

## Genetic scissors: a tool for rewriting the code of life

*Emmanuelle Charpentier and Jennifer Doudna are awarded the Nobel Prize in Chemistry 2020 for discovering one of gene technology's sharpest tools: the CRISPR/Cas9 genetic scissors. Researchers can use these to change the DNA of animals, plants and microorganisms with extremely high precision. This technology has revolutionised the molecular life sciences, brought new opportunities for plant breeding, is contributing to innovative cancer therapies and may make the dream of curing inherited diseases come true.*

One of the attractions of science is that it is unpredictable – you can never know in advance where an idea or a question may lead. Sometimes a curious mind will meet a dead end, sometimes it will encounter a thorny labyrinth that takes years to navigate. But, now and again, she realises she is the first person ever to gaze upon a horizon of untold possibility.

The gene editor called CRISPR-Cas9 is one such unexpected discovery with breathtaking potential. When Emmanuelle Charpentier and Jennifer Doudna started investigating the immune system of a *Streptococcus* bacterium, one idea was that they could perhaps develop a new form of antibiotic. Instead, they discovered a molecular tool that can be used to make precise incisions in genetic material, making it possible to easily change the code of life.



Figure 1. Using the genetic scissors, researchers can edit the genome of practically all living things.

### A powerful tool that affects everyone

Just eight years after their discovery, these genetic scissors have reshaped the life sciences. Biochemists and cell biologists can now easily investigate the functions of different genes and their possible role in the progression of disease. In plant breeding, researchers can give plants specific characteristics, such as the ability to withstand drought in a warmer climate. In medicine, this gene editor is contributing to new cancer therapies and the first studies attempting to cure inherited diseases.

There are almost endless examples of how CRISPR-Cas9 could be used, which also include unethical applications. As with all powerful technology, these genetic scissors need to be regulated. More on that later.

In 2011, neither Emmanuelle Charpentier nor Jennifer Doudna had any idea that their first meeting, in a café in Puerto Rico, was a life-changing encounter. We will start by presenting Charpentier, who initially proposed their collaboration.

## Charpentier is fascinated by pathogenic bacteria

Some people have called her driven, attentive and thorough. Others say that Emmanuelle Charpentier always looks for the unexpected. Herself, she quotes Louis Pasteur, “Chance favours the prepared mind”. The urge to make new discoveries and the desire to be free and independent have governed her path. Including her doctoral studies at the Institut Pasteur in Paris, she has lived in five different countries, seven different cities and worked at ten different institutions.

Her surroundings and approaches have shifted, but the majority of her research has one common denominator: pathogenic bacteria. Why are they so aggressive? How do they develop their resistance to antibiotics? And is it possible to find new treatments that can stop their progress?

In 2002, when Emmanuelle Charpentier started her own research group at the University of Vienna, she focused on one of the bacteria that cause the greatest harm to humanity: *Streptococcus pyogenes*. Every year, it infects millions of people, often causing easily treatable infections such as tonsillitis and impetigo. However, it can also cause life-threatening sepsis and break down the soft tissues in the body, giving it a reputation as a ‘flesh eater’.

To better understand *S. pyogenes*, Charpentier began by thoroughly investigating how this bacterium’s genes are regulated. This decision was the first step on the path to the discovery of the genetic scissors – but before we walk further along that road, we will find out more about Jennifer Doudna. Because while Charpentier is making detailed studies of *S. pyogenes*, Doudna hears – for the first time – an abbreviation that she thinks sounds like *crisper*.

## Science – as much adventure as a detective story

Even as a child growing up on Hawaii, Jennifer Doudna had a strong urge to know things. One day, her father placed James Watson’s book *The Double Helix* on her bed. This detective-style story about how James Watson and Francis Crick solved the structure of the DNA molecule was like nothing she had read in her school textbooks. She was captivated by the scientific process and realised that science is more than just facts.

However, when she started to solve scientific mysteries, her attention was not on DNA, but on its molecular sibling: RNA. In 2006 – when we meet her – she is leading a research group at the University of California, Berkeley, and has two decades’ experience of working with RNA. She has a reputation as a successful researcher with a nose for ground-breaking projects, and has recently entered an exciting new field: *RNA interference*.

For many years, researchers had believed that they understood the basic function of RNA, but they suddenly discovered lots of small RNA molecules that help regulate gene activity in cells. Jennifer Doudna’s involvement in RNA interference is the reason why, in 2006, she gets a phone call from a colleague in a different department.

## Bacteria carry an ancient immune system

Her colleague, who is a microbiologist, tells Doudna about a new discovery: when researchers compare the genetic material of vastly different bacteria, as well as archaea (a type of microorganism), they find repetitive DNA sequences that are surprisingly well preserved. The same code appears over and over again, but between the repetitions there are unique sequences that differ (figure 2). It is like the same word is being repeated between each unique sentence in a book.

These arrays of repeated sequences are called *clustered regularly interspaced short palindromic repeats*, abbreviated as CRISPR. The interesting thing is that the unique, non-repetitive sequences in CRISPR appear to match the genetic code of various viruses, so the current thinking is that this is one part of an ancient immune system that protects bacteria and archaea from viruses. The hypothesis is that if a bacterium has succeeded in surviving a virus infection, it adds a piece of the virus' genetic code into its genome as a memory of the infection.

No one yet knows how all this works, says her colleague, but the suspicion is that the mechanism used by bacteria to neutralise a virus is similar to that studied by Doudna: RNA interference.

## Doudna maps a complex machinery

This news is both remarkable and thrilling. If it is true that bacteria have an ancient immune system, then this is a big deal. Jennifer Doudna's sense of molecular intrigue comes to life and she starts to learn everything she can about the CRISPR system.

It turns out that, in addition to the CRISPR sequences, researchers have discovered special genes that they have called *CRISPR-associated*, abbreviated as *cas*. What Doudna finds interesting is that these genes are very similar to genes that code for already known proteins that specialise in unwinding and cutting up DNA. So do the Cas proteins have the same function? Do they cleave virus DNA?

She puts her research group to work and, after a few years, they have succeeded in revealing the function of several different Cas proteins. In parallel, a handful of other research groups at other universities are studying the newly discovered CRISPR/Cas system. Their mapping shows that bacteria's immune systems can take very different forms. The CRISPR/Cas system studied by Doudna belongs to class 1; it is a complex machinery that requires many different Cas proteins to disarm a virus. The class 2 systems are significantly simpler because they need fewer proteins. In another part of the world, Emmanuelle Charpentier has just come across such a system. Back to her.

## A new and unknown piece of the CRISPR-system puzzle

When we left Emmanuelle Charpentier she was living in Vienna, but in 2009 she moved to a position with good research opportunities at Umeå University in the north of Sweden. She was warned about moving to such a remote part of the world, but the long, dark winter allows her plenty of peace and quiet for work.

And she needs it. She is also interested in small, gene-regulating RNA molecules and, working with researchers in Berlin, she has mapped the small RNAs found in *S. pyogenes*. The results have given her a lot to think about, because one of the small RNA molecules that exists in large amounts in this bacterium is an as yet unknown variant, and the genetic code for this RNA is very close to the peculiar CRISPR sequence in the bacterium's genome.

The similarities between the two make Charpentier suspect that they are linked. Careful analysis of their genetic codes also reveals that one part of the small and unknown RNA molecule matches the part of CRISPR that is repeated. It is like finding two puzzle pieces that fit together perfectly (figure 2).

Charpentier had never worked with CRISPR, but her research group initiates some thorough microbiological detective work to map the CRISPR system in *S. pyogenes*. This system, which belongs to class 2, was already known to only require a single Cas protein, Cas9, to cleave virus DNA. Charpentier shows that the unknown RNA molecule, which is named *trans-activating crisper RNA* (*tracrRNA*), also has a decisive function; it is necessary for the long RNA that is created from the CRISPR sequence in the genome to mature into its active form (figure 2).

## *Streptococcus*' natural immune system against viruses: CRISPR/Cas9

When viruses infect a bacterium, they send their harmful DNA into it. If the bacterium survives the infection, it inserts a piece of the virus DNA in its genome, like a memory of the virus. This DNA is then used to protect the bacterium from new infections.

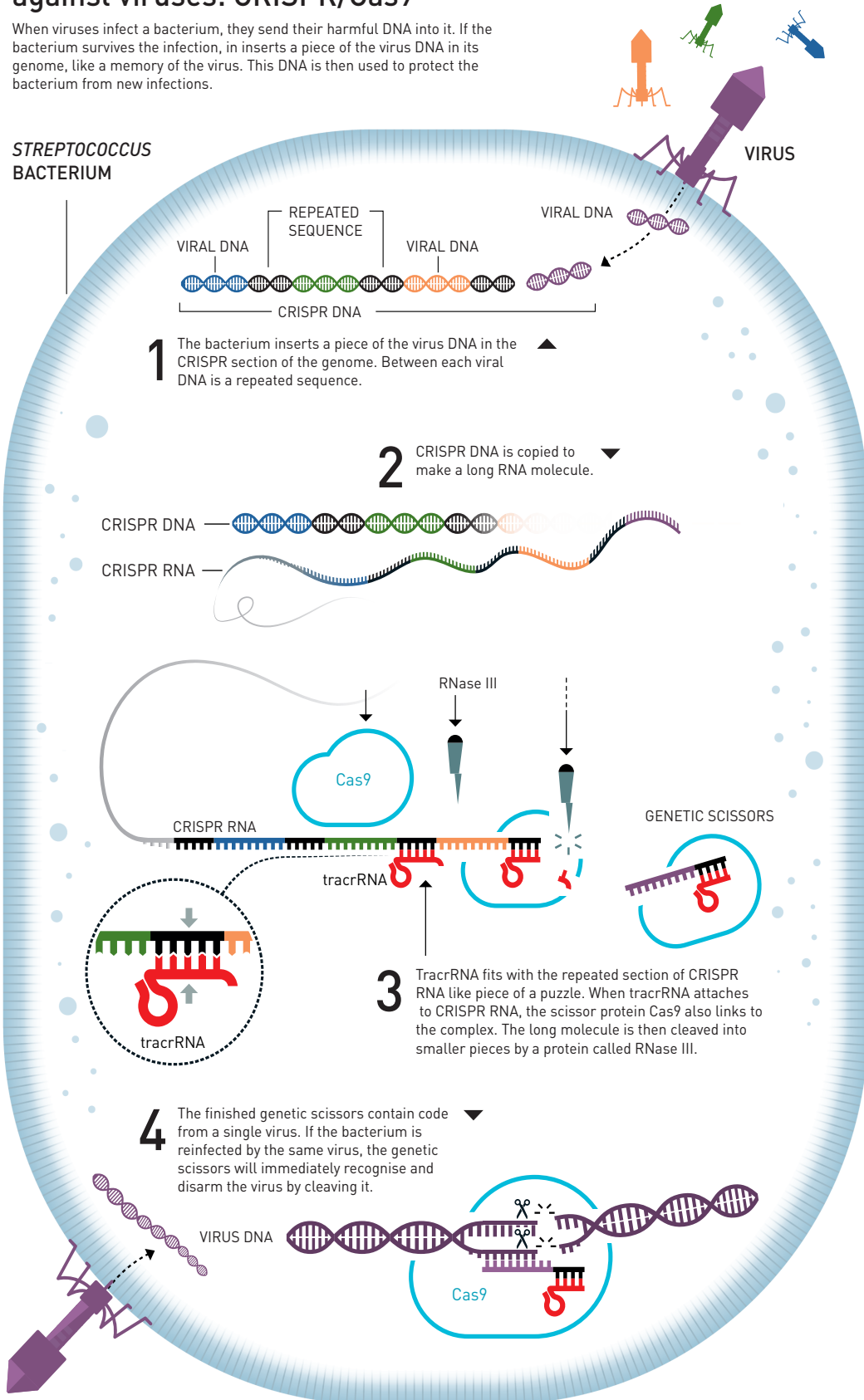


Figure 2

After intensive and targeted experimentation, Emmanuelle Charpentier publishes the discovery of tracrRNA in March 2011. She knows that she is on the heels of something very exciting. She has many years of experience in microbiology and in her continuing investigation of the CRISPR-Cas9 system she wants to cooperate with a biochemist. Jennifer Doudna is the natural choice. So that spring, when Charpentier is invited to a conference in Puerto Rico to talk about her findings, her aim is to meet this skilled Berkeley researcher.

### **A life-changing meeting in a Puerto Rican café**

By coincidence, they meet at a café on the second day of the conference. A colleague of Doudna introduces them to each other and, the following day, Charpentier proposes that they should explore the old parts of the capital city together. As they stroll along the cobbled streets, they start talking about their research. Charpentier wonders whether Doudna is interested in a collaboration – would she like to participate in studying the function of Cas9 in *S. pyogenes*' simple class 2 system?

Jennifer Doudna is intrigued, and they and their colleagues make plans for the project via digital meetings. Their suspicion is that CRISPR-RNA is needed to identify a virus' DNA, and that Cas9 is the scissor that cuts off the DNA molecule. However, nothing happens when they test this in vitro. The DNA molecule remains intact. Why? Is something wrong with the experimental conditions? Or does Cas9 have an entirely different function?

After a great deal of brainstorming and numerous failed experiments, the researchers finally add tracrRNA to their tests. Previously, they believed that tracrRNA was only necessary when CRISPR-RNA was cleaved into its active form (figure 2), but once Cas9 had access to tracrRNA what everyone was waiting for actually happened: the DNA molecule was cleaved into two parts.

Evolutionary solutions have often surprised researchers, but this was something extraordinary. The weapon that streptococci have developed as a protection from viruses is simple and effective, even brilliant. The history of genetic scissors could have stopped here; Charpentier and Doudna had uncovered a fundamental mechanism in a bacterium that causes great suffering for humanity. That discovery was astounding in itself, but chance favours prepared minds.

### **An epoch-making experiment**

The researchers decide to try to simplify the genetic scissors. Using their new knowledge about tracrRNA and CRISPR-RNA, they figured out how to fuse the two into a single molecule, which they named guide RNA. With this simplified variant of the genetic scissors, they then undertake an epoch-making experiment: they investigate whether they can control this genetic tool so that it cuts the DNA at a location decided by the researchers.

By this time, the researchers know that they are close to a major breakthrough. They take a gene that is already in a freezer in Doudna's laboratory and select five different places where the gene should be cleaved. They then change the CRISPR part of the scissors so that its code matches the code where the cuts are to be made (figure 3). The result was overwhelming. The DNA molecules were cleaved in exactly the right places.



## The CRISPR/Cas9 genetic scissors

When researchers are going to edit a genome using the genetic scissors, they artificially construct a guide RNA, which matches the DNA code where the cut is to be made. The scissor protein, Cas9, forms a complex with the guide RNA, which takes the scissors to the place in the genome where the cut will be made.

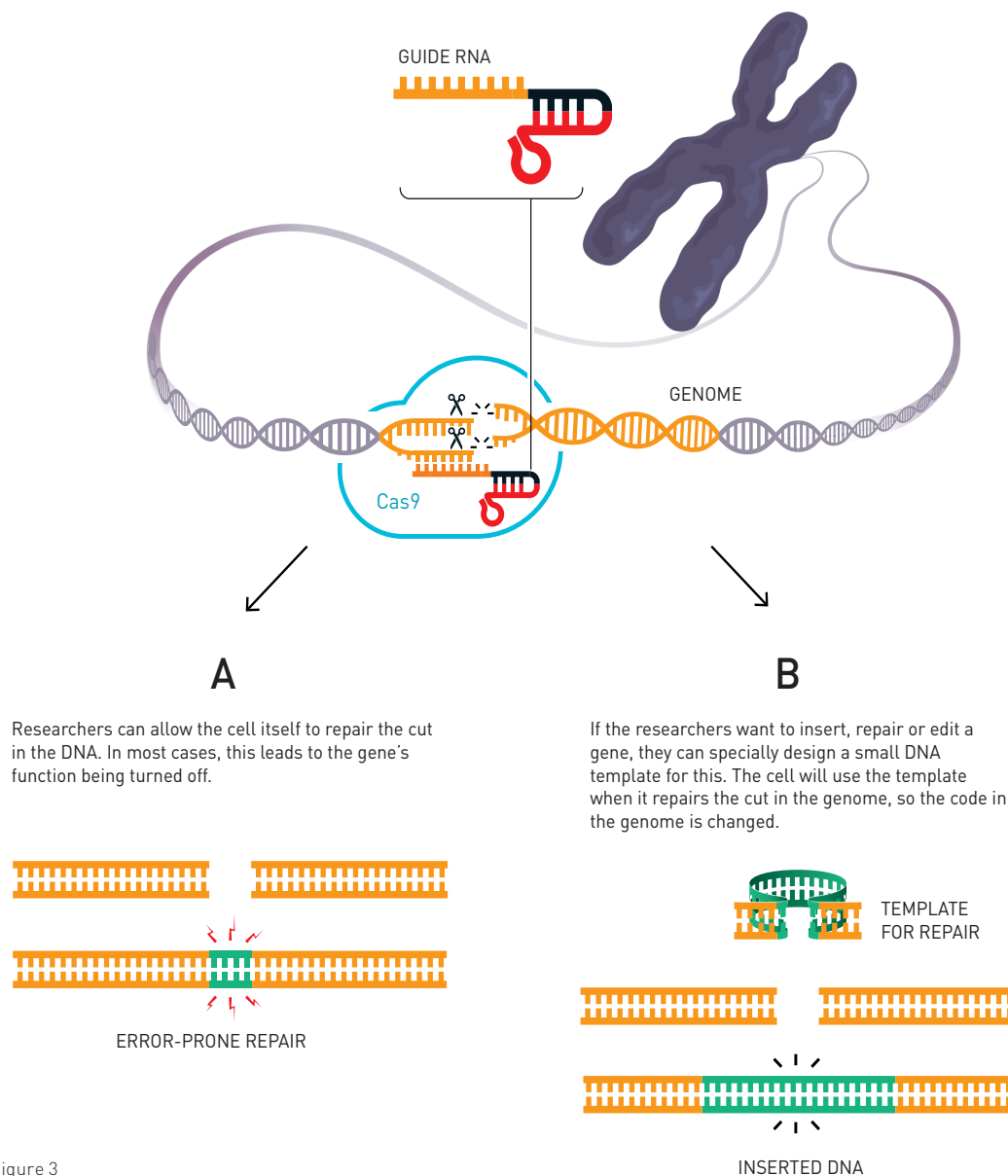


Figure 3

## Genetic scissors change the life sciences

Soon after Emmanuelle Charpentier and Jennifer Doudna publish their discovery of the CRISPR/Cas9 genetic scissors in 2012, several research groups demonstrate that this tool can be used to modify the genome in cells from both mice and humans, leading to explosive development. Previously, changing the genes in a cell, plant or organism was time-consuming and sometimes impossible. Using the genetic scissors, researchers can – in principle – make cuts in whichever genome they wish. After this, it is easy to utilise the cell's natural systems for DNA repair so that they rewrite the code of life (figure 3).

Because this gene tool is so easy to use, it is now widespread in basic research. It is used to change the DNA of cells and laboratory animals for the purpose of understanding how different genes function and interact, such as during the course of a disease.

Genetic scissors have also become a standard tool in plant breeding. The methods previously used by researchers to modify plant genomes often required the addition of genes for antibiotic resistance. When the crops were planted, there was a risk of this antibiotic resistance spreading to the surrounding microorganisms. Thanks to the genetic scissors, researchers no longer need to use these older methods as they can now make very precise changes to the genome. Among other things, they have edited the genes that make rice absorb heavy metals from the soil, leading to improved rice varieties with lower levels of cadmium and arsenic. Researchers have also developed crops that better withstand drought in a warmer climate, and which resist insects and pests that would otherwise have to be dealt with using pesticides.

### **Hope of curing inherited diseases**

In medicine, the genetic scissors are contributing to new immunotherapies for cancer and trials are underway to make a dream come true – curing inherited diseases. Researchers are already performing clinical trials to investigate whether they can use CRISPR/Cas9 to treat blood diseases such as sickle cell anaemia and beta thalassaemia, as well as inherited eye diseases.

They are also developing methods for repairing genes in large organs, such as the brain and muscles. Animal experiments have shown that specially designed viruses can deliver the genetic scissors to the desired cells, treating models of devastating inherited diseases such as muscular dystrophy, spinal muscular atrophy and Huntington's disease. However, the technology needs further refinement before it can be tested on humans.

### **The power of genetic scissors requires regulation**

Alongside all their benefits, genetic scissors can also be misused. For example, this tool can be used to create genetically modified embryos. However, for many years there have been laws and regulations that control the application of genetic engineering, which include prohibitions on modifying the human genome in a way that allows the changes to be inherited. Also, experiments that involve humans and animals must always be reviewed and approved by ethical committees before they are carried out.

One thing is certain: these genetic scissors affect us all. We will face new ethical issues, but this new tool may well contribute to solving many of the challenges now facing humanity. Through their discovery, Emmanuelle Charpentier and Jennifer Doudna developed a chemical tool that has taken life sciences into a new epoch. They have made us gaze out onto a vast horizon of unimagined potential and, along the way – as we explore this new land – we are guaranteed to make new and unexpected discoveries.

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## FURTHER READING

Additional information on this year's prizes, including a scientific background in English, is available on the website of the Royal Swedish Academy of Sciences, [www.kva.se](http://www.kva.se), and at [www.nobelprize.org](http://www.nobelprize.org), where you can watch video footage of the press conferences, the Nobel Lectures and more. Information on exhibitions and activities related to the Nobel Prizes and the Prize in Economic Sciences is available at [www.nobelprizemuseum.se](http://www.nobelprizemuseum.se)

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## The Royal Swedish Academy of Sciences has decided to award the Nobel Prize in Chemistry 2020 to

### **EMMANUELLE CHARPENTIER**

Born 1968 in Juvisy-sur-Orge, France.  
Ph.D. 1995 from Institut Pasteur, Paris, France. Director of the Max Planck Unit for the Science of Pathogens, Berlin, Germany.

### **JENNIFER A. DOUDNA**

Born 1964 in Washington, D.C, USA.  
Ph.D. 1989 from Harvard Medical School, Boston, USA. Professor at the University of California, Berkeley, USA and Investigator, Howard Hughes Medical Institute.

*“for the development of a method for genome editing”*

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