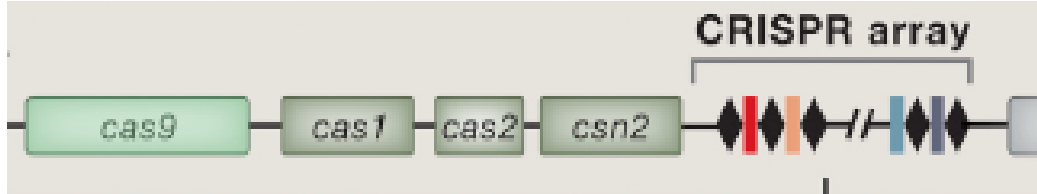
- Characterized CRISPR locus in 1993.
- Coined the term CRISPR through correspondence with Ruud Jansen 2002.
- CRISPR snippets from the genomes of bacteriophage .
- hypothesized, correctly, that CRISPR is an adaptive immune system.



# Discovery of Cas9 and PAM

May, 2005 — Alexander Bolotin, French National Institute for Agricultural Research (INRA)

- Noticed unusual CRISPR locus in *Streptococcus thermophilis*.
- Found **Cas 9 gene** encoding large Cas 9 nuclease.
- Furthermore, they noted that the all spacers share a common sequence at one end the **protospacer adjacent motif (PAM)**,



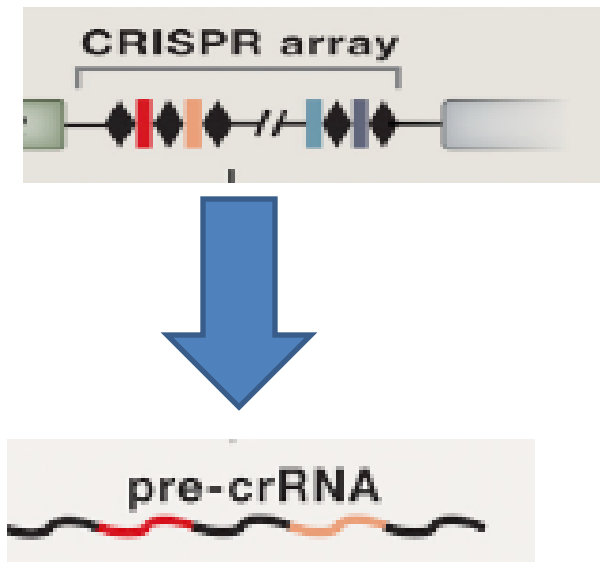
# Hypothetical scheme of adaptive immunity

March, 2006 — Eugene Koonin, US National Center for Biotechnology Information, NIH

Koonin was studying clusters of orthologous groups of proteins by computational analysis and proposed a hypothetical scheme for CRISPR cascades as bacterial immune system based on inserts homologous to phage DNA in the natural spacer array, abandoning previous hypothesis that the Cas proteins might comprise a novel DNA repair system.

**Experimental demonstration of adaptive immunity.** March, 2007 — Philippe Horvath, Danisco France SAS

- *S. thermophilus* is widely used in the dairy industry to make yogurt and cheese.
- showed experimentally that CRISPR systems are indeed an adaptive immune system
- showed that Cas9 is likely the only protein required for interference, the process by which the CRISPR system inactivates invading phage, details of which were not yet known.



**Spacer sequences are transcribed into guide RNAs**

August, 2008 — John van der Oost, University of Wageningen, Netherlands

- showed that in *Escherichia coli*, spacer sequences, which are derived from phage, are transcribed into small RNAs, termed CRISPR RNAs (crRNAs), that guide Cas proteins to the target DNA.

## **CRISPR acts on DNA targets**

*December, 2008 — Luciano Marraffini and Erik Sontheimer, Northwestern University, Illinois*

- Demonstrated that the **target molecule is DNA, not RNA**. This was somewhat surprising, as many people had considered CRISPR to be a parallel to eukaryotic RNAi silencing mechanisms, which target RNA.

## **Cas9 cleaves target DNA**

*December, 2010 — Sylvain Moineau, University of Laval, Quebec City, Canada*

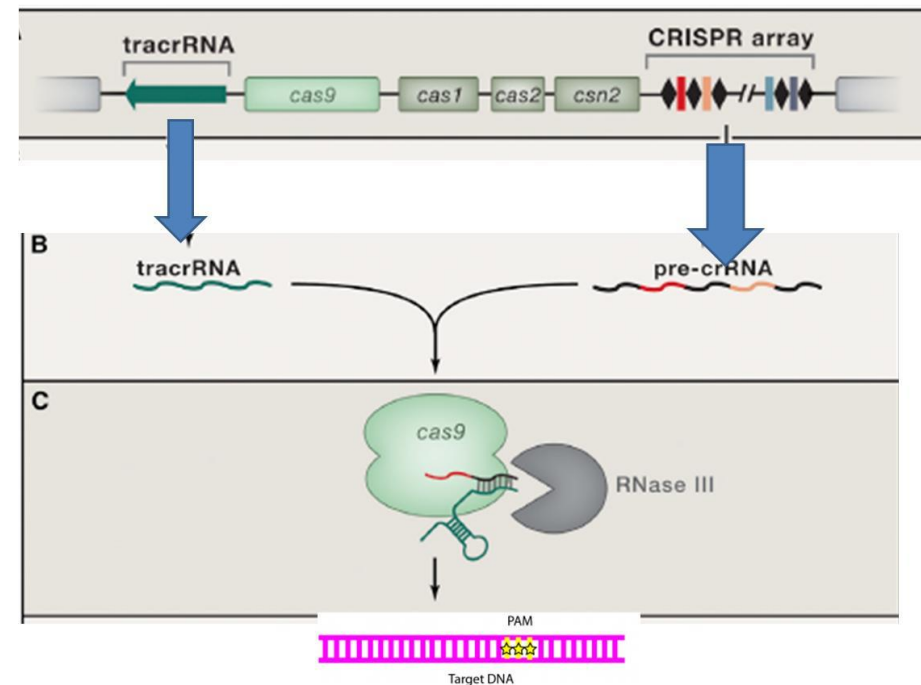
- Demonstrated that **CRISPR-Cas9 creates double-stranded breaks in target DNA at precise positions, 3 nucleotides upstream of the PAM**.
- Confirmed that Cas9 is the only protein required for cleavage in the CRISPR-Cas9 system.

**CRISPR is an adaptive immune system.**

## Discovery of tracrRNA for Cas9 system

*March, 2011 — Emmanuelle Charpentier, Umea University, Sweden and University of Vienna, Austria*

- Discovered that in addition to the crRNA, a second small RNA exists.
- Named it **trans-activating CRISPR RNA (tracrRNA)**.
- Demonstrated that **tracrRNA forms a duplex with crRNA**, and that it is this duplex that guides Cas9 to its targets.

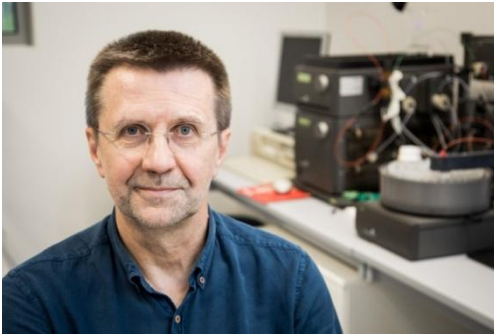


**CRISPR-Cas9 :  
FROM FUNDAMENTALS TO APPLICATION**

## CRISPR systems can function heterologously in other species

July, 2011 — Virginijus Siksnys, Vilnius University, Lithuania

- Cloned the entire CRISPR-Cas locus from *S. thermophilus* (a Type II system) and expressed it in *E. coli* (which does not contain a Type II system).

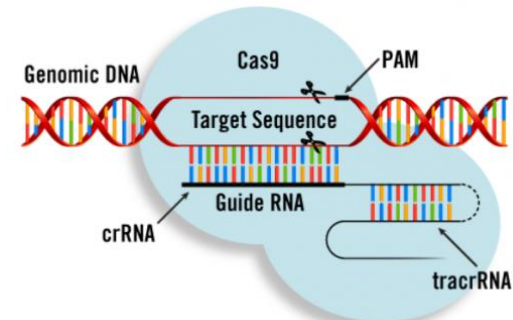
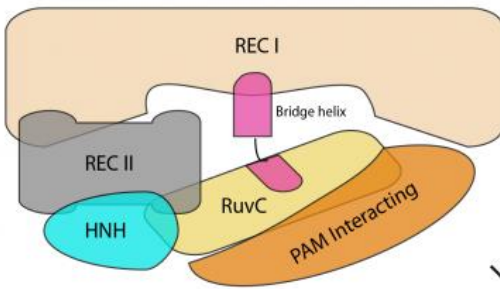


## Biochemical characterization of Cas9-mediated cleavage

September, 2012 — , Vilnius University, Lithuania

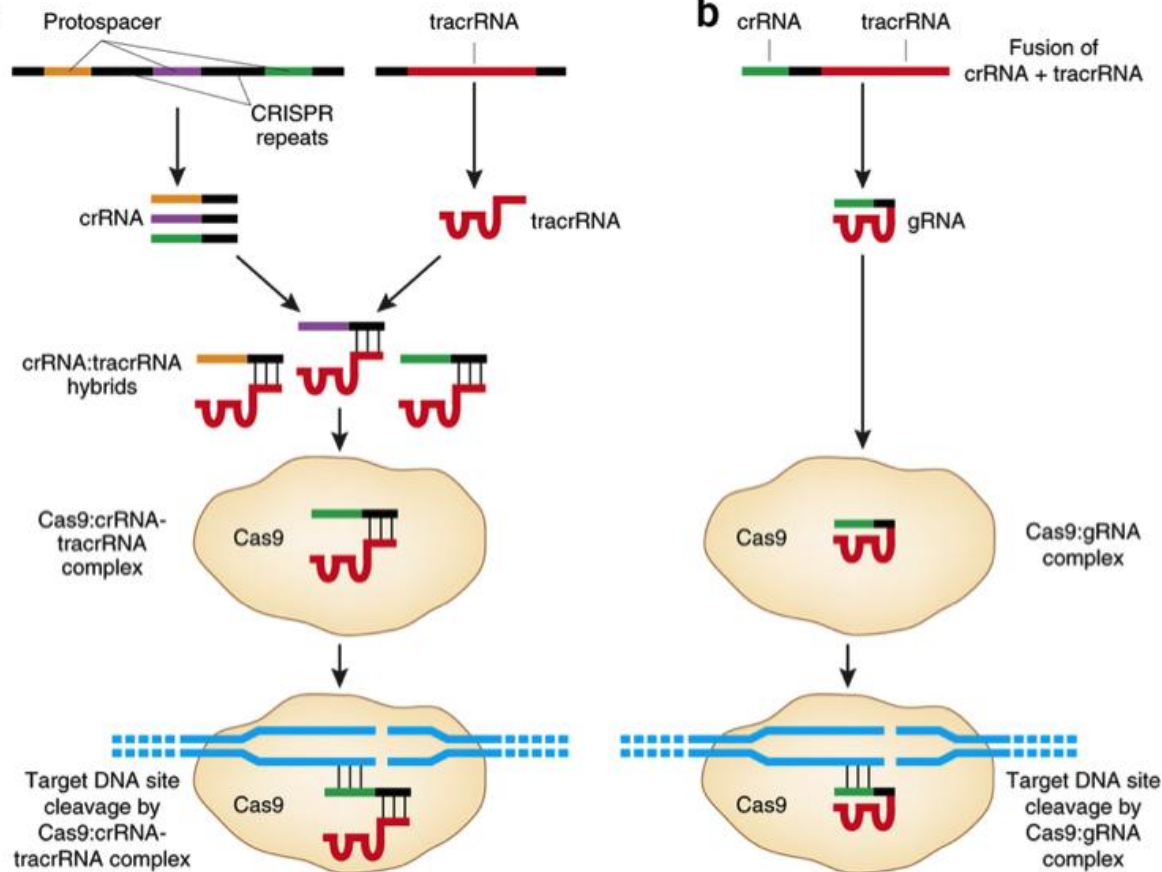
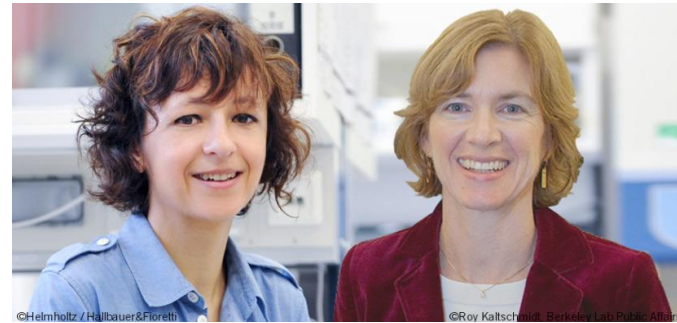
- Purified Cas9 in complex with crRNA from the *E. coli* strain engineered to carry the *S. thermophilus* CRISPR locus.
- Mechanistically characterize Cas9's mode of action .
- Verified the cleavage site and the requirement for the PAM.
- RuvC domain cleaves the non-complementary strand while the HNH domain cleaves the complementary site.
- crRNA could be trimmed down to a 20-nt stretch sufficient for efficient cleavage.
- Reprogramming of Cas9 to target a site by changing the sequence of the crRNA.

Cas9 Complex (Inactive)



June, 2012 — **Charpentier and Jennifer Doudna**, University of California, Berkeley

- Reported that the crRNA and the tracrRNA could be fused together to create a single, synthetic guide, further simplifying the system



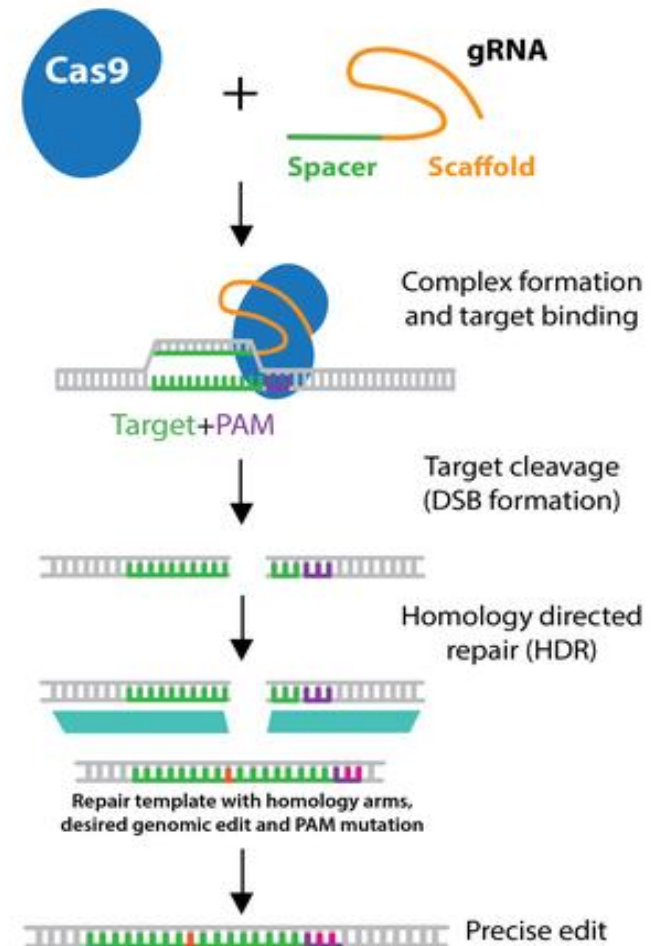


# CRISPR-Cas9 harnessed for genome editing

January, 2013 — Feng Zhang, Broad Institute of MIT and Harvard, McGovern Institute for Brain Research at MIT, Massachusetts



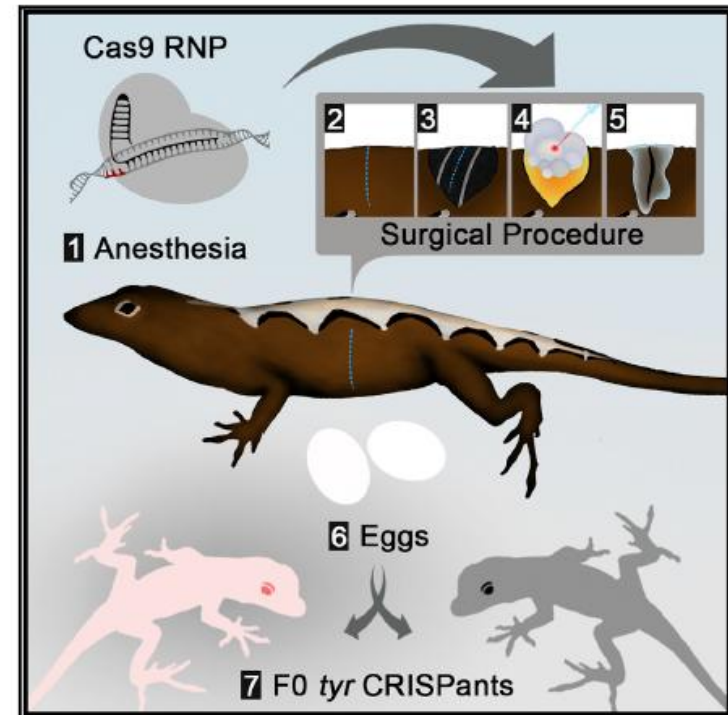
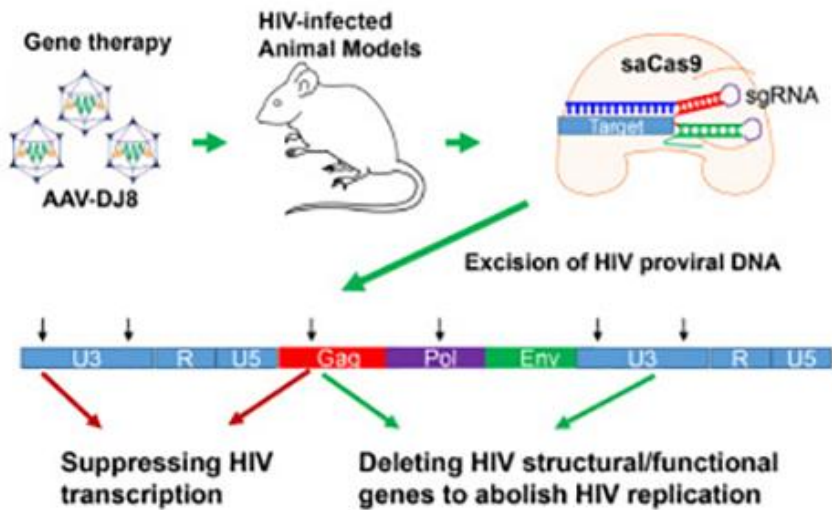
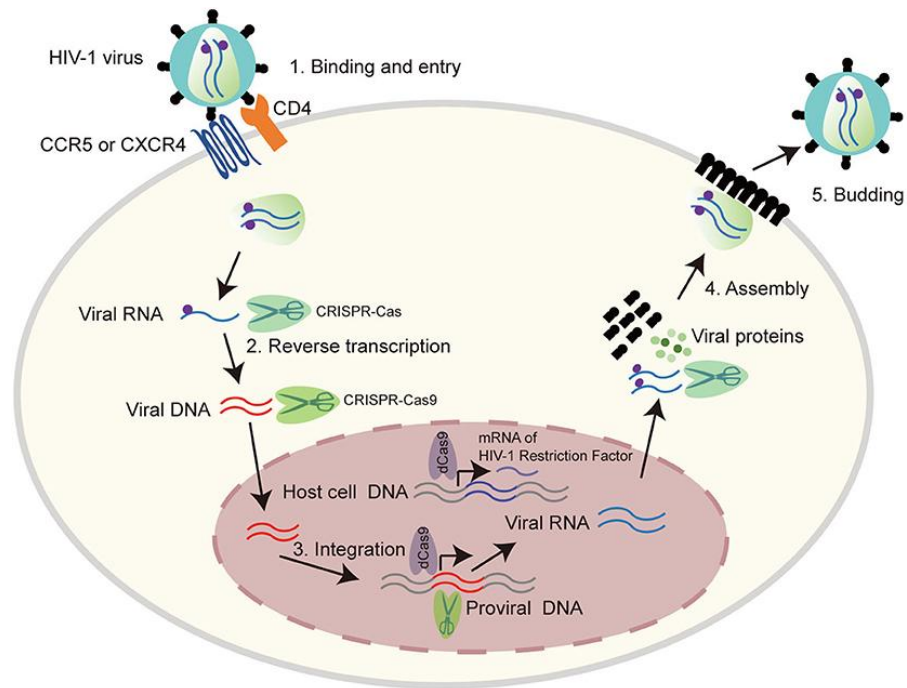
- Engineered two different Cas9 orthologs (from *S. thermophilus* and *S. pyogenes*) and demonstrated **targeted genome cleavage in human and mouse cells.**
- showed that the system (i) could be programmed to target multiple genomic loci, and (ii) could drive homology-directed repair.



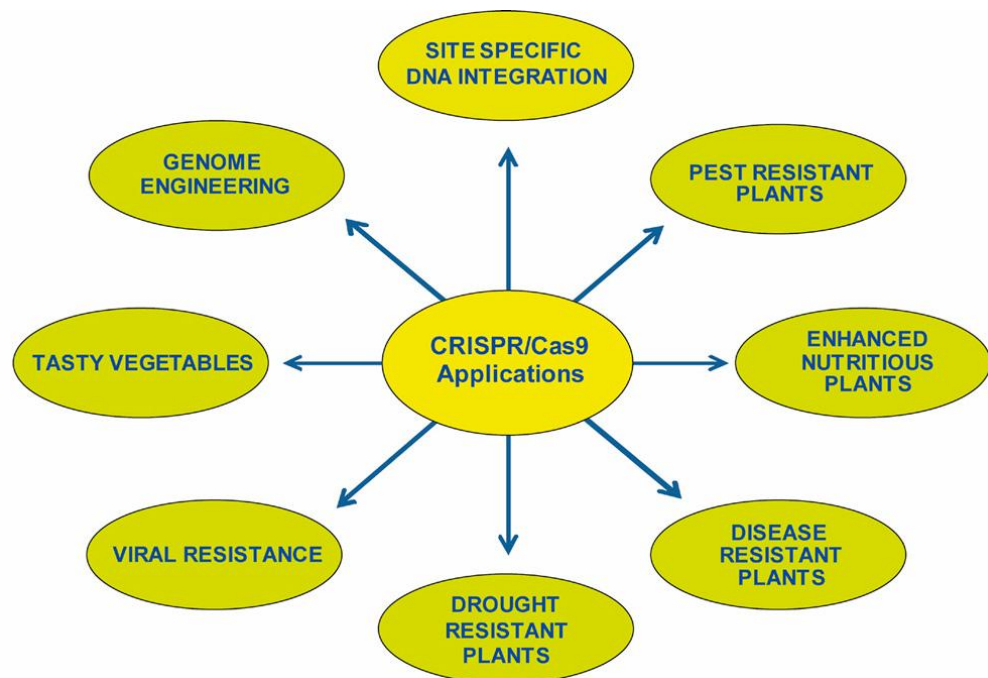
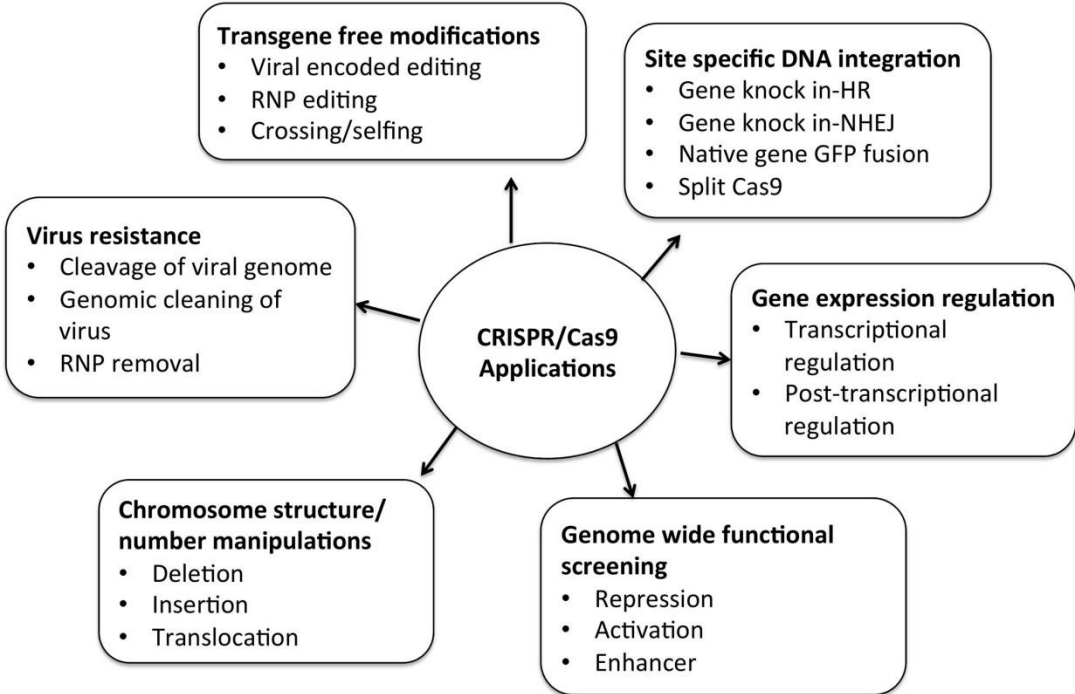
[Nature Protocol.](#) 2013 Nov;8(11):2281-2308

**Genome engineering using the CRISPR-Cas9 system.**

Ran FA, Hsu PD, Wright J, Agarwala V, Scott D, Zhang F.



Highlights



**GENOME**

**READ**

**COPY**

**and now u can REWRITE ....**



**THANK U ALL**