



# The CRISPR Revolution: From Reading to Editing Genomes

One Health Research Symposium  
Kansas City  
August 19, 2018

**Kevin Davies PhD**  
Executive Editor,  
*The CRISPR Journal*





# CRISPR



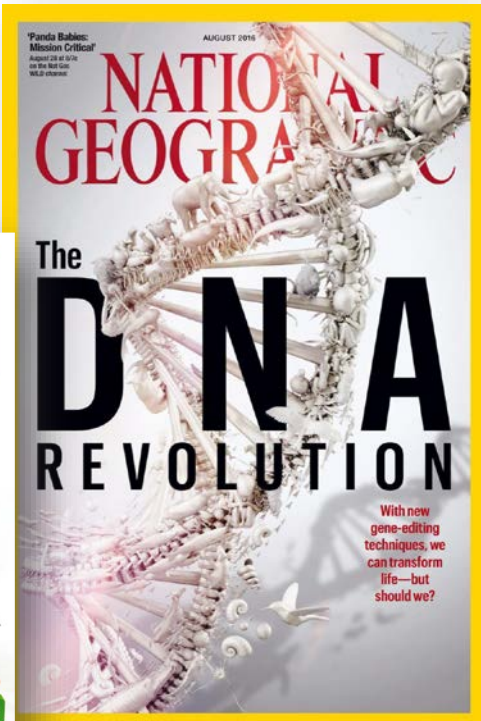
**The Economist**  
AUGUST 22ND - 29TH 2015  
economist.com

How Russians cope with recession  
No-go for NGOs in China  
Islamic State's taste for slavery  
Commodities: the binge, the hangover  
India's poet-politicians

## Editing humanity

The prospect of genetic enhancement

No baldness  
Sprinter  
Perfect pitch  
High IQ  
Low risk of Alzheimer's, breast cancer and strokes  
20/20 vision



SEPTEMBER 7, 2015  
**C&EN**  
CHEMICAL & ENGINEERING NEWS

**DISRUPTING PRIONS**  
Small molecules target progressive diseases P.37  
**BOSTON IN PHOTOS**  
Sights from the fall 2015 ACS national meeting P.48

## EDITING THE GENOME

CRISPR/Cas9 takes gene manipulation into a new era P.14  
PUBLISHED BY THE AMERICAN CHEMICAL SOCIETY



The risks of using CRISPR to edit embryos



60  overtime  
MINUTES

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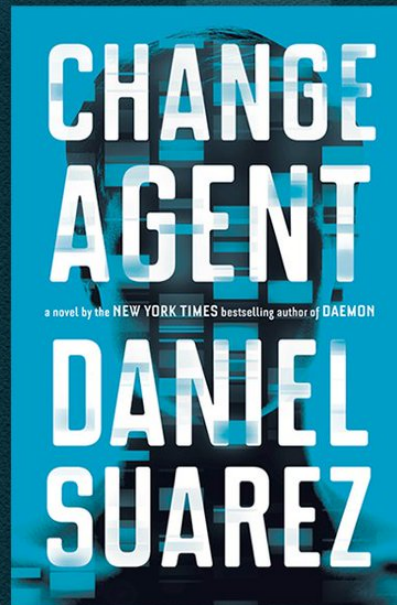
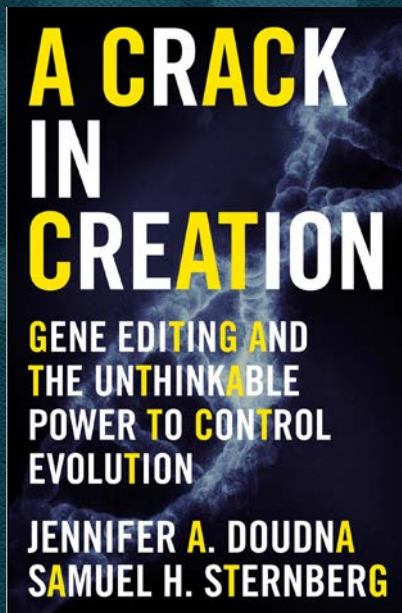




**C**RUNCHY  
**R**ECTUMS  
**I**N  
**S**ASSY  
**P**INK  
**R**AY-BANS



# CRISPR in the Public Eye



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OCTOBER 18, 2016 2:31pm PT by Lesley Goldberg

## Jennifer Lopez Sets Futuristic Bio-Terror Drama at NBC (Exclusive)

Amanda Edwards/WireImage  
Jennifer Lopez

'C.R.I.S.P.R.' — aka "clustered regularly interspaced short palindromic repeats" — marks her latest project for the network.

NBC is reteaming with Jennifer Lopez for a futuristic procedural.

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**EMMY WORTHY CAST**  
— MOVIE PILOT



BIG  
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BIGGER

*"Are you familiar with CRISPR?"*

DWAYNE JOHNSON

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We've gone from reading DNA...







genome.gov

National Human Genome Research Institute

April 14, 2003

## Newsroom

News Releases ▶ International Consortium Completes Human Genome Project



National Human Genome Research Institute  
National Institutes of Health  
Department of Health and Human Services  
and  
Office of Science  
U.S. Department of Energy

### International Consortium Completes Human Genome Project

#### *All Goals Achieved; New Vision for Genome Research Unveiled*

**BETHESDA, Md.,** April 14, 2003 - The International Human Genome Sequencing Consortium, led in the United States by the National Human Genome Research Institute (NHGRI) and the Department of Energy (DOE), today announced the successful completion of the Human Genome Project more than two years ahead of schedule.

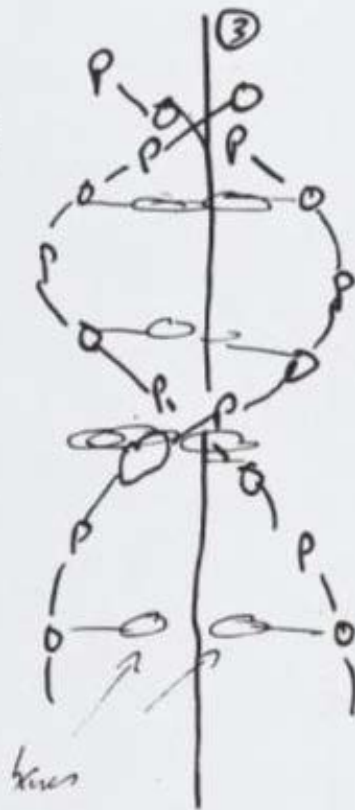
Also today, NHGRI unveiled its bold new vision for the future of genome research, officially ushering in the era of the genome. The vision will be published in the April 24 issue of the journal *Nature*, coinciding with the 50th anniversary of *Nature's* publication of the landmark paper by Nobel Laureates James Watson and Francis Crick that described DNA's double helix. Dr. Watson also was the first leader of the [Human Genome Project](#).

The international effort to sequence the 3 billion DNA letters in the human genome is considered by many to be one of the most ambitious scientific undertakings of all time, even compared to splitting the atom or going to the moon.

"The Human Genome Project has been an amazing adventure into ourselves, to understand our own DNA instruction book, the shared inheritance of all humankind," said NHGRI Director Francis S. Collins, M.D., Ph.D., leader of the Human Genome Project since 1993. "All of the project's goals have been completed successfully - well in advance of the original deadline and for a cost substantially less than the original estimates."

“My

like this



19 March 1953

Jim Watson and I have probably made a most important discovery...

Our structure is very beautiful. D.N.A. can be thought of roughly as a very long chain with flat bits sticking out. The flat bits are called the 'bases'...

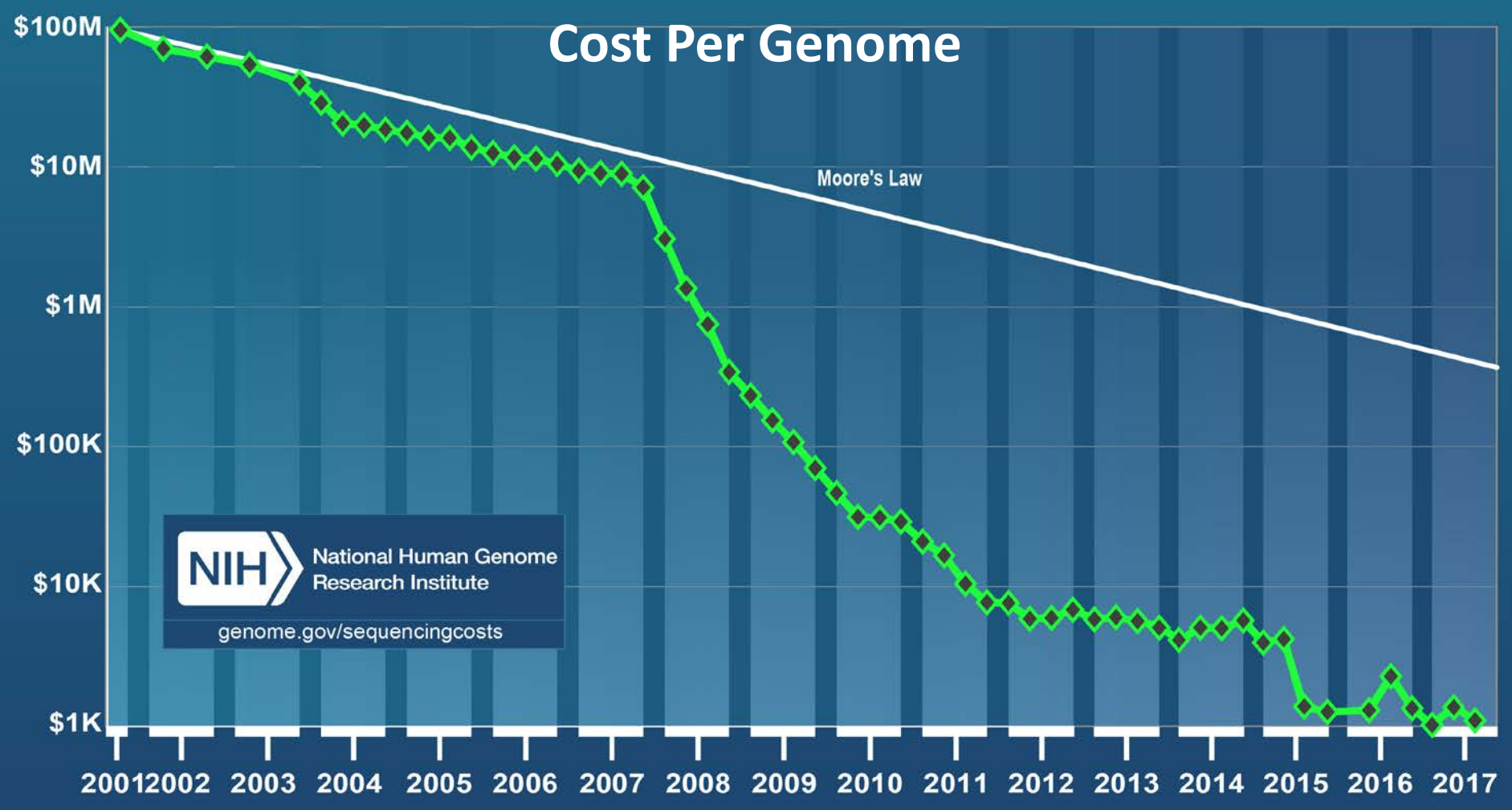
... Lots of love, Daddy”





Cas Kramer (Univ Leicester)

# Cost Per Genome



**NIH** National Human Genome Research Institute  
[genome.gov/sequencingcosts](http://genome.gov/sequencingcosts)



## “The Chemists’ Pub”



**Sir Shankar Balasubramanian & David Klenerman**

*The Panton Arms / University of Cambridge, UK*

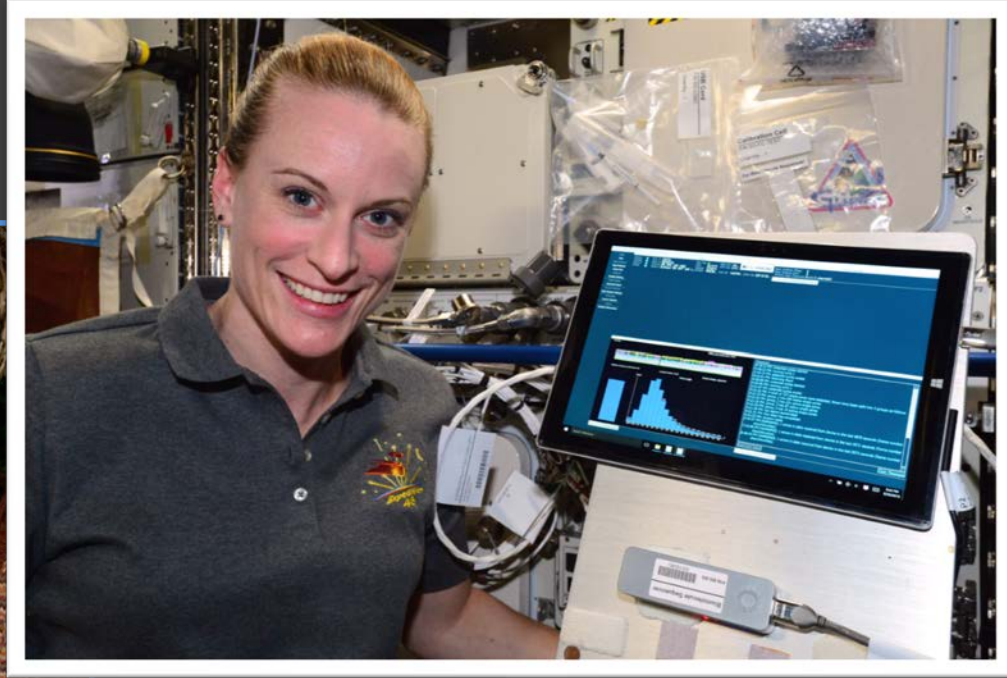
## “Project Jim”



Houston  
31 May 2007



# The Portable Sequencer



Quick et al. *Nature* February 2016



... to writing DNA...



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GP-write

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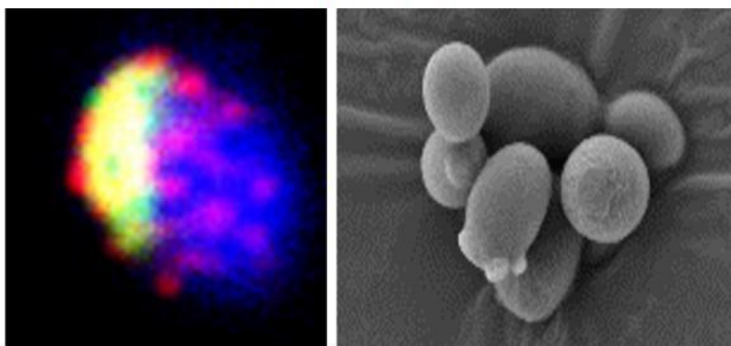
## Introducing GP-write: A Grand Challenge

The Genome Project-write (GP-write) is an open, international research project led by a multi-disciplinary group of scientific leaders who will oversee a reduction in the costs of engineering and testing large genomes in cell lines more than 1,000-fold within ten years.

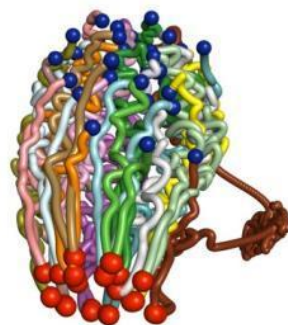
[Download the GP-write White Paper](#)

[Submit a Pilot](#)

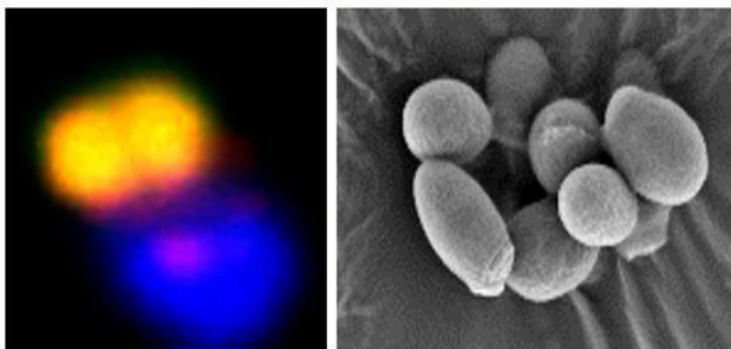
## 天然酿酒酵母



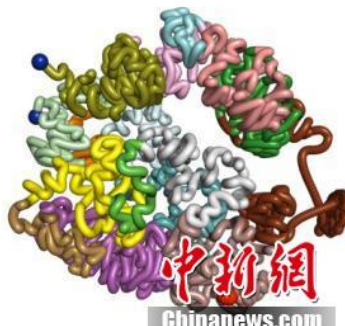
## 16条染色体



## 人造酿酒酵母



## 1条巨大染色体



Chinanews.com

## MINIMAL YEAST

Two separate teams have managed to pack nearly all of yeast's genetic material — usually spread over 16 chromosomes — into just one and two, respectively.



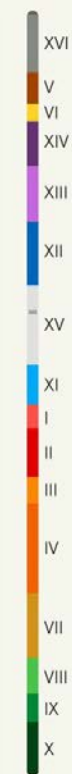
### TWO CHROMOSOMES

A US team used CRISPR to remove genetic material at the ends and middle of chromosomes, known as telomeres and centromeres, respectively, and relied on yeast's natural DNA-repair mechanisms to produce a strain with just two chromosomes.



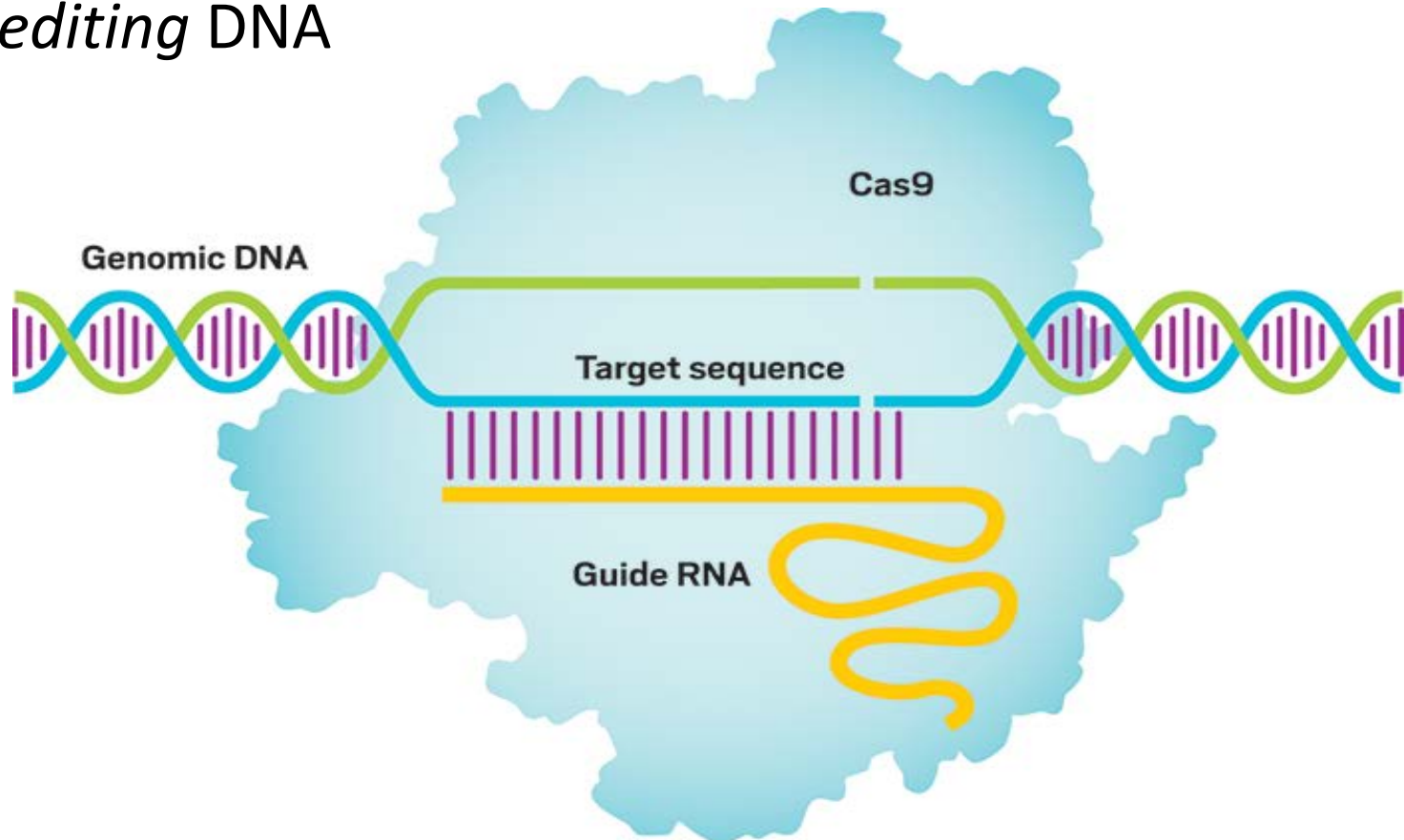
### ONE CHROMOSOME

A team in China used CRISPR to remove 15 centromeres and 30 telomeres, in addition to 19 long, repeated sequences.



©nature

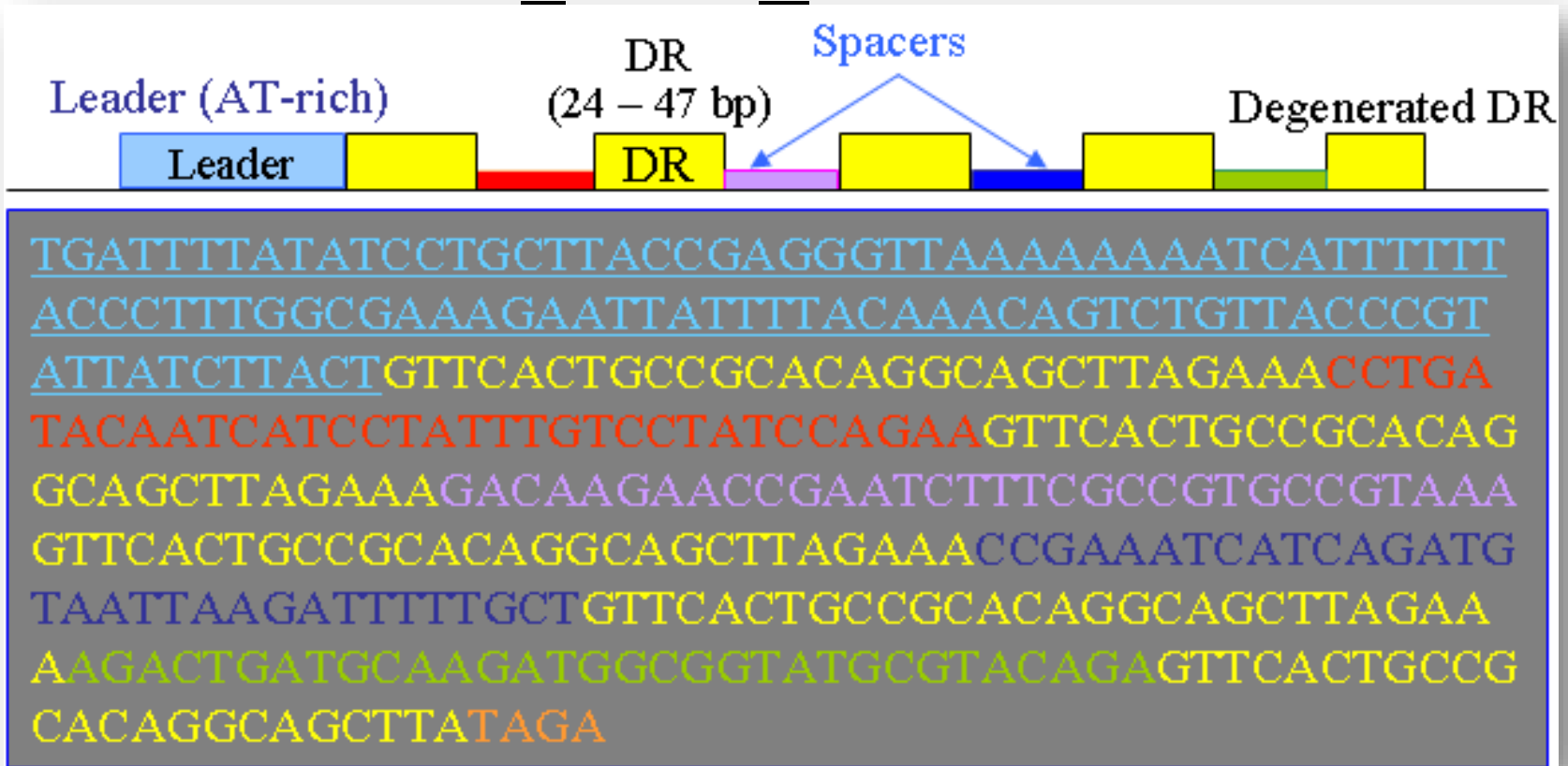
... to *editing* DNA



CRISPR-Cas9: PRECISION EDITING OF  
DNA



# CRISPR: Clustered Regularly Interspaced Short Palindromic



Francisco Mojica  
Salt Lakes of Santa Pola



**Asunto: Re: Acronym**

**Fecha:** Wed, 21 Nov 2001 16:39:06 +0100

**De:** "Ruud Jansen" <R.Jansen@vet.uu.nl>

**Empresa:** Diergeneeskunde

**A:** "Francisco J. Martínez Mojica" <fmojica@ua.es>

Dear Francis

What a great acronym is CRISPR.

I feel that every letter that was removed in the alternatives made it less crispy so I prefer the snappy CRISPR over SRSR and SPIDR. Also not unimportant is the fact that in MedLine CRISPR is a unique entry, which is not true for some of the other shorter acronyms.

---

JOURNAL OF **MOLECULAR  
EVOLUTION**

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## **Intervening Sequences of Regularly Spaced Prokaryotic Repeats Derive from Foreign Genetic Elements**

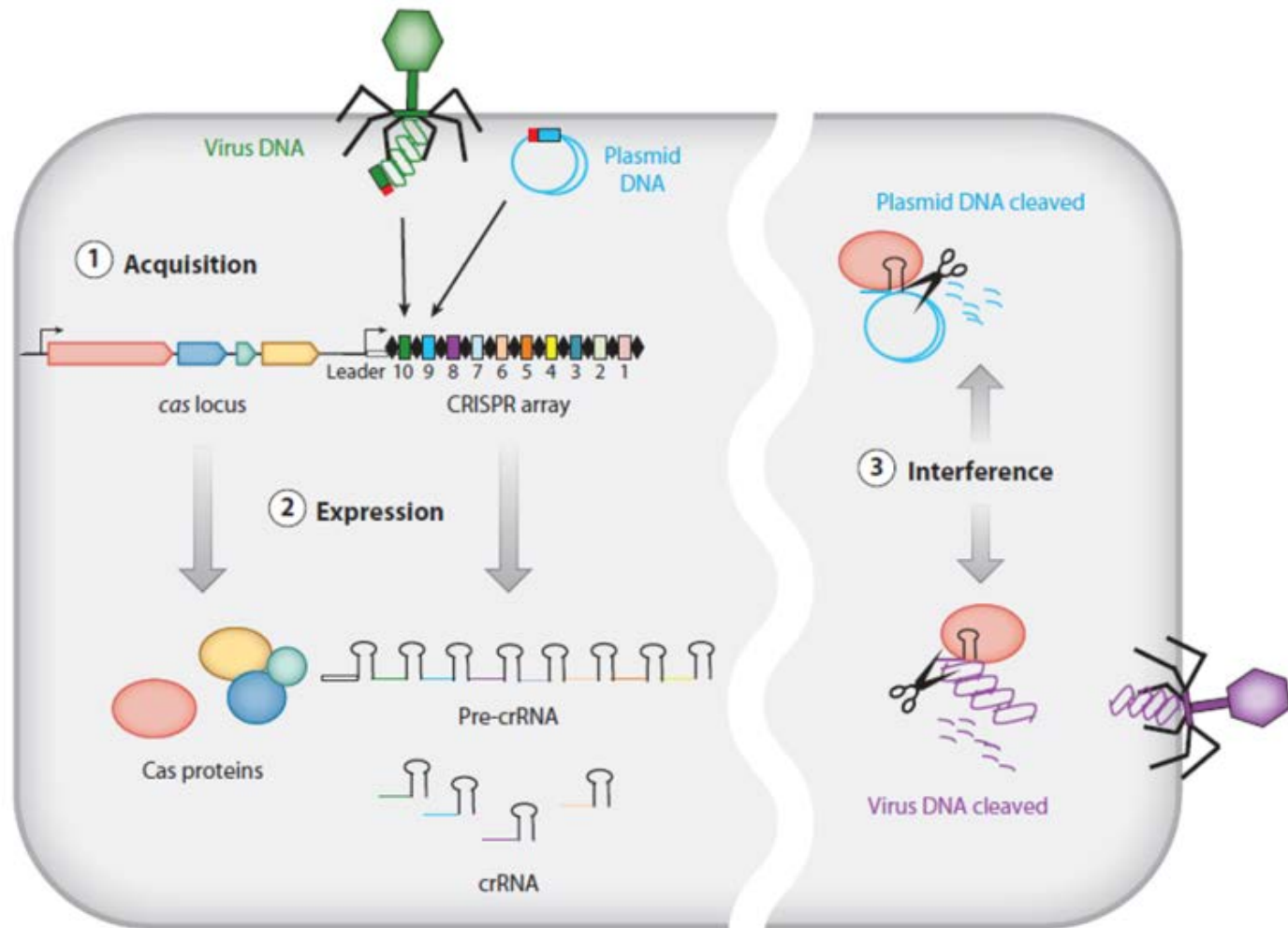
**Francisco J.M. Mojica, César Díez-Villaseñor, Jesús García-Martínez, Elena Soria**

División de Microbiología, Departamento de Fisiología, Genética y Microbiología, Universidad de Alicante, Campus de San Vicente, E-03080, Spain

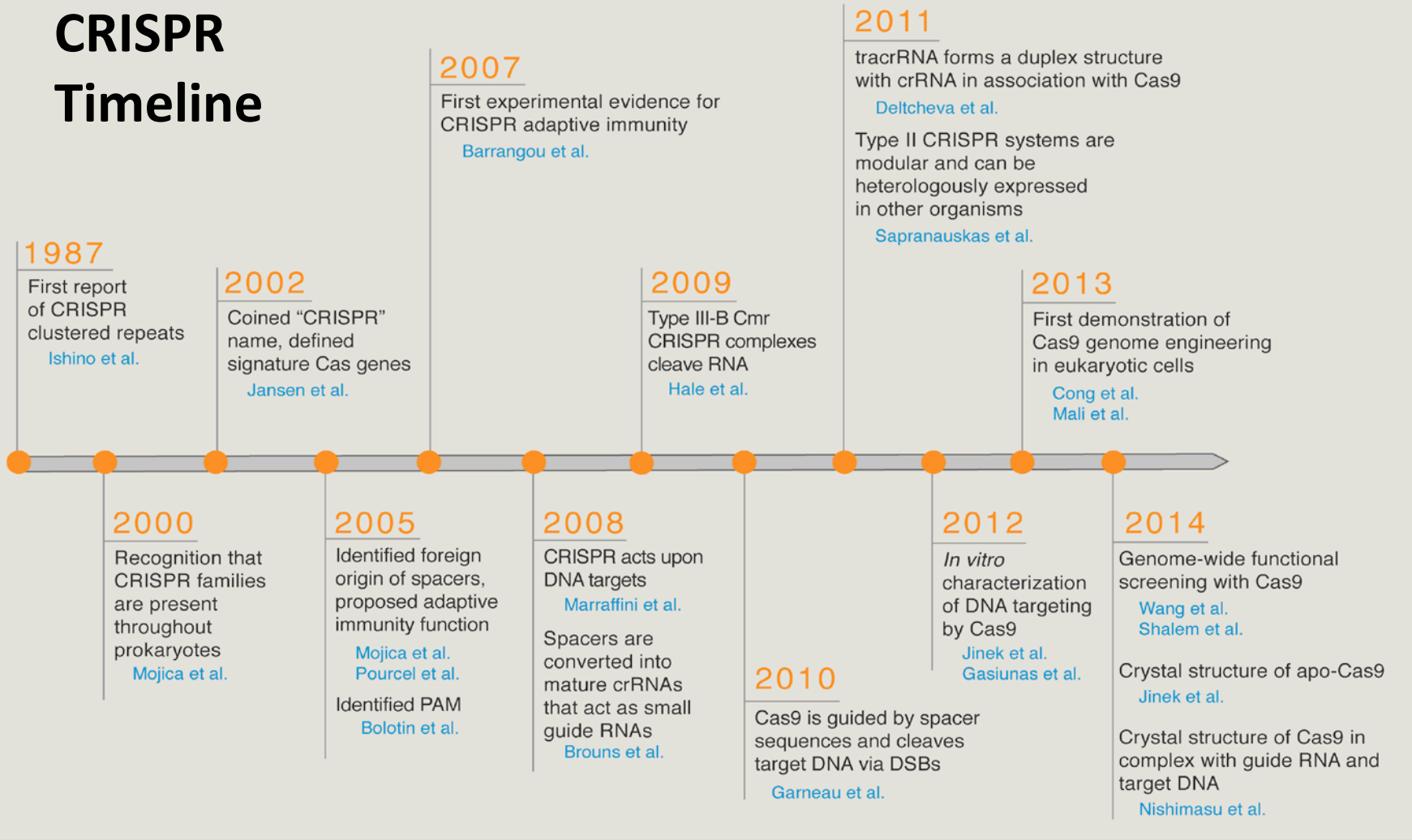
Received: 6 February 2004 / Accepted: 1 October 2004 [*Reviewing Editor:* Dr. John Huelsenbeck]



**CRISPR** is a naturally occurring, bacterial immune defense system that provides a means to recognize, remember and destroy viral invaders



# CRISPR Timeline



# Heroes of CRISPR

Ishino



Mojica



Koonin



Barrangou



Horvath



Makarova



Chylinski



Fremaux



Moineau



Marraffini



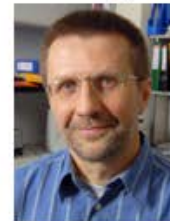
Sontheimer



Gasiunas



Siksnys



Jinek



Cong



Mali



Charpentier



Doudna



Zhang



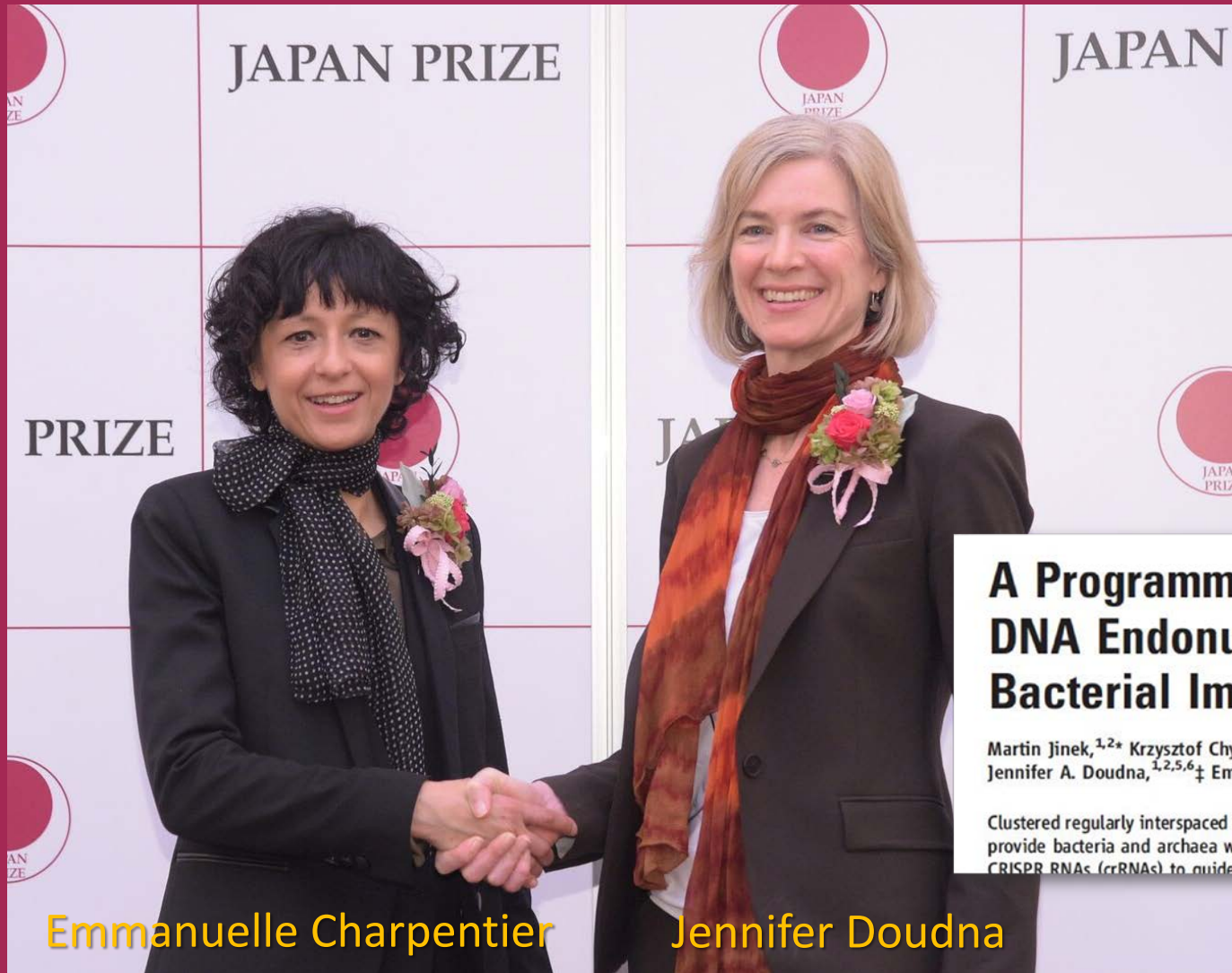
Garneau



Church







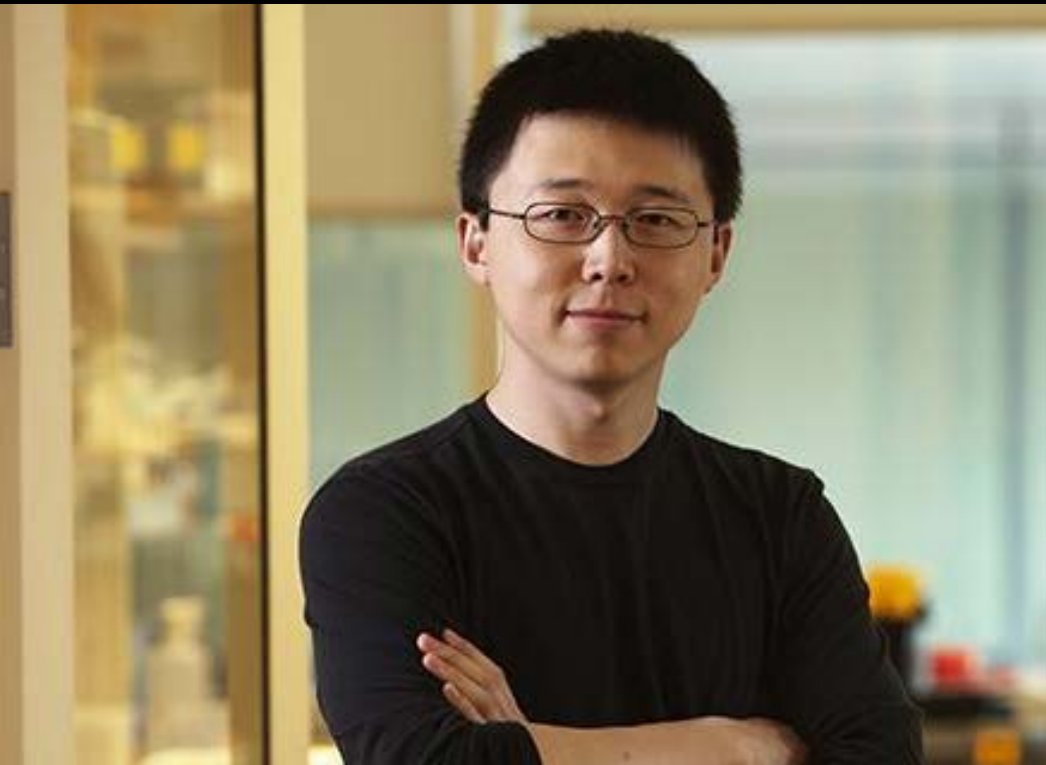
Emmanuelle Charpentier

Jennifer Doudna

## A Programmable Dual-RNA–Guided DNA Endonuclease in Adaptive Bacterial Immunity

Martin Jinek,<sup>1,2\*</sup> Krzysztof Chylinski,<sup>3,4\*</sup> Ines Fonfara,<sup>4</sup> Michael Hauer,<sup>2,†</sup> Jennifer A. Doudna,<sup>1,2,5,6,‡</sup> Emmanuelle Charpentier<sup>4,‡</sup>

Clustered regularly interspaced short palindromic repeats (CRISPR)/CRISPR-associated (Cas) systems provide bacteria and archaea with adaptive immunity against viruses and plasmids by using CRISPR RNAs (crRNAs) to guide the silencing of invading nucleic acids. We show here that in



Feng Zhang (MIT/Broad Institute)

## Multiplex Genome Engineering Using CRISPR/Cas Systems

Le Cong,<sup>1,2\*</sup> F. Ann Ran,<sup>1,4\*</sup> David Cox,<sup>1,3</sup> Shuailiang Lin,<sup>1,5</sup> Robert Barretto,<sup>6</sup> Naomi Habib,<sup>1</sup> Patrick D. Hsu,<sup>1,4</sup> Xuebing Wu,<sup>7</sup> Wenyang Jiang,<sup>8</sup> Luciano A. Marraffini,<sup>8</sup> Feng Zhang<sup>1†</sup>

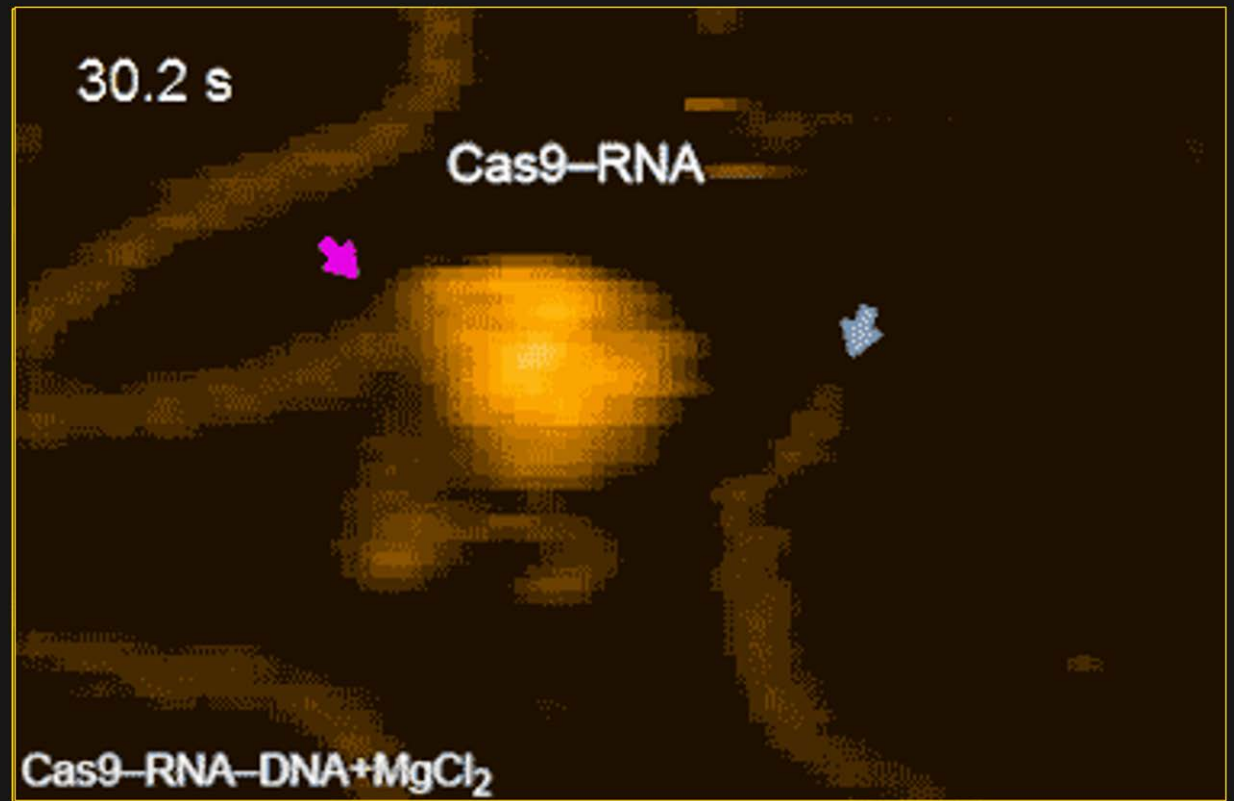
Functional elucidation of causal genetic variants and elements requires precise genome editing technologies. The type II prokaryotic CRISPR (clustered regularly interspaced short palindromic repeats)/Cas adaptive immune system has been shown to facilitate RNA-guided site-specific DNA cleavage. We engineered two different type II CRISPR/Cas systems and demonstrate that Cas9 nucleases can be directed by short RNAs to induce precise cleavage at endogenous genomic loci in human and mouse cells. Cas9 can also be converted into a nicking enzyme to facilitate homology-directed repair with minimal mutagenic activity. Lastly, multiple guide sequences can be encoded into a single CRISPR array to enable simultaneous editing of several sites within the mammalian genome, demonstrating easy programmability and wide applicability of the RNA-guided nuclease technology.

Precise and efficient genome-targeting technologies are needed to enable systematic reverse engineering of causal genetic variations by allowing selective perturbation of individual genetic elements. Although genome-editing technologies such as designer zinc fingers (ZFs) (1–4), transcription activator–like effectors (TALEs) (4–10), and homing meganucleases (11) have be-

gun to enable targeted genome modifications, there remains a need for new technologies that are scalable, affordable, and easy to engineer. Here, we report the development of a class of precision genome-engineering tools based on the RNA-guided Cas9 nuclease (12–14) from the type II prokaryotic clustered regularly interspaced short palindromic repeats (CRISPR) adaptive immune system (15–18).

Lights. Camera. Action... **CUT!**

CRISPR-Cas9 visualized  
by high-speed atomic  
force microscopy



M. Shibata, H. Nishimasu *et al.*  
*Nature Communications* 8, 1430 (2017)

Hiroshi Nishimasu (Univ Tokyo)



## The Heroes of CRISPR

Eric S. Lander<sup>1,2,3,\*</sup>

<sup>1</sup>Broad Institute of MIT and Harvard, 415 Main Street, Cambridge, MA

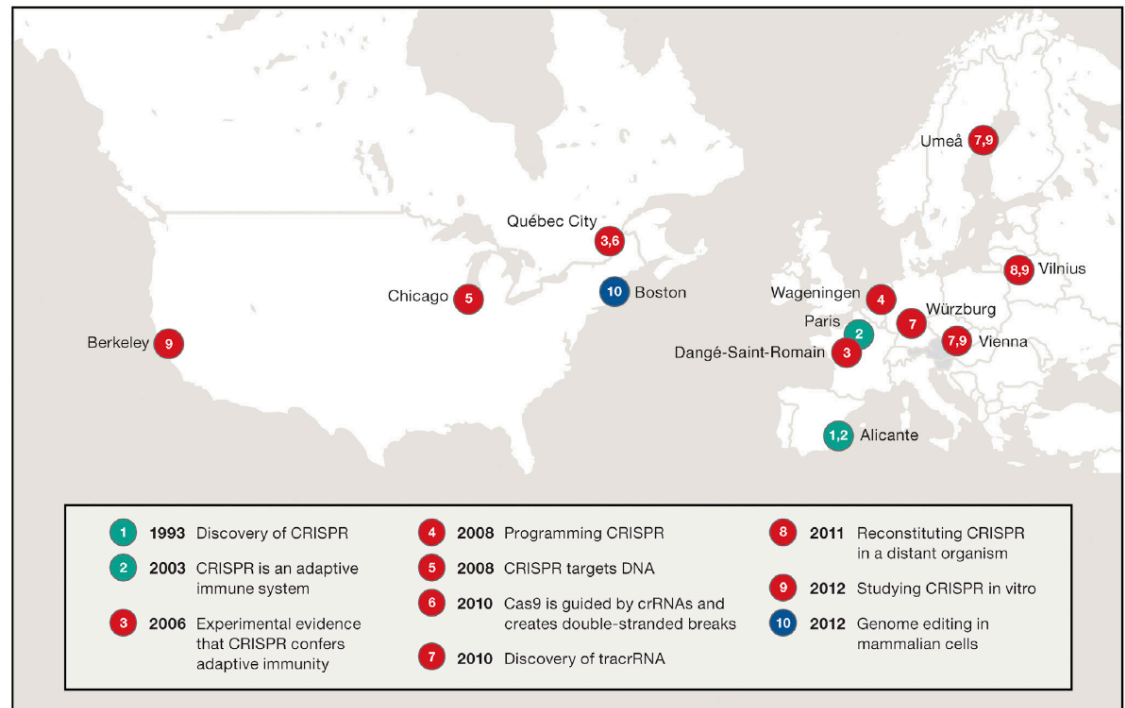
<sup>2</sup>Department of Biology, Massachusetts Institute of Technology, Cambridge, MA

<sup>3</sup>Department of Systems Biology, Harvard Medical School, Boston, MA

\*Correspondence: [lander@broadinstitute.org](mailto:lander@broadinstitute.org)

<http://dx.doi.org/10.1016/j.cell.2015.12.041>

Three years ago, scientists reported that CRISPR genome editing in living eukaryotic cells. Since then, by storm, with thousands of labs using it for applications preceding 20-year journey—the discovery of a natural system as an adaptive immune system; its biological engineering—remains little known. This Perspective and the stories of pioneers—and draw lessons from a scientific discovery.

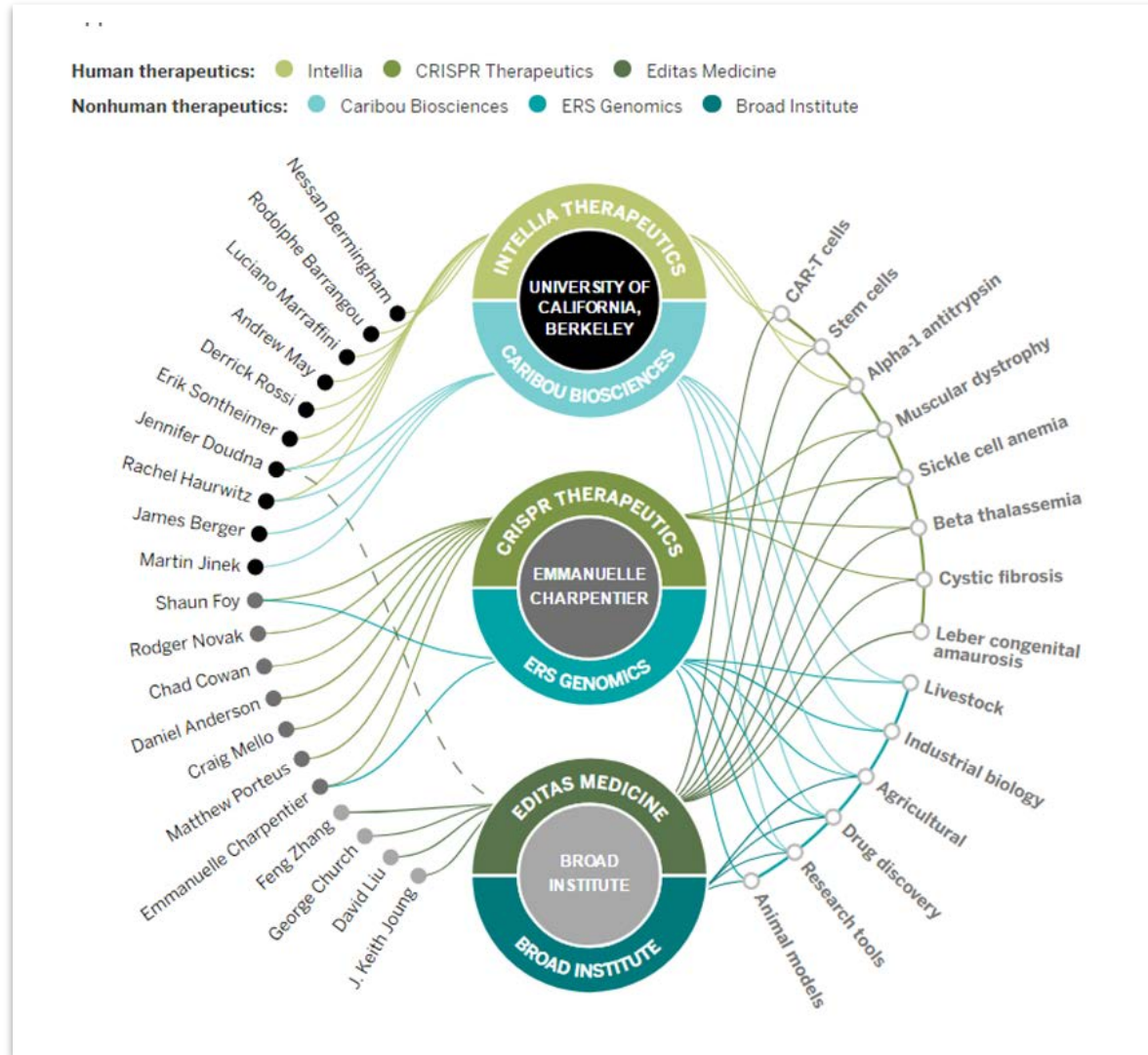
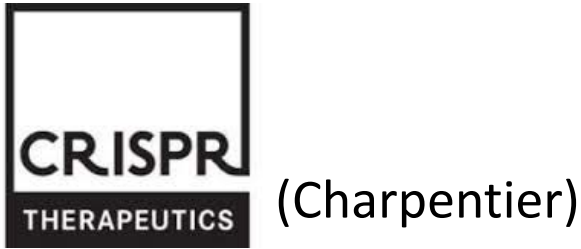


Lander's essay "is his masterwork, at once so evil and yet so brilliant that I find it hard not to stand in awe even as I picture him cackling loudly in his Kendall Square lair, giant laser weapon behind him poised to destroy Berkeley if we don't hand over our patents."

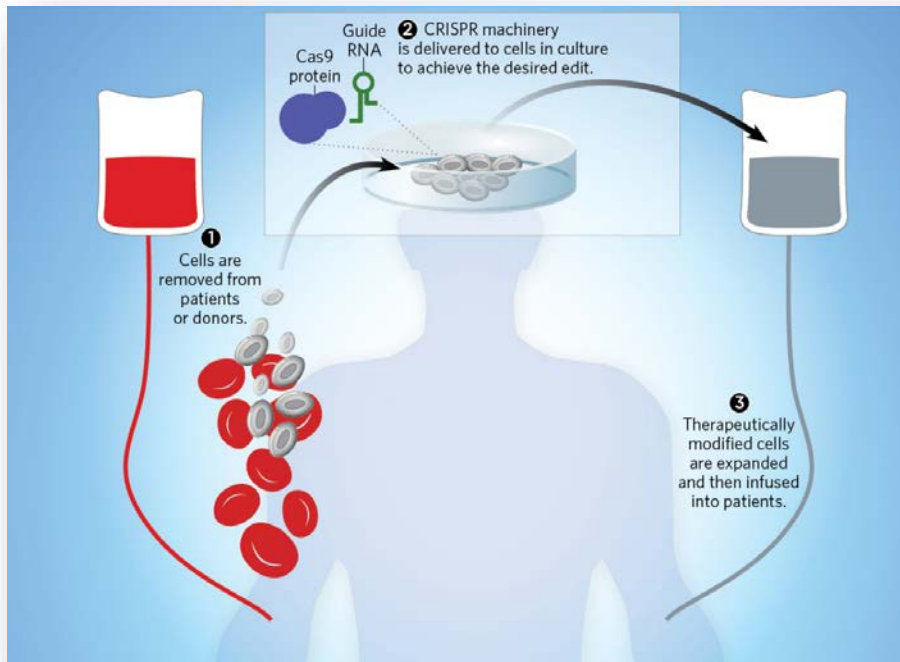
--- **Michael Eisen** (HHMI/UC

Berkeley)

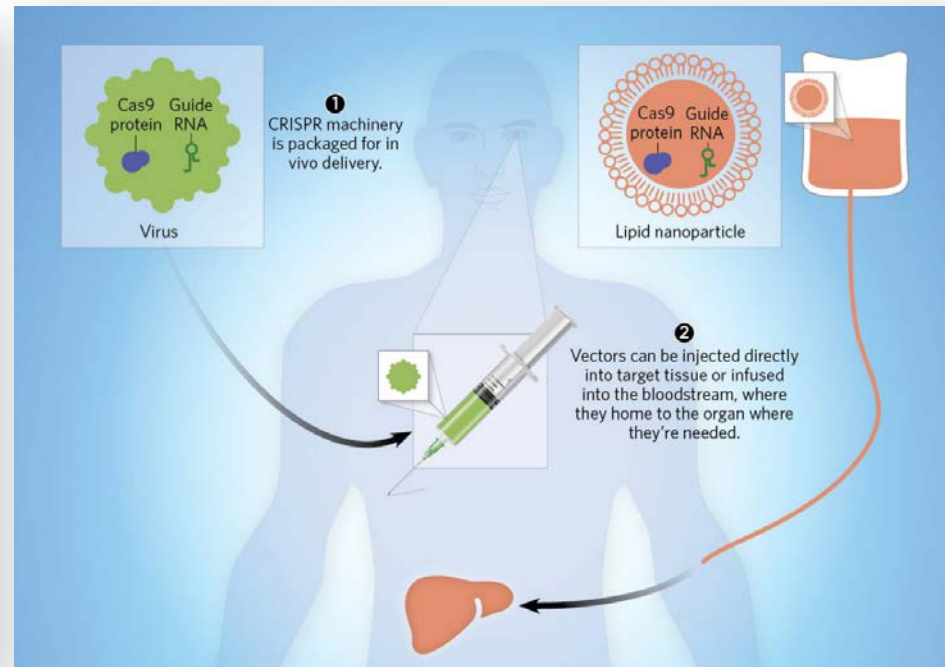
# CRISPR Business



# CRISPR Gene Therapy



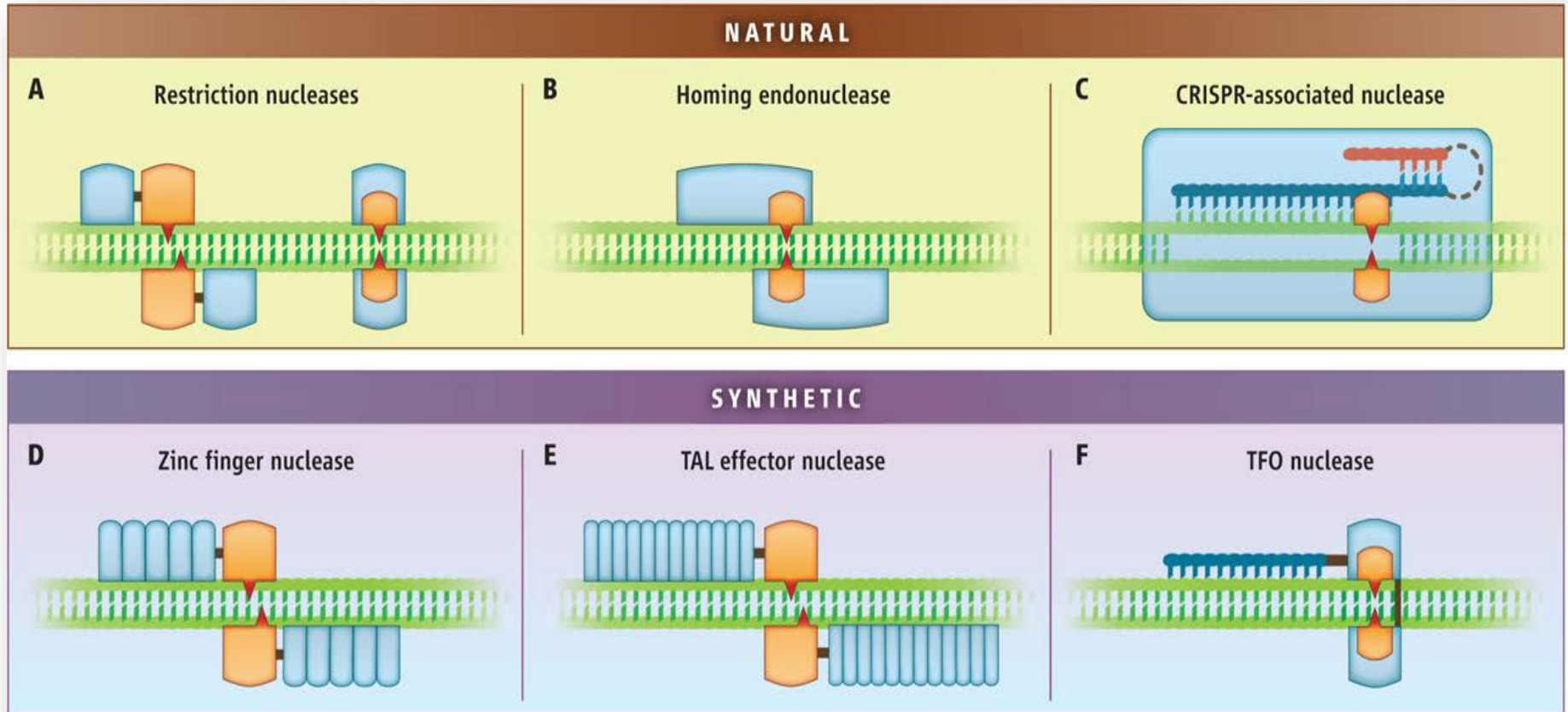
ex vivo



in situ



# Genome Editing B.C. (Before CRISPR)



# “Invisible Mending”

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## AP Exclusive: US scientists try 1st gene editing in the body

**AP** By The Associated Press  
November 15, 2017 7:48 am



Brian Madeux, 44, uses an infrared device to look at his veins as nurse Siobhan Field prepares an IV line for the first human gene editing therapy for NPS, at the UCSF Benioff Children's Hospital Oakland in Oakland, Calif., on Monday, Nov. 6, 2017. Madeux, who...

OAKLAND, Calif. (AP) — Scientists for the first time have tried editing a gene inside the body in a bold attempt to permanently change a person's DNA to try to cure a disease.

“It’s kind of humbling. I’m willing to take that risk. Hopefully it will help me and other people.”

--- Brian Madeux, 44  
Hunter Syndrome patient

Associated Press, Nov 15, 2017

New hope for China's  
left-behind kids p. 1226

How pesticides should  
be regulated p. 1232

A twist on photoemission  
delay pp. 1239 & 1274

# Science

\$15  
22 SEPTEMBER 2017  
sciencemag.org

AAAS

## CRISPR PIGS

Eliminating endogenous  
retrovirus in a step toward  
xenotransplantation



Luhan Yang (eGenesis)

### REPORT

## Inactivation of porcine endogenous retrovirus in pigs using CRISPR-Cas9

Dong Niu<sup>1,2,\*</sup>, Hong-Jiang Wei<sup>3,4,\*</sup>, Lin Lin<sup>5,\*</sup>, Haydy George<sup>1,\*</sup>, Tao Wang<sup>1,\*</sup>, I-Hsiu Lee<sup>1,\*</sup>, Hong-Ye Zhao<sup>3</sup>, Yong Wang<sup>6</sup>, Yanan Kan<sup>1</sup>, Ellen Shrock<sup>7</sup>, Emal Lasha<sup>1</sup>, Gang Wang<sup>1</sup>, Yonglun Luo<sup>5</sup>, Yubo Qing<sup>3,4</sup>, Deling Jiao<sup>3,4</sup>, Heng Zhao<sup>3,4</sup>, Xiaoyang Zhou<sup>6</sup>, Shouqi Wang<sup>8</sup>, Hong Wei<sup>6</sup>, Marc Güell<sup>1,†</sup>, George M. Church<sup>1,7,9,†</sup>, Luhan Yang<sup>1,†,‡</sup>

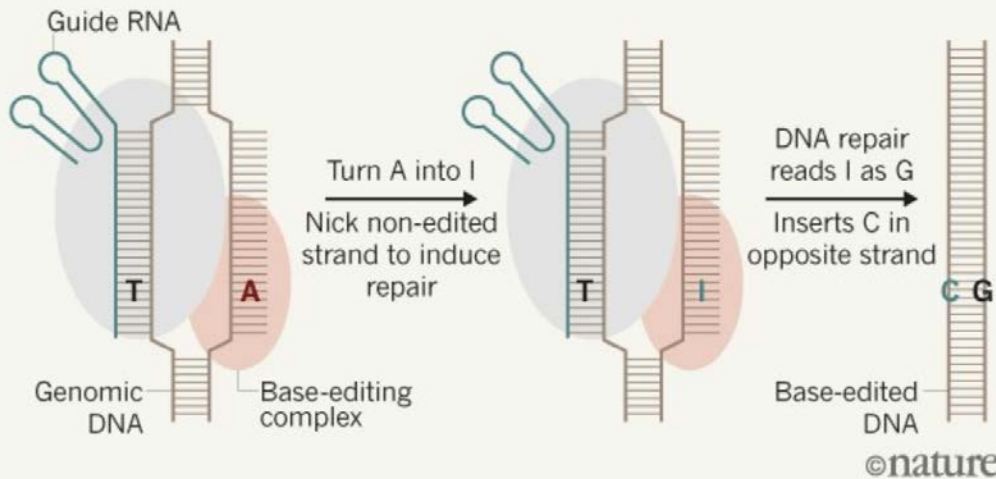
<sup>1</sup>eGenesis, Inc., Cambridge, MA 02139, USA.



# CRISPR 2.0: Base Editing

## CHANGING BASES

Researchers have devised several ways of making pinpoint changes in DNA and RNA. One technique uses a modified CRISPR–Cas9 system to edit single DNA base pairs.



David Liu (Harvard)

## Programmable base editing of A·T to G·C in genomic DNA without DNA cleavage

Nicole M. Gaudelli<sup>1,2,3</sup>, Alexis C. Komor<sup>1,2,3</sup>, Holly A. Rees<sup>1,2,3</sup>, Michael S. Packer<sup>1,2,3</sup>, Ahmed H. Badran<sup>1,2,3</sup>, David I. Bryson<sup>1,2,3</sup> & David R. Liu<sup>1,2,3</sup>

doi:10.1038/nature24644

# Is CRISPR Safe?



Cold  
Spring  
Harbor  
Laboratory

bioRxiv

THE PREPRINT SERVER FOR BIOLOGY

Confirmatory Results

## No unexpected CRISPR-Cas9 off-target activity gene-edited mice

Vivek Iyer, Katharina Boroviak, Mark Thomas, Brendan Doe, Edw

doi: <https://doi.org/10.1101/263129>

This article is a preprint and has not been peer-reviewed [what does this

### Abstract

CRISPR-Cas technologies have transformed genome-editing of experimental organisms and have immense therapeutic potential. Despite significant advances in our understanding of the CRISPR-Cas9 system, concerns remain over the potential for off-target effects. Recent studies have addressed these concerns using whole-genome sequencing (WGS) of gene-edited embryos or animals to search for de novo mutations (DNMs), which may represent candidate changes induced by poor editing fidelity. Critically, these studies used strain-matched but not pedigree-matched controls and thus were unable to reliably distinguish generational or colony-related differences from true DNMs. Here we used a trio design and whole genome sequenced 8 parents and 19 embryos, where 10 of the embryos were mutagenised with well-characterised gRNAs targeting the coat colour Tyrosinase (Tyr) locus. Detailed analyses of these whole genome data allowed us to conclude that if CRISPR mutagenesis were causing SNV or indel off-target mutations in treated embryos, then the number of these mutations is not statistically distinguishable from the background rate of DNMs occurring due to other processes.

New Results

## Identification of Pre-Existing Adaptive Immunity to Cas9 Proteins in Humans

Carsten Trevor Charlesworth, Priyanka S Deshpande, Daniel P Dever, Beruh Dejene, Natalia Gomez-Ospina, Sruthi Mantri, Mara Pavel-Dinu, Joab Camarena, Kenneth I Weinberg, Matthew H Porteus

doi: <https://doi.org/10.1101/243345>

This article is a preprint and has not been peer-reviewed [what does this mean?].

Abstract

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### Abstract

The CRISPR-Cas9 system has proven to be a powerful tool for the precise modification of specific DNA sequences. We are currently underway to use the CRISPR-Cas9 system to correct human genetic diseases. The most widely used homologous recombination derived from the bacterium *Streptococcus pyogenes* (C

STAT

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IN THE LAB

## CRISPR hits a snag: Our immune systems may attack the treatment

By ANDREW JOSEPH @DrewQJoseph / JANUARY 8, 2018

## Uh Oh—CRISPR Might Not Work in People

A sampling of human blood has turned up a surprise: most people could be immune to one of the world's biggest advances in genetic engineering.



nature  
biotechnology

Repair of double-strand breaks induced by  
CRISPR–Cas9 leads to large deletions and complex  
rearrangements

Michael Kosicki, Kärt Tomberg & Allan Bradley



**“This is the first systematic assessment of unexpected events resulting from CRISPR/Cas9 editing in therapeutically relevant cells, and we found that changes in the DNA have been seriously underestimated before now. It is important that anyone thinking of using this technology for gene therapy proceeds with caution, and looks very carefully to check for possible harmful effects.”**

**Allan Bradley**  
Director Emeritus  
Wellcome Sanger  
Institute

# ARTICLE

doi:10.1038/nature23305

## Correction of a pathogenic gene mutation in human embryos

Hong Ma<sup>1\*</sup>, Nuria Marti-Gutierrez<sup>1\*</sup>, Sang-Wook Park<sup>2\*</sup>, Jun Wu<sup>3\*</sup>, Yeonmi Lee<sup>1</sup>, Keiichiro Suzuki<sup>3</sup>, Amy Koski<sup>1</sup>, Dongmei Ji<sup>1</sup>, Tomonari Hayama<sup>1</sup>, Riffat Ahmed<sup>1</sup>, Hayley Darby<sup>1</sup>, Crystal Van Dyken<sup>1</sup>, Ying Li<sup>1</sup>, Eunju Kang<sup>1</sup>, A.-Reum Park<sup>2</sup>, Daesik Kim<sup>4</sup>, Sang-Tae Kim<sup>2</sup>, Jianhui Gong<sup>5,6,7,8</sup>, Ying Gu<sup>5,6,7</sup>, Xun Xu<sup>5,6,7</sup>, David Battaglia<sup>1,9</sup>, Sacha A. Krieg<sup>9</sup>, David M. Lee<sup>9</sup>, Diana H. Wu<sup>9</sup>, Don P. Wolf<sup>1</sup>, Stephen B. Heitner<sup>10</sup>, Juan Carlos Izpisua Belmonte<sup>3,§</sup>, Paula Amato<sup>1,9,§</sup>, Jin-Soo Kim<sup>2,4,§</sup>, Sanjiv Kaul<sup>10,§</sup> & Shoukhrat Mitalipov<sup>1,10,§</sup>

Genome editing of human embryos using CRISPR-Cas9 technology has been shown to be accurate and efficient. In this study, the authors demonstrate the correction of a pathogenic gene mutation in human embryos using CRISPR-Cas9 technology. The study shows that the DSB was repaired accurately and the resulting embryos were viable and healthy. The authors also show that the correction of the mutation was permanent and heritable. This study is a significant step towards the use of genome editing in human embryos for the treatment of genetic diseases.



Shoukhrat Mitalipov (OHSU)

50p

THE 1 PAPER · BRITAIN'S FIRST AND ONLY CONCISE QUALITY TITLE

**i world exclusive**

# One giant step for designer babies

*The essential daily briefing*

THURSDAY 27 JULY 2017

News.co.uk

### Twin Peaty

Swimming hero wins second gold

» **Revealed** Era of genetically modified babies moves closer, as scientists prove they can safely alter human embryos

» Inherited diseases caused by defective genes can be corrected in the earliest stage of life, revolutionary technique shows

» Same technology could be used to select stronger muscles or better eyesight, prompting fierce ethical debate

» 'They've done it. The quality of the work is high,' top scientist tells

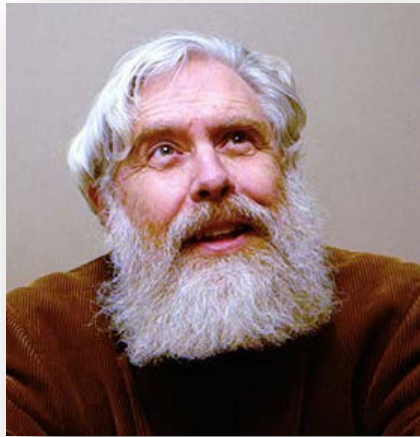
» Religious organisations likely to oppose groundbreaking research

**SPECIAL REPORT SYSTEME CONNOR, PAGES 6-7**

INSIDE CAR BAN BACKLASH | SCIENCE | TV & RADIO | GAMES

●● The concept of altering the human germline in embryos for clinical purposes has been debated over many years from many different perspectives, and has been viewed almost universally as a line that should not be crossed... Designer babies make great Hollywood movies. They make really bad science, and I think they are really bad ethics.”

---



●● Gene editing is already used to make GM-persons [cancer, HIV]... If by GM-persons, we limit our meaning to heritable DNA changes, then the most likely use would be changing deadly DNA variants into their common healthy versions. This should be far safer than testing new drugs, which impact complex human systems in unknown ways.”



# NEWS RELEASES

Tuesday, January 23, 2018

## NIH to launch genome editing research program

*Somatic Cell Genome Editing aims to develop tools for safe and effective genome editing in humans.*



The National Institutes of Health will launch an effort aimed at removing barriers that slow the adoption of genome editing for treating patients. This program, Somatic Cell Genome Editing, plans to award researchers approximately \$190 million over six years beginning this year, pending availability of funds. These researchers will collaborate to improve the delivery mechanisms for targeting gene editing tools in patients, develop new and improved genome editors, develop assays for testing the safety and efficacy of the genome editing tools in animal and human cells, and assemble a genome editing toolkit containing the resulting knowledge, methods, and tools to be shared with the scientific community. The program is funded by NIH's Common Fund.

The National Academies of  
SCIENCES • ENGINEERING • MEDICINE

REPORT

# Human Genome Editing

**SCIENCE,  
ETHICS,  
AND  
GOVERNANCE**

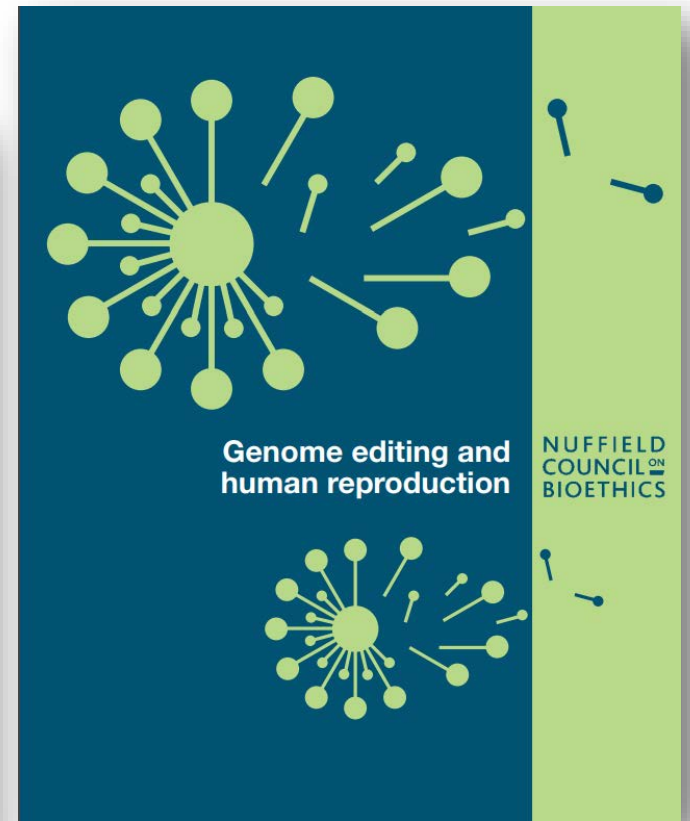
NATIONAL ACADEMY OF SCIENCES  
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“We say proceed with all due caution, but we don’t prohibit germline, after considerable discussion and debate. We’re talking only about fixing diseases.”

-- Rick Hynes (HHMI/MIT)

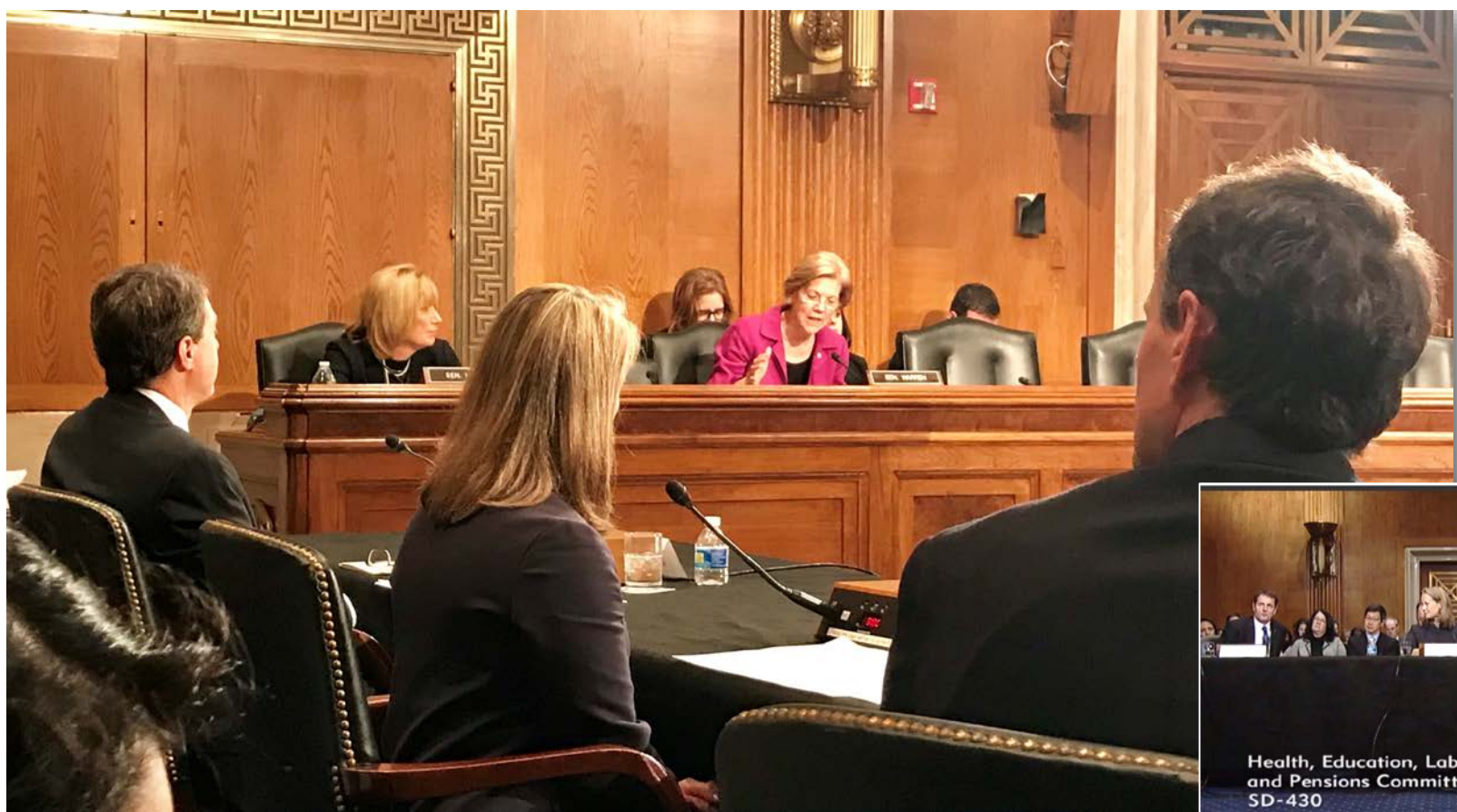
# Genetically modified babies given go ahead by UK ethics body

The Nuffield Council on Bioethics says changing the DNA of a human embryo could be 'morally permissible' if it is in the child's best interests





# CRISPR Goes to Washington





Corteva

## What Is a Genetically Modified Crop? A European Ruling Sows Confusion

In Europe, plants created with gene-editing technologies will be stringently regulated as G.M.O.'s. But older crops whose DNA has been altered will be left alone.



**Matt Ridley**  
@mattwidley

Follow

Catastrophic decision of the ECJ to go against the advice of its advocate general and treat gene-edited crops as GMOs.

Will ensure European farming is less competitive, less innovative, more dependent on chemicals and worse for the environment.



**Ewan Birney**  
@ewanbirney

Follow

So placing CRISPR in the same bucket of all GMO and uv, gamma, chemical mutations in another bucket is like the Catholic Church classifying ducks as fish



**Jonathan Pettitt**  
@genotripe

Follow

#ECJ statement is beyond dumb: "organisms obtained by mutagenesis are GMOs within the meaning of the GMO Directive". All plants are GMOs by this definition - mutagenesis is an unavoidable consequence of DNA replication. #GMOs #CRISPR



**Owen Paterson MP**  
@OwenPaterson

Follow

All too predictable ruling from the European Court which sadly "threatens research on gene-edited crops in the bloc", & will see investment move elsewhere as the EU condemns itself to become the Museum of World Farming [nature.com/articles/d4158...](https://www.nature.com/articles/d4158...)



**Kevin Folta**  
@kevinfolta

Follow

Hey EU! Politicians have decided you won't get gene edited crops. Fire up the Cesium 137 source and order some ethyl methanesulfonate - you're doing it old school. Condolences to EU colleagues & farmers that can't benefit from the best technology.



**Mark Lynas**  
@mark\_lynas

Follow

European court plumbs the depths of scientific absurdity with today's gene editing decision - random mutagenesis is OK, while precision editing is a 'GMO' and therefore borderline illegal. Go figure! Like saying doctors can use blunderbuss but not scalpel.



**Clive G. Brown**  
@Clive\_G\_Brown

Follow

Central also is the misanthropic idea that there is a 'Nature' that exists in a state of perfected harmony and that humans and all their works are corrupting it. If only these twits realized that all of their beloved vegetables and most farm animals are hideous mutants.

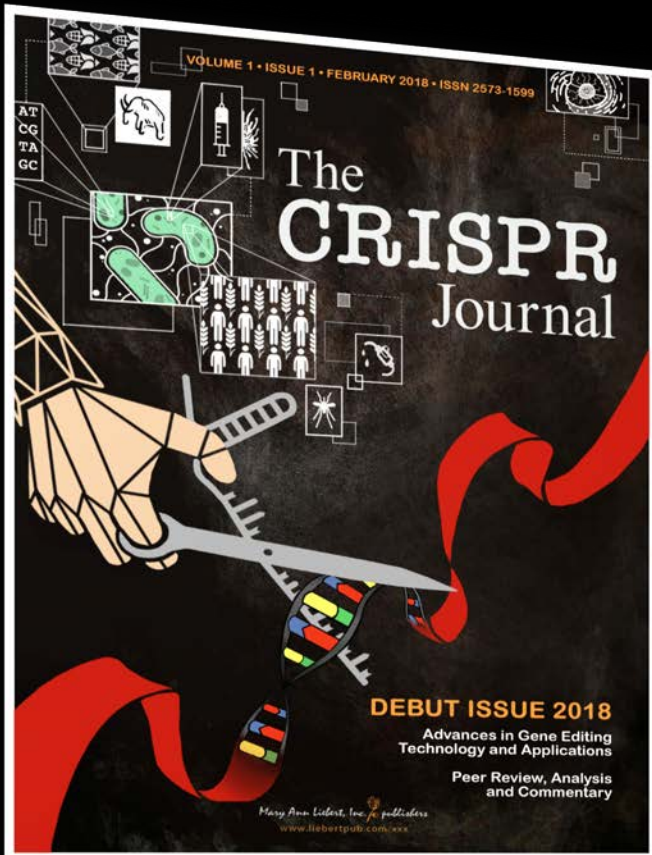




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- Cutting edge of CRISPR/genome editing
- Research, commentary and analysis
- Intended for (and edited by) the global CRISPR community
- Speed of peer review and production
- Print (6x/year), online and OA options
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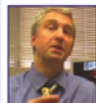
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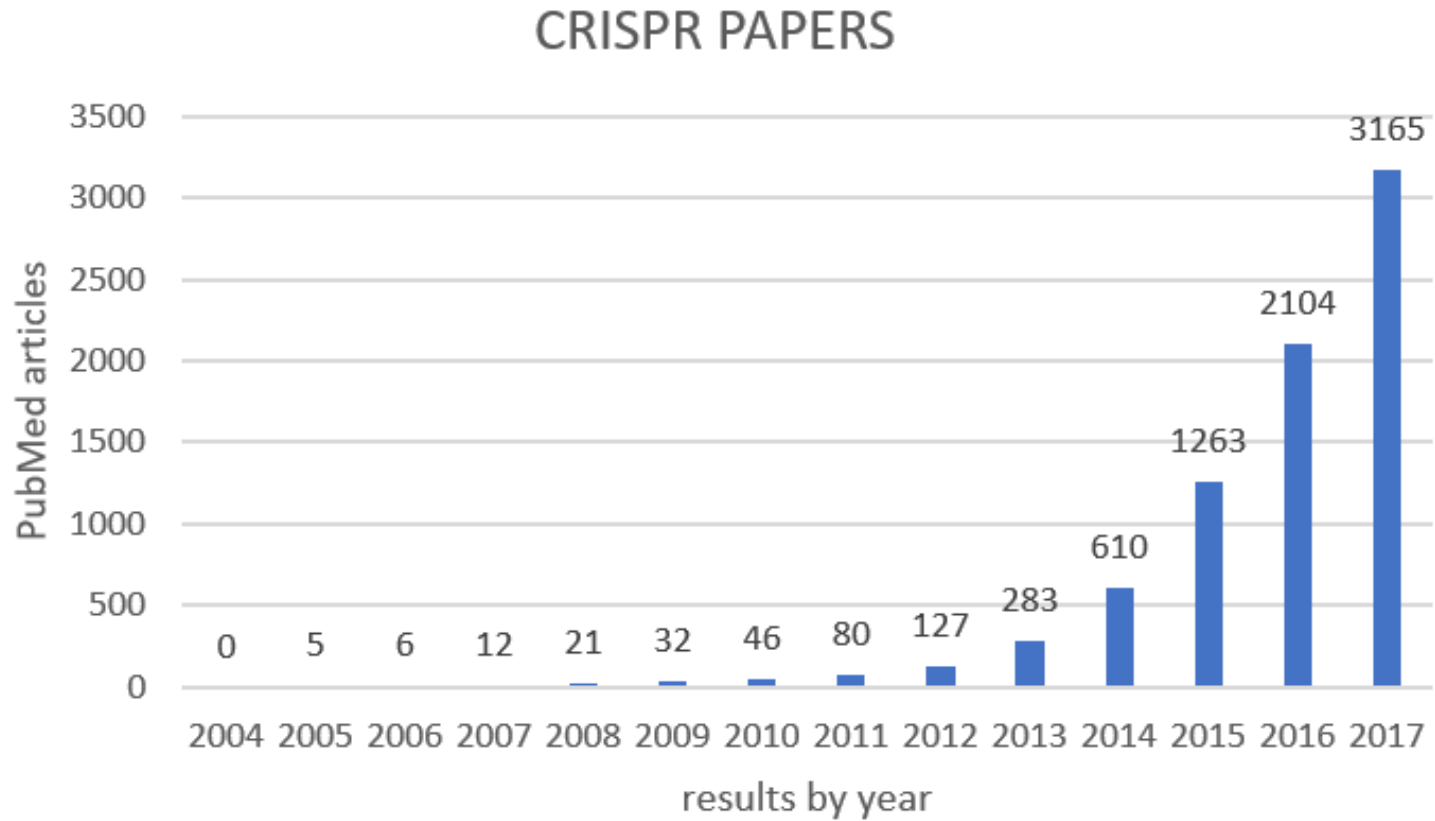
**Huimin Zhao**  
University of Illinois



**Jian-Kang Zhu**  
Shanghai Institute for  
Biological Sciences



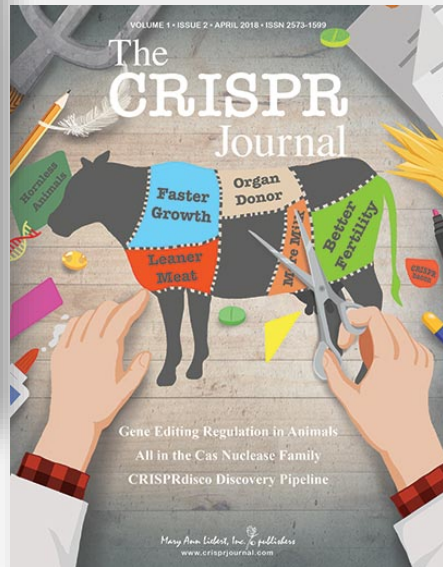
# CRISPR on the Rise



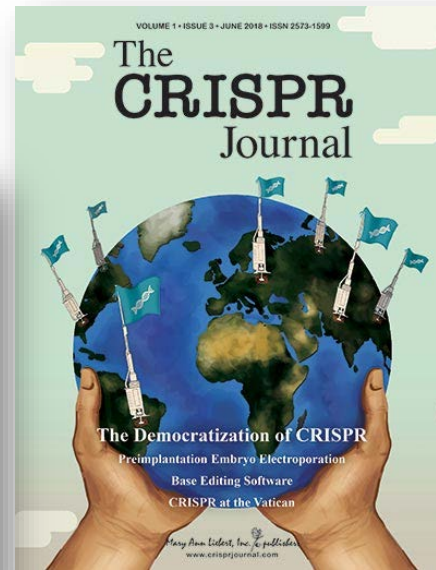
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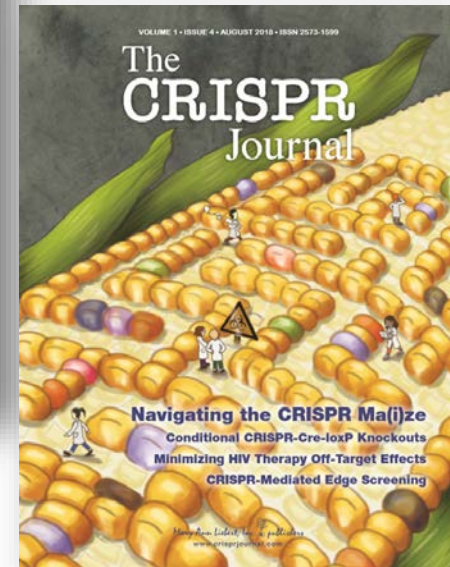
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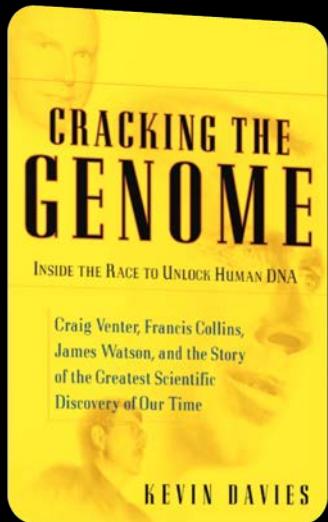
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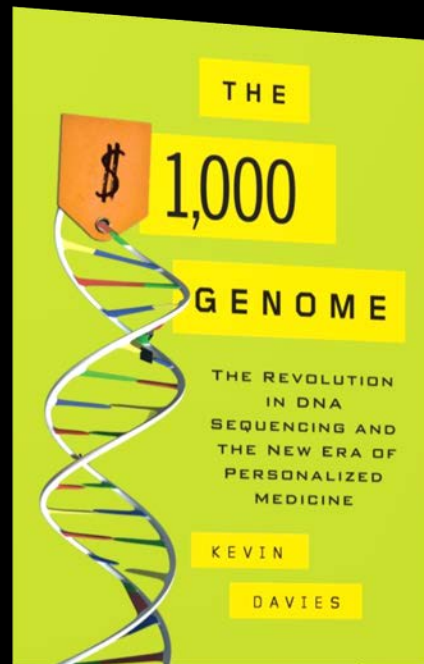
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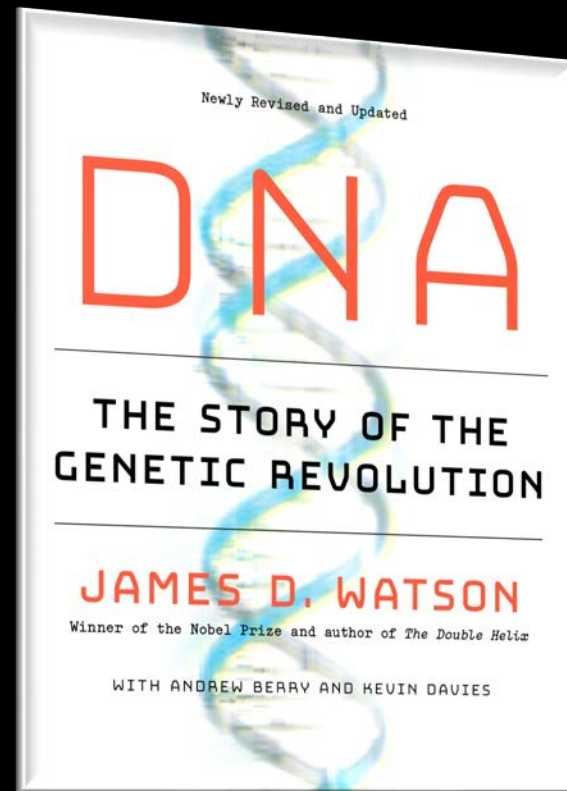
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