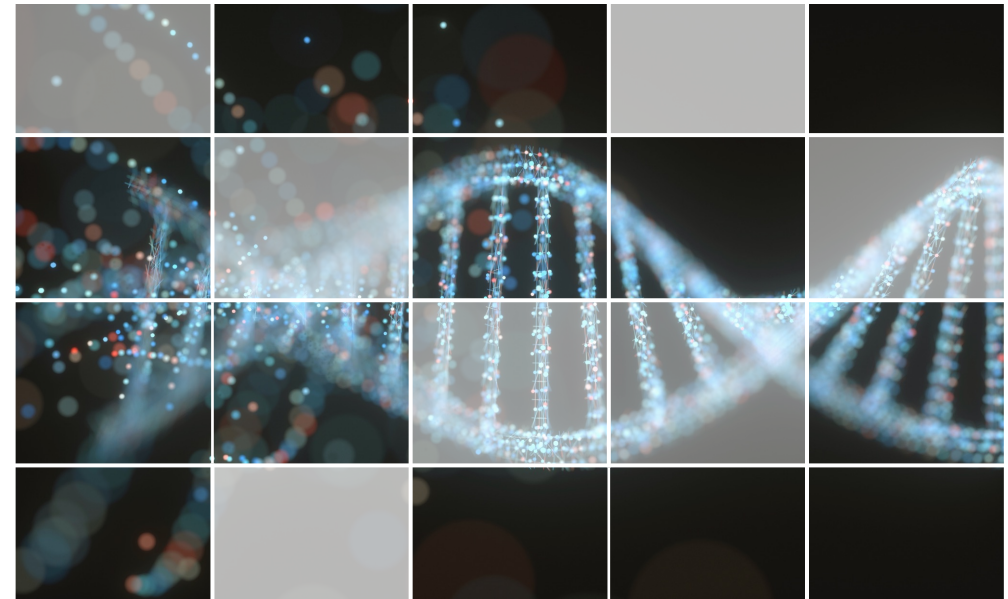




# Hipolit's Biotech Breakdown: **Gene Editing 101 & CRISPR-Cas9**



**Hipolit Cichocki 2022.08**  
**Dragon Gate Investment Partners LLC**

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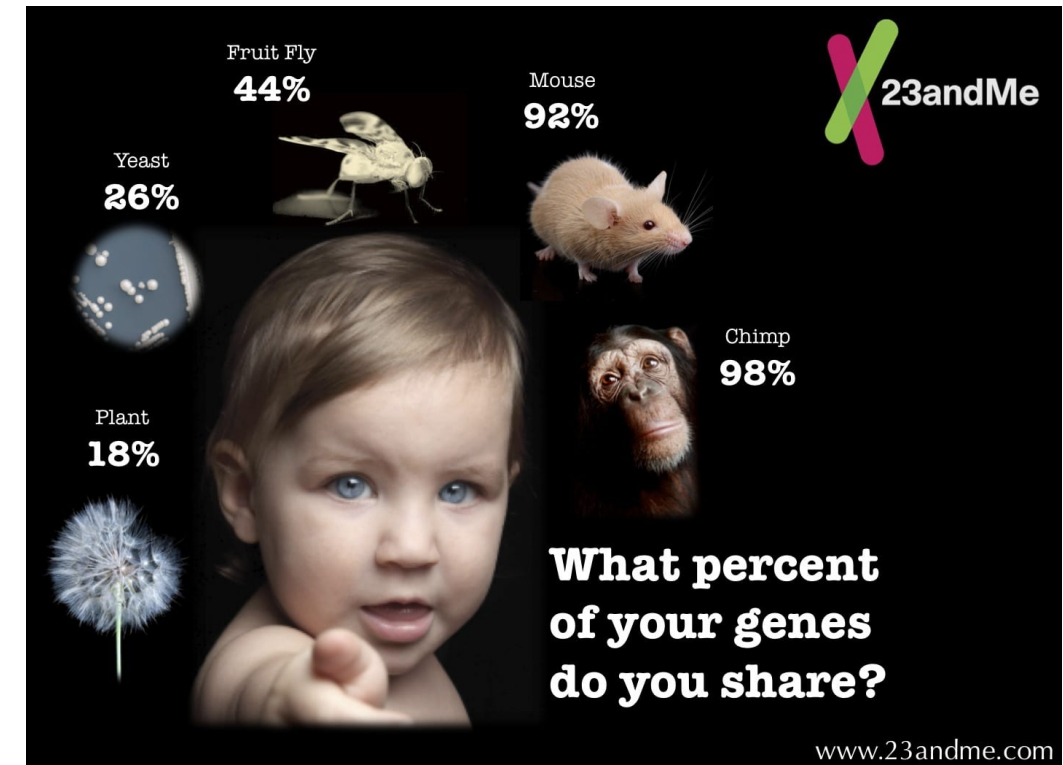
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# Gene Editing Overview



# Every Living Thing Has Genes

- **What do genes and DNA do?**
  - **DNA**- deoxyribonucleic acid contains the biological information that makes every living thing unique
  - **Genes are made of DNA**, which carries the instructions for us to develop, survive, reproduce, and make proteins which do most of the work in our bodies.
  - **Genes are a basic unit of inheritance.**
- **Can our DNA change over time?**
  - As we age, some of our genes experience epigenetic changes or chemical variations that can turn genes off or on without changing information.
  - Over long periods of time our DNA can change through mutation and natural selection.
- **Is it possible to edit DNA?**
  - **Scientists are now able to make changes in DNA** to solve problems.



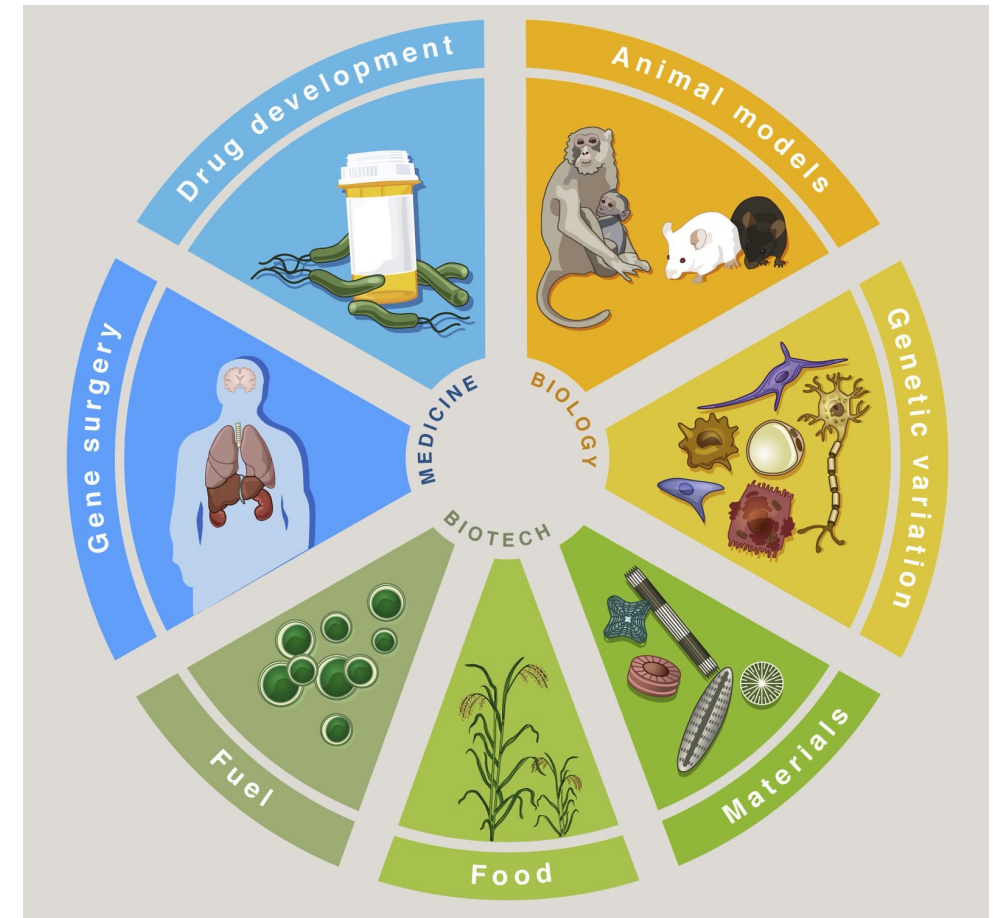
Source: 23andMe.



# The World of DNA

- **Why Does DNA Matter?**
  - **Understanding DNA and gene editing is important.** It allows us to better understand what we already have and have the knowledge needed to help shape the growing field.
- **What can be done with DNA?**
  - DNA has become something that scientists are able to decipher, identify specific traits carried within it, and even edit.
- **Potential Impacts**
  - Cell and gene therapies
  - Diagnostics
  - Agriculture
  - Bioenergy

*"You could turn someone into a freakin butterfly"– Elon Musk*



Source: Cell: Hsu et al 2014.

# Return of the Woolly Mammoth

- **Can we bring back extinct species?**
  - **Through gene editing, it is possible to bring back extinct species such as the Woolly Mammoth**
- **The Woolly Mammoth project**
  - **George Church, a geneticist at Harvard and MIT, and his team have raised \$15M to bring back the woolly mammoth.**
  - **Church was optimistic that he could rewrite the Asian elephant's DNA to produce something that looks and behaves like a mammoth through CRISPR gene editing.**
  - **By tweaking certain genes to produce denser hair or a thicker layer of fat, the team hopes to create an animal with mammoth-like characteristics.**
  - They are optimistic this can be done within the next few years and hope to have a complete woolly mammoth within the decade.



Source: Flying Puffin via Wikicommons.

# Reading Our DNA: The Human Genome Project





# Key Definitions (Human Genome Project)

## Human Genome Project

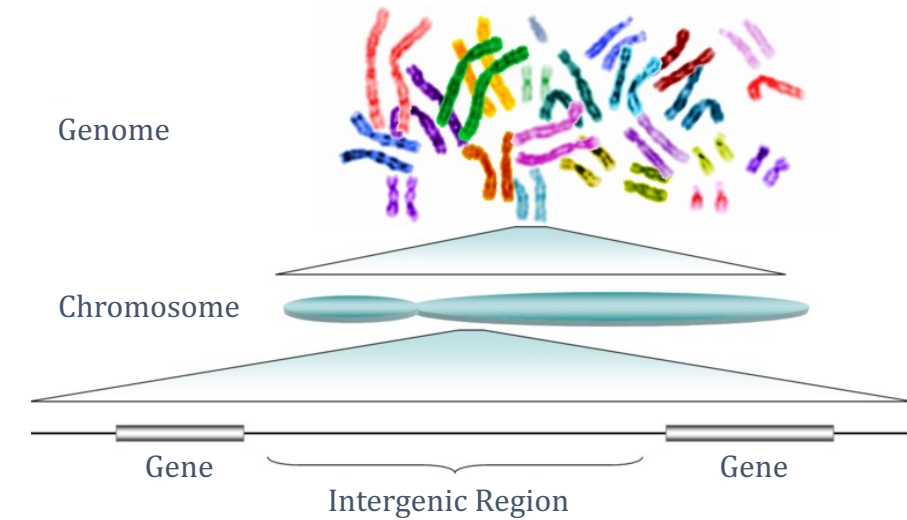
- The Human Genome Project (HGP) was an international, collaborative research program whose goal was the complete mapping and understanding of all human genes.

## Dimer

- A molecule composed of two identical, simpler molecules.

## Genome

- A genome is an organism's complete set of deoxyribonucleic acid (DNA), a chemical compound that contains the genetic instructions needed to develop and direct the activities of every organism.



Source: SITN: Harvard University.



## Key Definitions Continued (Human Genome Project)

### Chromosomes

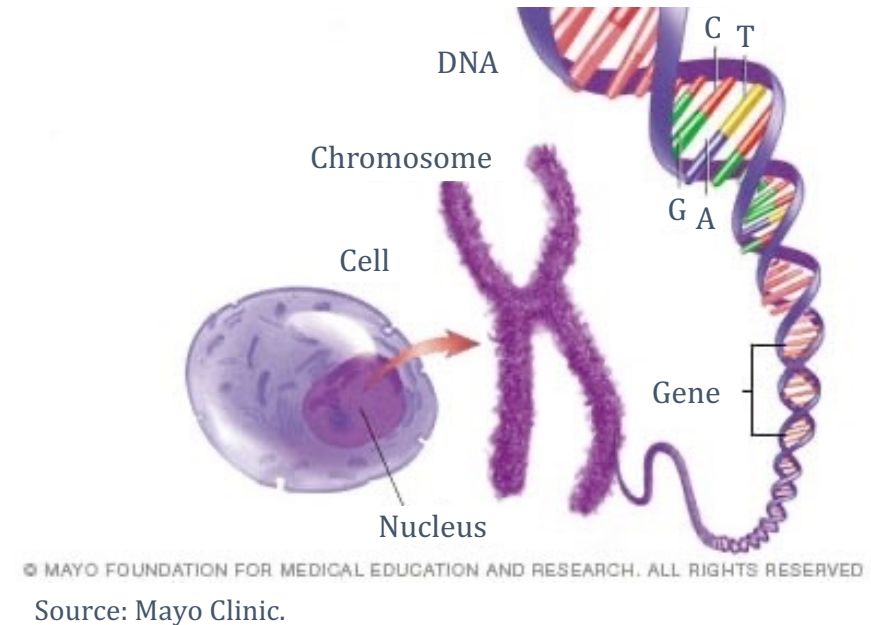
- Threadlike structures made of protein and a single molecule of DNA that serve to carry the genomic information from cell to cell. Humans have 22 pairs of numbered chromosomes (autosomes) and one pair of sex chromosomes (XX or XY), for a total of 46.

### Genes

- The gene is considered the basic unit of inheritance. Genes are passed from parents to offspring and contain the information needed to specify physical and biological traits. Most genes code for specific proteins or segments of proteins, which have differing functions within the body.

### Nucleotide bases

- Nucleotides are the four chemical units that make up the strands in a DNA molecule. The bases are adenine (A), thymine (T), guanine (G) and cytosine (C). Bases on opposite strands pair specifically; an A always pairs with a T, and a C always with a G.





# Human Genome Project

- **Project mission**

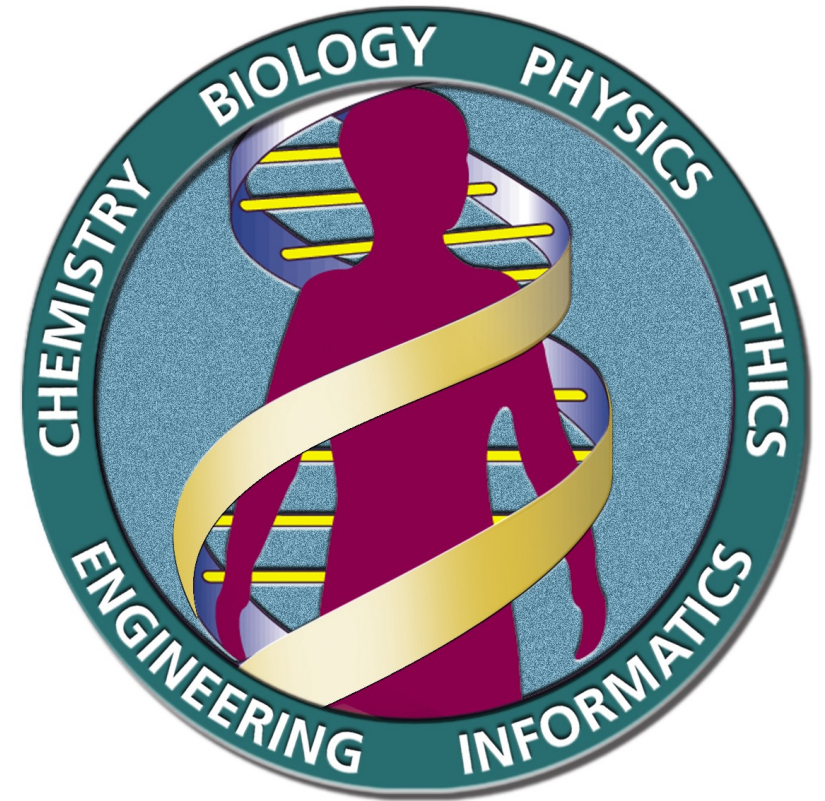
- Decipher the chemical sequence of the entire genome, identify all 50,000 to 100,000 genes contained within the genome, and provide the research tools needed to analyze all this genetic information.

- **Isolating and analyzing the genetic material**

- At the time, this study could potentially provide scientists with powerful new approaches to understand disease development and to create new strategies for their prevention and treatment.

- **A great scientific undertaking**

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



Source: U.S. Department of Energy, Human Genome Project.



# Timeline

1984

Early meetings **assess the feasibility of a Human Genome Project.**

1988

The **main goals** of the Human Genome Project were **first articulated by a special committee of the U.S. National Academy of Sciences**, and later adopted through a detailed series of five-year plans jointly written by the National Institutes of Health (NIH) and the Department of Energy (DOE).

1990

The **initial planning stage was completed** with the publication of a joint research plan, "Understanding Our Genetic Inheritance: The Human Genome Project, The First Five Years, FY 1991-1995." This initial research plan set out specific goals for the first five years of what was then projected to be a 15-year research effort.

1995

Human Genome Project **researchers publish a physical map of the human genome.**

1999

The Human Genome Project **successfully completes the pilot phase of sequencing the human genome.** Human Genome Project **researchers decode the DNA sequence of the first human chromosome.**

2003

The Human Genome **Project is completed.**

2004

The **International Human Genome Sequence Consortium publishes their finished human genome sequence.**

# Human Genome Project Contributors and Findings

- **Project Contributors**

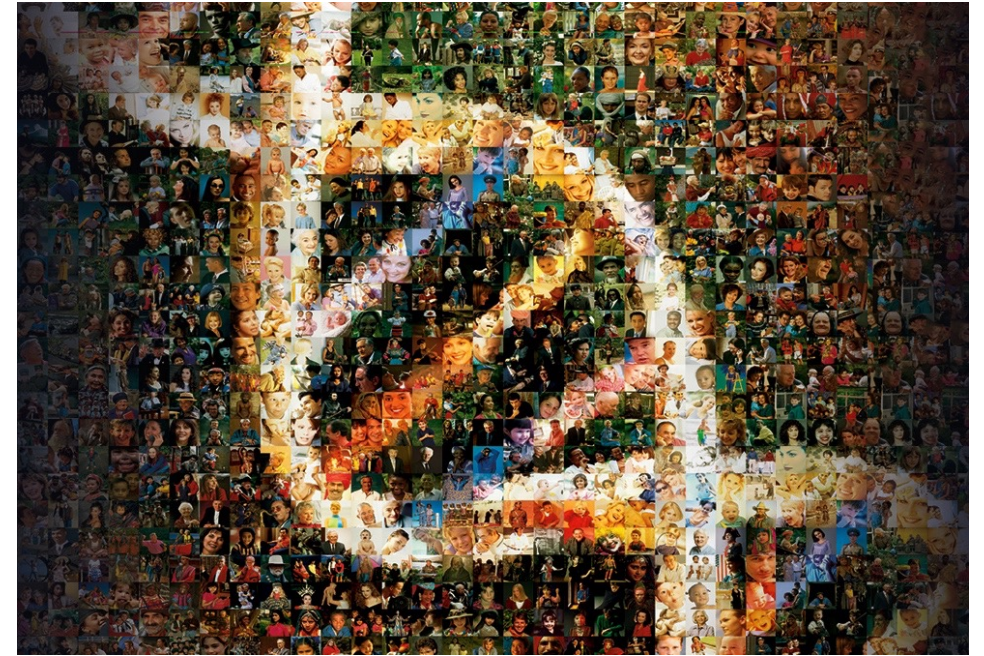
- Contributors include the NIH, DOE, and numerous universities and research centers throughout the US, UK, France, Germany, Japan, and China.

- **What was accomplished?**

- The finished sequence covers about 99 percent of the human genome's gene-containing regions has been sequenced to an accuracy of 99.99 percent.
- In 2003, an accurate and complete human genome sequence was finished and made available to scientists and researchers two years ahead of the original schedule and under the original estimated budget.

- **Completing the last 8%**

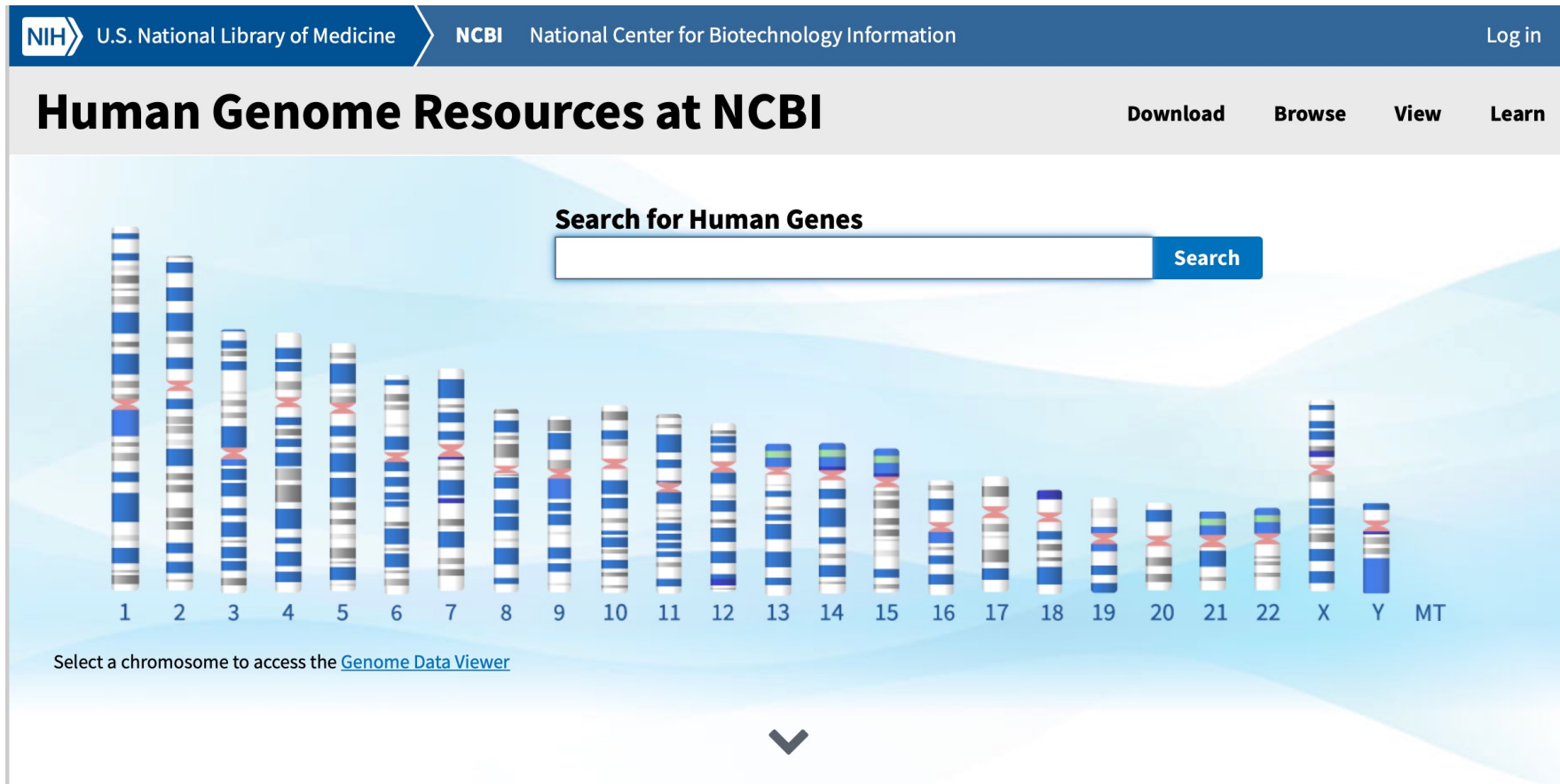
- About 8% of our genome was not sequenced due to technological limitations. The T2T (Telomere 2 Telomere) team took on this challenge and recently completed it, with 6 papers being posted on Science in 2022.



Source: NIH National Human Genome Research Institute.

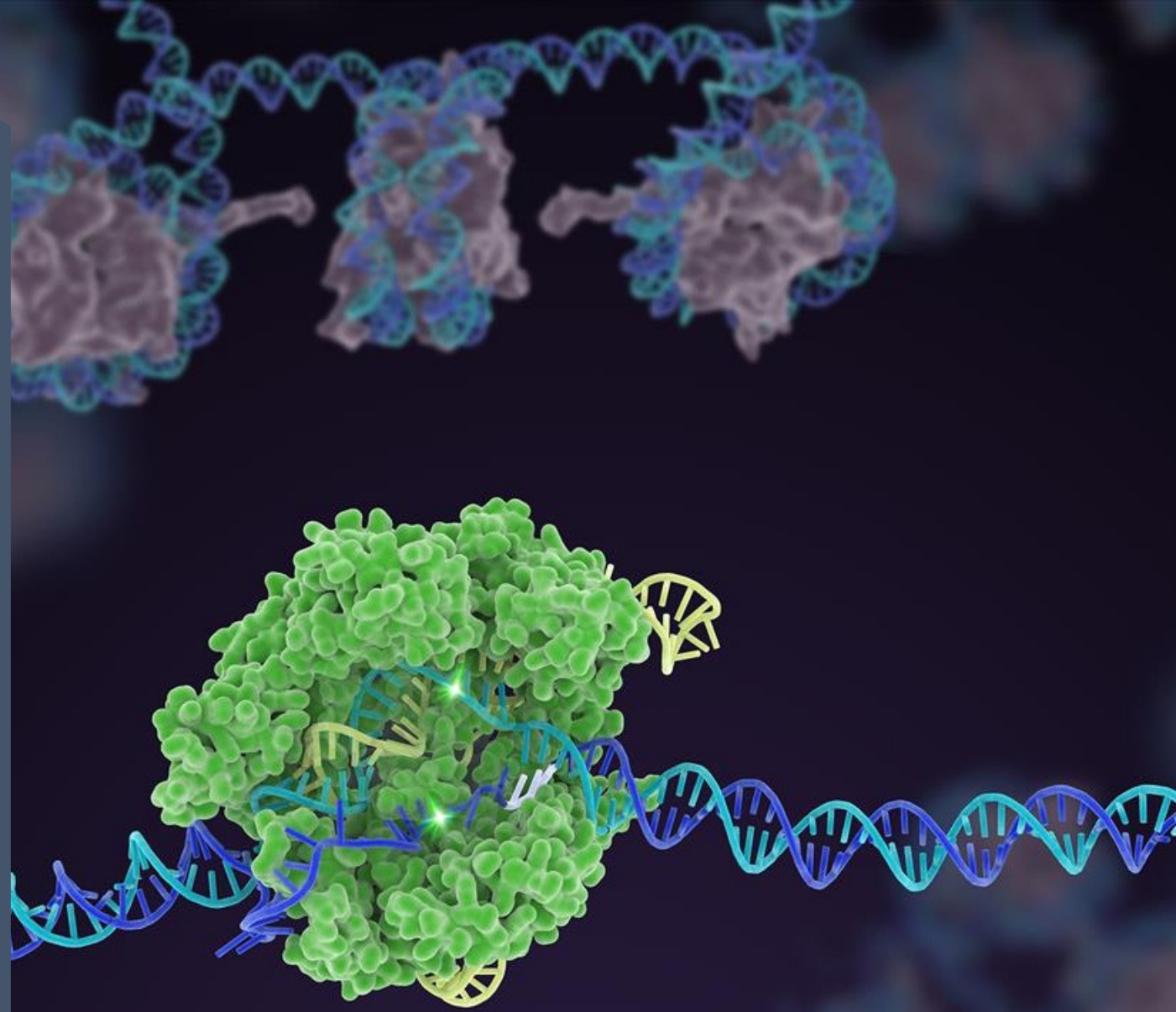
# Accessing the Information

- The information is accessible on the NIH website: <https://www.ncbi.nlm.nih.gov/projects/genome/guide/human/>



The screenshot shows the "Human Genome Resources at NCBI" webpage. At the top, there is a navigation bar with the NIH logo, "U.S. National Library of Medicine", the NCBI logo, "National Center for Biotechnology Information", and a "Log in" link. Below this is a header section with the title "Human Genome Resources at NCBI" and four action links: "Download", "Browse", "View", and "Learn". The main content area features a large graphic of human chromosomes, numbered 1 through 22, plus X, Y, and MT. Above the chromosomes is a search bar with the text "Search for Human Genes" and a "Search" button. Below the chromosomes, there is a link that says "Select a chromosome to access the [Genome Data Viewer](#)". A large downward-pointing arrow is centered at the bottom of the page.

# CRISPR-Cas9: Genetic Scissors





## Definitions (CRISPR-Cas9)

### CRISPR-Cas9

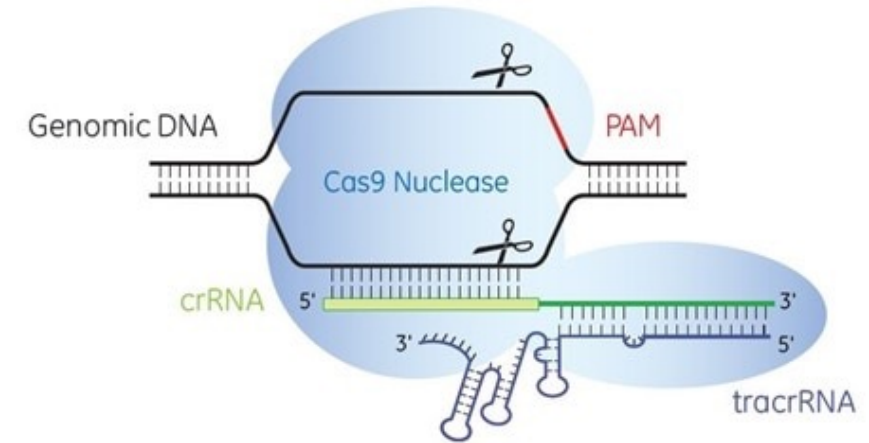
- CRISPR-Cas9 is a gene editing tool that was adapted from a naturally occurring genome editing system that bacteria use as an immune defense.

### Guide RNA

- A two-piece molecule that Cas9 binds and uses to identify a complementary DNA sequence. Composed of the CRISPR RNA (crRNA) and trans-activating CRISPR RNA (tracrRNA). Cas9 uses the tracrRNA portion of the guide as a handle, while the crRNA spacer sequence directs the complex to a matching DNA sequence.

### Endonuclease

- An enzyme that breaks down a nucleotide chain into two or more shorter chains by cleaving the internal covalent bonds linking nucleotides.



Source: horizon A PerkinElmer company.



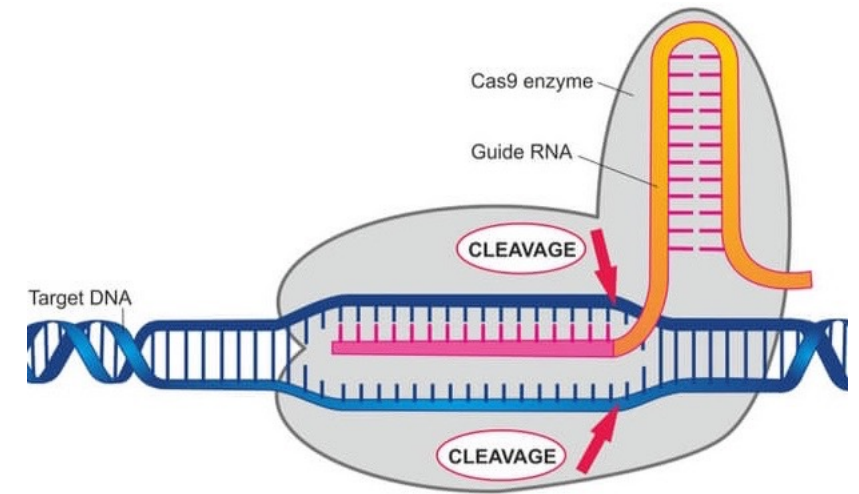
## Definitions Continued (CRISPR-Cas9)

### Enzyme

- A substance that acts as a catalyst in living organisms, regulating the rate at which chemical reactions proceed without itself being altered in the process.

### RNA

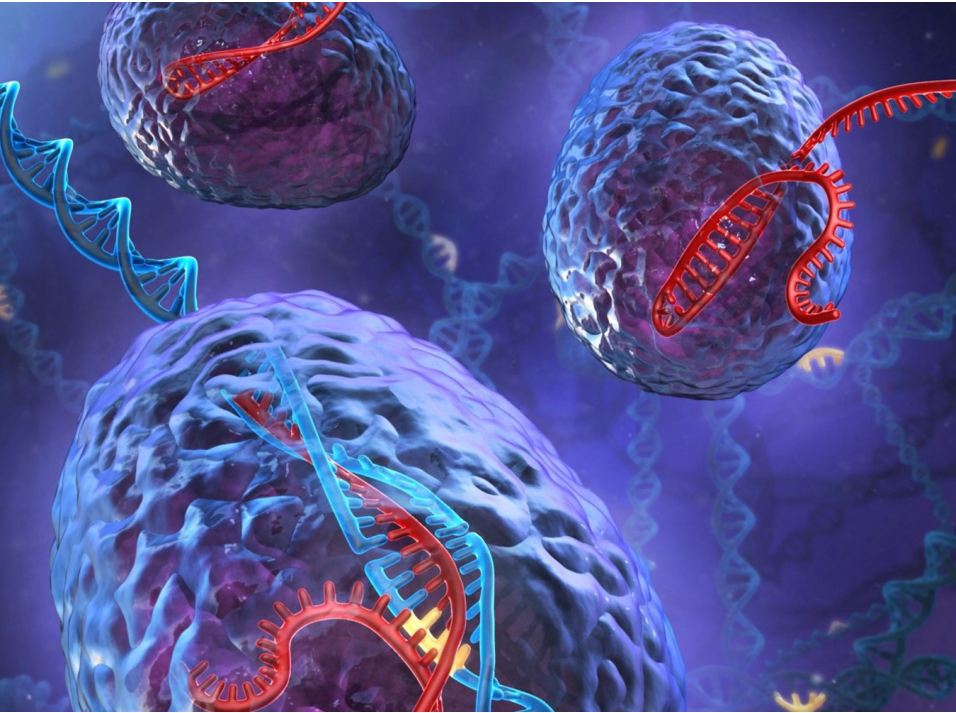
- Ribonucleic acid contains information that has been copied from DNA. Cells make several different forms of RNA, and each form has a specific job in the cell. Many forms of RNA have functions related to making proteins. RNA is also the genetic material of some viruses instead of DNA.



Source: Cell: LABIOTECH.eu.



## CRISPR-Early Discovery



Source: Stephen Dixon.

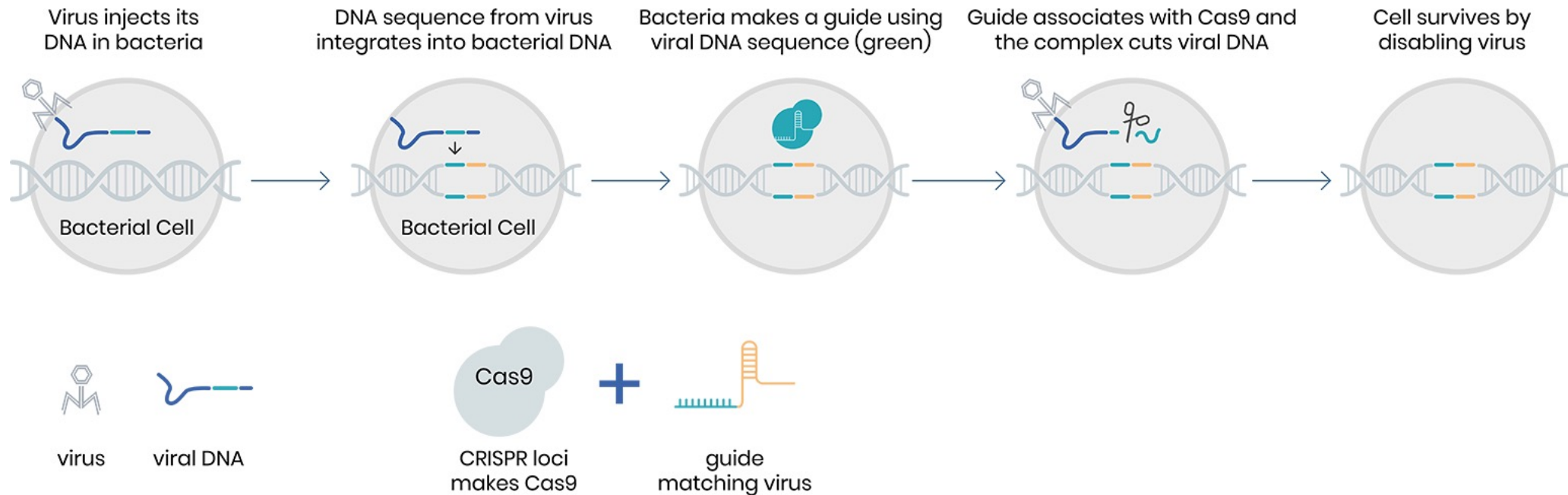
- **CRISPR was first observed in 1987.**
  - Researchers in Japan noticed a weird, repeating sequence **in the DNA of E. coli bacteria.**
- **Later studies found repeating segments of DNA in other microbial species.**
  - These mysterious repeats consisted of a **short sequence of genetic code and a similar sequence in reverse.** This peculiar **palindrome pattern was dubbed CRISPR** — “clustered regularly interspaced short palindromic repeats.”
  - **Further research led to the discovery of CRISPR-associated (Cas) genes, which produce Cas enzymes that can slice through DNA.**
  - Scientists eventually realized that bacteria have been using CRISPR-Cas complexes for billions of years to attack and destroy enemy viruses, and that this ancient bacterial immune system could be adapted for use in genetic engineering.
- **International recognition**
  - In 2020, Emmanuelle Charpentier and Jennifer Doudna were awarded the **Nobel Prize** in Chemistry for their work on CRISPR-Cas9.



# How Does CRISPR Work?

- **Bacterial immune system**

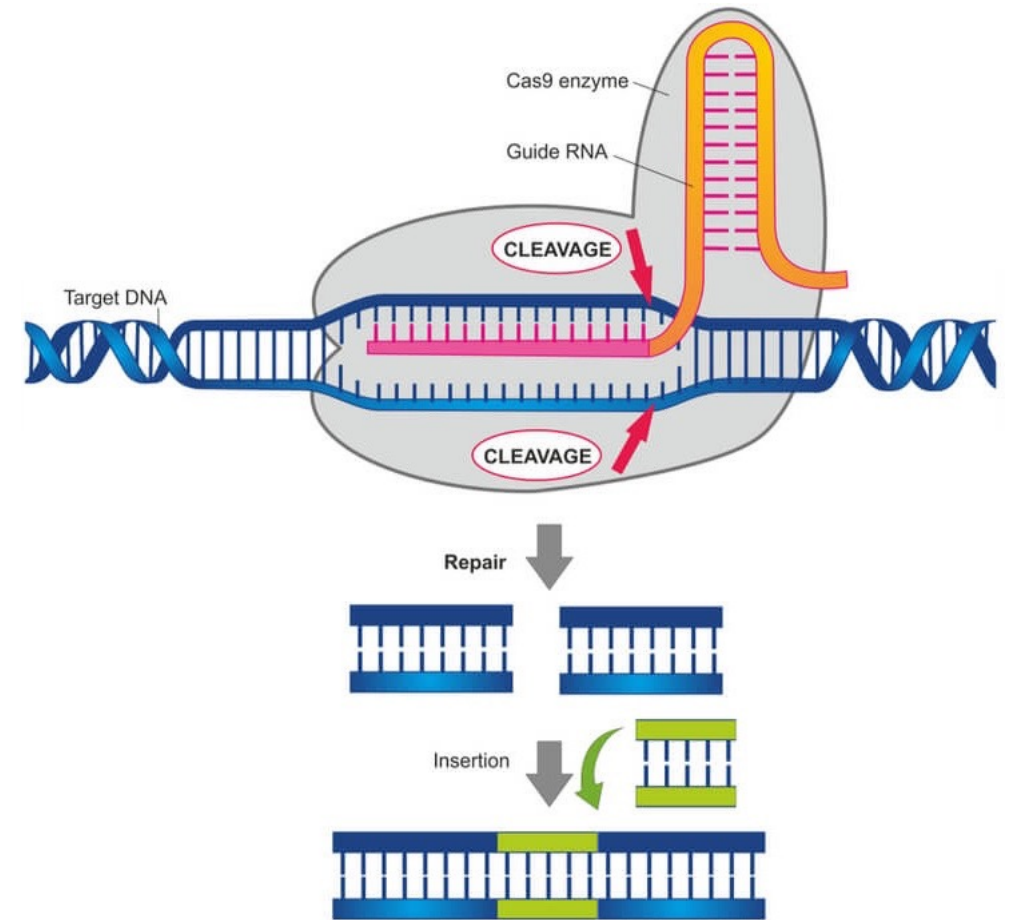
- In bacteria, **CRISPR-Cas9 is a naturally occurring genome editing system that bacteria use as an immune defense.**
- The patterns of integrated viral DNA are known as CRISPR arrays and help bacteria remember the virus, in case of another attack. **Researchers adapted this immune defense system to edit DNA.**



Source: Intellia Therapeutics.

# CRISPR-Cas9 Gene Editing Overview

- **Design an RNA molecule**
  - The molecule matches the mutated DNA sequence in the gene.
- **Combine the RNA with a Cas9 enzyme**
  - Cas9 can cut through DNA like scissors.
- **The cut is made**
  - The RNA acts like a very fast GPS — it guides the Cas9 enzyme to the mutated DNA sequence. The enzyme then binds to the sequence and deletes it.”
- **The final repair is made**
  - Benign virus is engineered to deliver and insert the correct DNA sequence into the edited gene.
  - Use the cell's own DNA repair machinery to add or delete pieces of genetic material, or to make changes to the DNA by replacing an existing segment with a customized DNA sequence



Source: Cell: LABIOTECH.eu.

# CRISPR vs Past Gene Editing Techniques

| Feature                       | CRISPR                                                      | TALEN's                                                             | ZFN's                                          |
|-------------------------------|-------------------------------------------------------------|---------------------------------------------------------------------|------------------------------------------------|
| Construction                  | 22bp + pam sequence                                         | 30-40bp                                                             | 9-18bp                                         |
| Target sequence recognition   | sgRNA, RNA-DNA interactions                                 | Repeat variable diresidues (RVDs) repeats, protein-DNA interactions | Zinc fingers protein, protein-DNA interactions |
| Endonuclease                  | Cas                                                         | FokI                                                                | FokI                                           |
| Endonuclease construction     | sgRNA synthesis or cloning                                  | 8-31RVD repeats                                                     | 3-4 Zinc fingers domains                       |
| Delivery                      | sgRNA complementary to the target sequence with CAS protein | 2 TALENs around the target sequence required                        | 2 ZFNs around the target sequence              |
| DNA sequence recognition size | 17-20bp +NGG *1                                             | (8-31bp) *2                                                         | (9 or 12bp) *2                                 |
| Targeting efficiency          | High                                                        | Moderate                                                            | Low                                            |
| Affordability                 | Highly Affordable                                           | Affordable                                                          | Resource intensive                             |
| Timeliness                    | Rapid                                                       | Time consuming                                                      | Time Consuming                                 |

## Definitions

- **ZFN**- Zinc-finger nucleases are fusions of the nonspecific DNA cleavage domain from the FokI restriction endonuclease with zinc-finger proteins. ZFN dimers induce targeted DNA DSBs that stimulate DNA damage response pathways.
- **TALEN**- Transcription activator-like effector nucleases are fusions of the FokI cleavage domain and DNA-binding domains derived from TALE proteins. TALENs induce targeted DSBs that activate DNA damage response pathways and enable custom alterations.

# CRISPR Gene Editing Procedures



## Step 1: Determine the Genetic Tool to Use

|                                                                                    |                                |                                                                                                                                   |
|------------------------------------------------------------------------------------|--------------------------------|-----------------------------------------------------------------------------------------------------------------------------------|
|  | Gene Knockout                  | <ul style="list-style-type: none"> <li>• NHEJ- non homologous end joining</li> </ul>                                              |
|                                                                                    | Gene Editing                   | <ul style="list-style-type: none"> <li>• HDR-homology directed repair</li> <li>• Base editing</li> <li>• Prime editing</li> </ul> |
|                                                                                    | Gene Activation and Inhibition | <ul style="list-style-type: none"> <li>• CRISPRa- CRISPR activation</li> <li>• CRISPRi- CRISPR inhibition</li> </ul>              |
|                                                                                    |                                |                                                                                                                                   |





## Step 2: Identify the Target Sequence

### Direct Sequencing

- Identify each individual base pair, in sequence, and compare the sequence to a normal gene.

### DNA Hybridization Models

- Use of the strong binding of a lone DNA strand to a strand whose sequence is the perfect complement of the first. Binding can differentiate between normal and mutated DNA.

### Restriction Enzyme Digestion or Hybridization

- Restriction enzymes are specialized enzymes that recognize and cut the DNA wherever they encounter a specific, very short sequence.



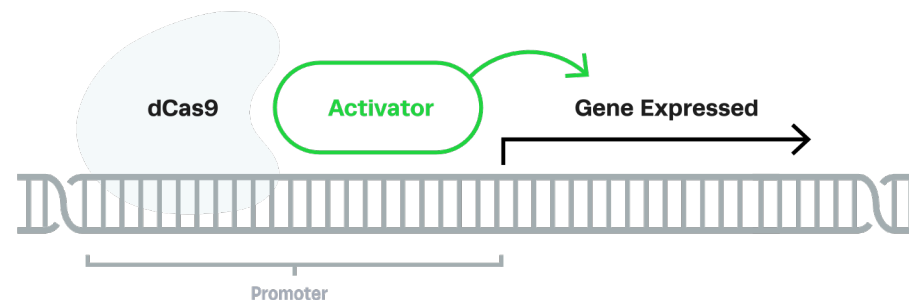
## Step 3: Create the Guide RNA for (CRISPRa & CRISPRi)

**Note: Perfect gRNA does not exist.**

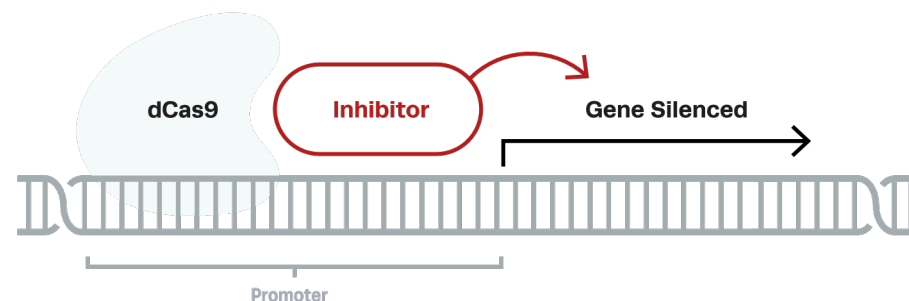
### Designing Guide RNAs for CRISPRa & CRISPRi

- The critical parameters for gRNA design are different for CRISPRa and CRISPRi.
- These tools involve gene activation or inhibition by targeting the promoter of the target gene.
- The location target DNA range for guide RNA is therefore quite narrow, requiring a balance of complementary sequence and optimized location while designing.

#### CRISPRa



#### CRISPRi

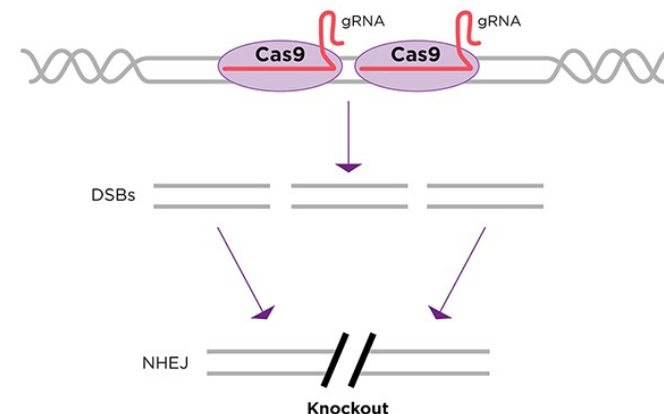


Source: Synthego.

## Step 3: Create the Guide RNA for (CRISPR Knockouts & Knock-ins)

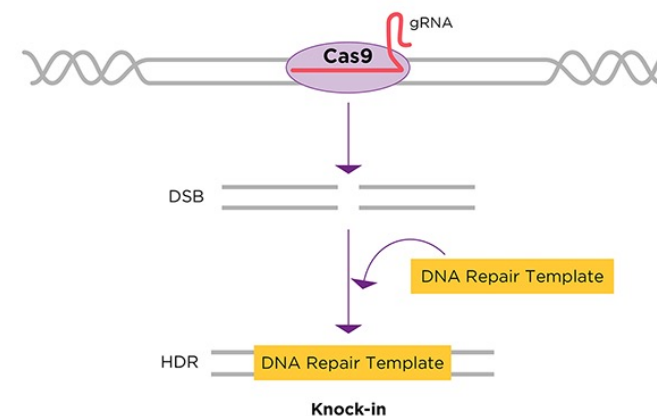
### Designing gRNA for Gene Knockouts

- CRISPR-mediated knockouts are commonly done through the non-homologous end joining (NHEJ) repair pathway.
- In this method, the Cas creates a double-strand break (DSB). The DSB is fixed by NHEJ.
- **This pathway** often results in insertion or deletion (indel) of one or more nucleotides at the break, which **leads to frameshift mutations that completely knock out gene function.**



### Guides for CRISPR Knock-ins

- A knock-in is done by inserting a new DNA fragment into the genome.
- **Researchers use** a more sophisticated repair pathway, known as the homology-directed repair (HDR). In HDR, the cell copies the sequence of donor DNA template to fill the broken piece accurately.
- During knock-ins, researchers supply a surplus of template DNA containing the desired genomic change, improving the chance that the cell will pick this DNA for repair.



Source: Taconic Biosciences.



## Step 4: Choose a Tool to Design Guide RNA (Tool List)

Geneious

Benchling

Guides

Broad  
Institute GPP

Horizon  
Discovery

CasOFFinder

IDT

CHOPCHOP

Off-Spotter

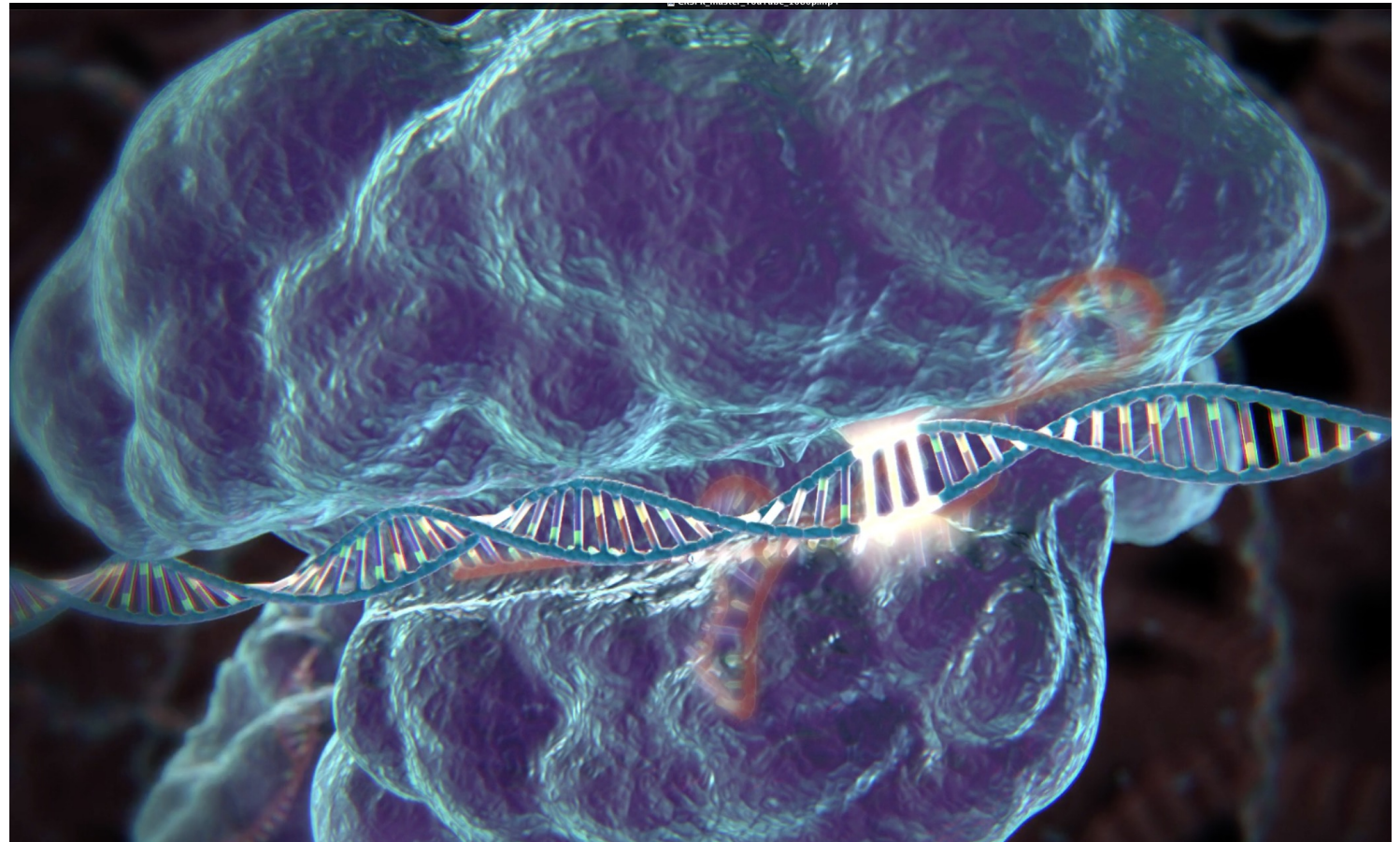
CRISPOR

Synthego

Deskgen

TrueDesign

E-CRISP

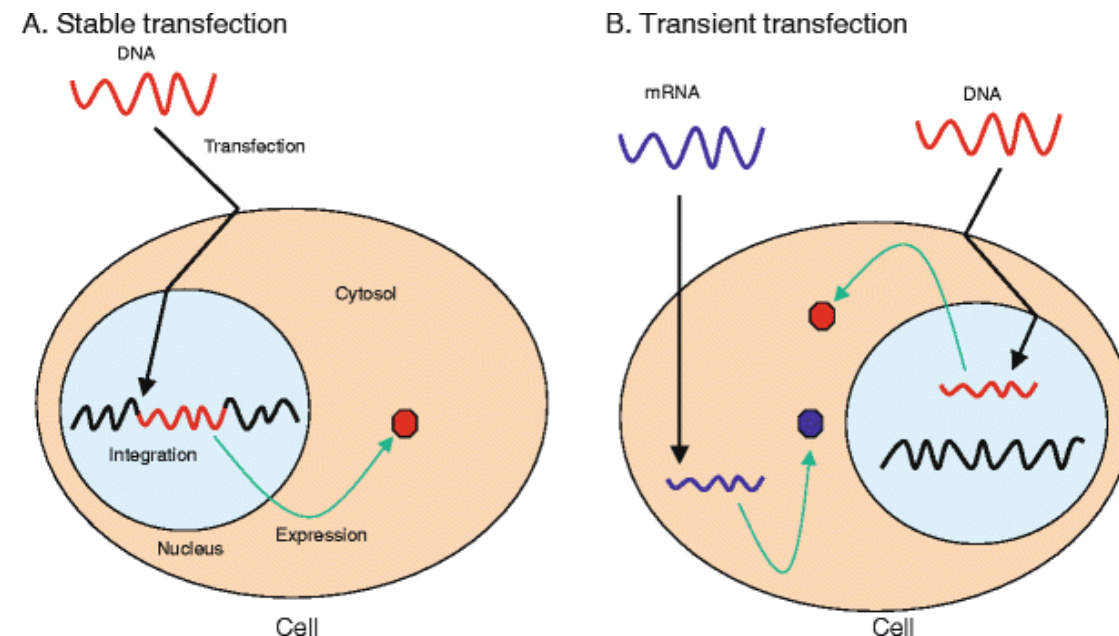


Source: McGovern Institute.

## Step 5: Insert CRISPR to the Target

### • Transient vs. Stable Transfection

- The method depends if the goal is permanent or temporary expression of Cas9 and/or the gRNA in cells.
- **Transient transfection: CRISPR components are introduced into the cell, but no DNA encoding gRNA or Cas9 are incorporated into the cell's genome.**
  - CRISPR-Cas9 can only cleave the DNA for a limited time.
- **Stable transfections: most effectively achieved by using a viral vector but can occur at low frequency after introducing plasmid DNA.**
  - Typically, DNA encoding Cas9 (but not guide RNA) is transfected to produce a Cas9-expressing cell line. Then, gRNAs can be introduced to Cas9-expressing cells through transient transfection.
- **Format of CRISPR Components-** Format of CRISPR components may affect which transfection method to use.
  - DNA
  - RNA
  - Ribonucleoprotein complex (RNP)



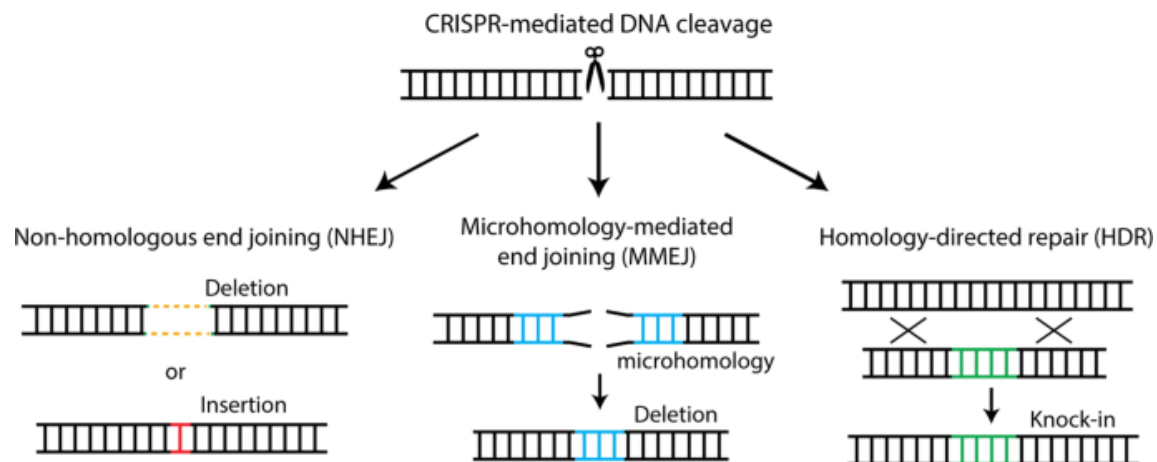
Source: Springer Link: Mammalian cell transfection: the present and the future.

## Step 5 Continued: Physical Transfection of CRISPR

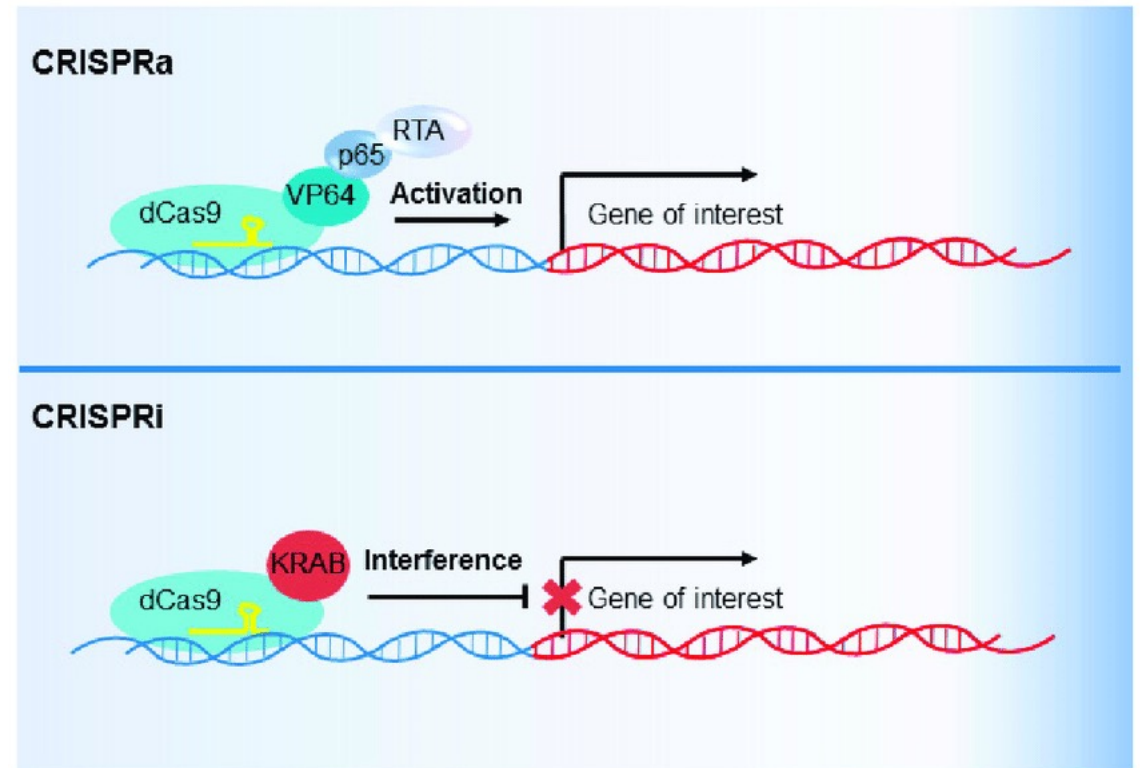
|             | Lipofection                                                    | Electroporation                                                      | Nucleofection                                                                             | Microinjection                                                          | Virus                                                                   |
|-------------|----------------------------------------------------------------|----------------------------------------------------------------------|-------------------------------------------------------------------------------------------|-------------------------------------------------------------------------|-------------------------------------------------------------------------|
| Principle   | Lipid complexed with genetic material fuses with cell membrane | Electric pulse forms pores in cell membrane for entry of DNA/RNA.RNP | An electrorotation-based optimized for nuclear delivery, pre-optimized for each cell type | Microneedle injects CRISPR components inside cells, oocytes, or zygotes | DNA/RNA is packaged into infectious particles and introduced into cells |
| Advantages  | Cost-effective, high throughput                                | Easy, fast, high efficiency                                          | Easy, fast, high efficiency                                                               | High efficiency                                                         | High efficiency                                                         |
| Limitations | Less efficient                                                 | Requires optimization                                                | Requires reagents and equipment                                                           | Time-consuming, technically demanding, low throughput                   | Time-consuming, safety requirements, expensive                          |
| Cell Types  | Few                                                            | Numerous                                                             | Numerous                                                                                  | Few                                                                     | Numerous                                                                |

## Step 6: Fixing the Gene

- Once the gene is cut, the gene error is resolved through
  - A knockout
  - Edit
  - Activation or inhibition



Source: Nature: Current trends in gene recovery mediated by the CRISPR-Cas system.



Source: Researchgate: CRISPR-Cas9 Genetic Analysis of Virus-Host Interactions.



## Step 7: Check the Results

### CRISPR Analysis Using Next-Generation Sequencing (NGS)

- Pros- Gold standard, extremely sensitive for detecting editing outcomes, high-throughput sequence-based data provides a comprehensive view of the indels (insertions or deletions) generated
- Cons- Time and labor intensive, expensive, best with large samples

### Inference of CRISPR Edits (ICE) from Synthego

- Pros- New, user-friendly, online tool that uses Sanger sequencing data to determine the relative abundance and levels of indels due to CRISPR editing.
- Cons- Deletion and insertion detection limitations, Sanger sequencing affects sensitivity, assumes edits use spCas9

### Tracking of Indels by Decomposition (TIDE)

- Pros- Lower cost alternative to NGS, Sanger sequencing, like ICE, TIDE software aligns your sgRNA sequence to unedited and edited samples and conducts a comparative analysis
- Cons: modifying parameters in model can be difficult for user

### T7 Endonuclease 1 (T7E1) Assay

- Pros- Does not use DNA sequencing to determine CRISPR editing efficiency, takes advantage of the T7 endonuclease, which preferentially cleaves mismatched DNA, the small fragments can be viewed by agarose gel electrophoresis.
- Cons: not quantitate, no info on different indel sequences generated



# Minimizing Off-Target Effects

## Ensure On-Target Activity of Guide RNA

- **Doench et al.** analyzed the specificity of thousands of guide RNAs while creating genome-wide mice and human libraries. Using sophisticated computational biology tools, the team **established scoring rules to predict the on-target activity of gRNAs**, and the “Doench rules” are now implemented in several online gRNA design tools.

## Minimize gRNA Off-Target Effects

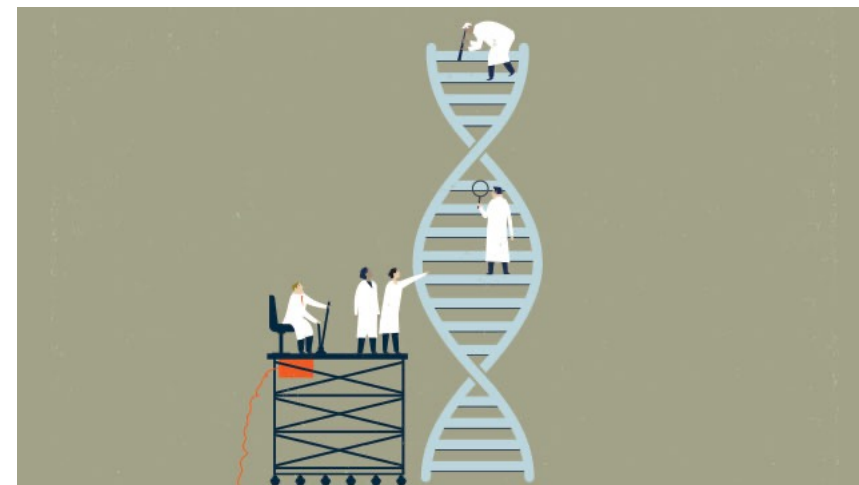
- The **Doench study** also yielded another useful metric: the **off-target activity score**. This score of a designed gRNA indicates how likely it is that the gRNA will bind non-intended targets.

## Multiple gRNAs to Improve CRISPR KO

- Multiple gRNAs targeting the same gene have been shown to improve editing efficiency and greatly increase the chance of generating a knockout.

## Choose the Best CRISPR Design Tool

- Details explained in pg. 26





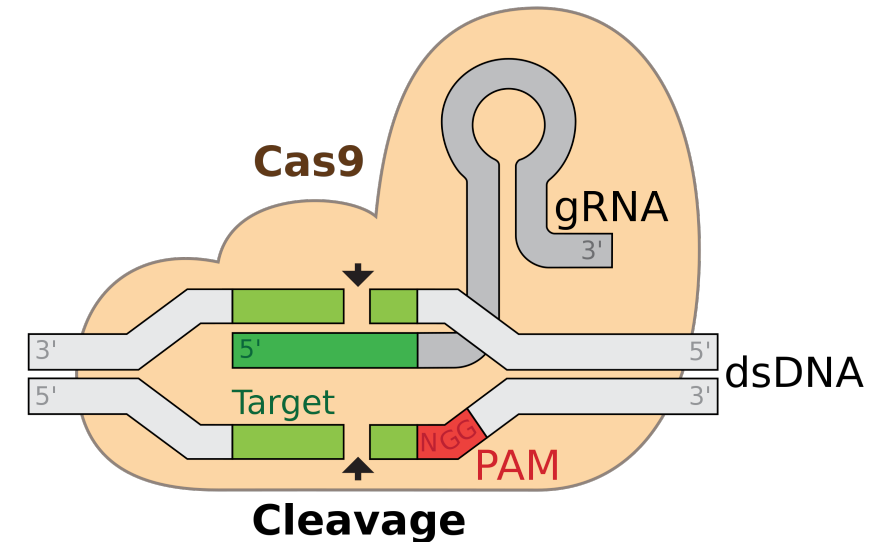
# Guide RNA, PAM, & CAS

## Guide RNA

- **Cas9 is guided** to its target sites with **by two RNAs: tracrRNA and crRNA**
- crRNA defines the genomic target for Cas9 and tracrRNA links the crRNA to Cas9 and facilitates processing of mature crRNAs from pre-crRNAs derived from CRISPR arrays.
- **In most CRISPR systems, two RNAs have been condensed into one RNA known as the guide RNA (gRNA) or single guide RNA (sgRNA).**

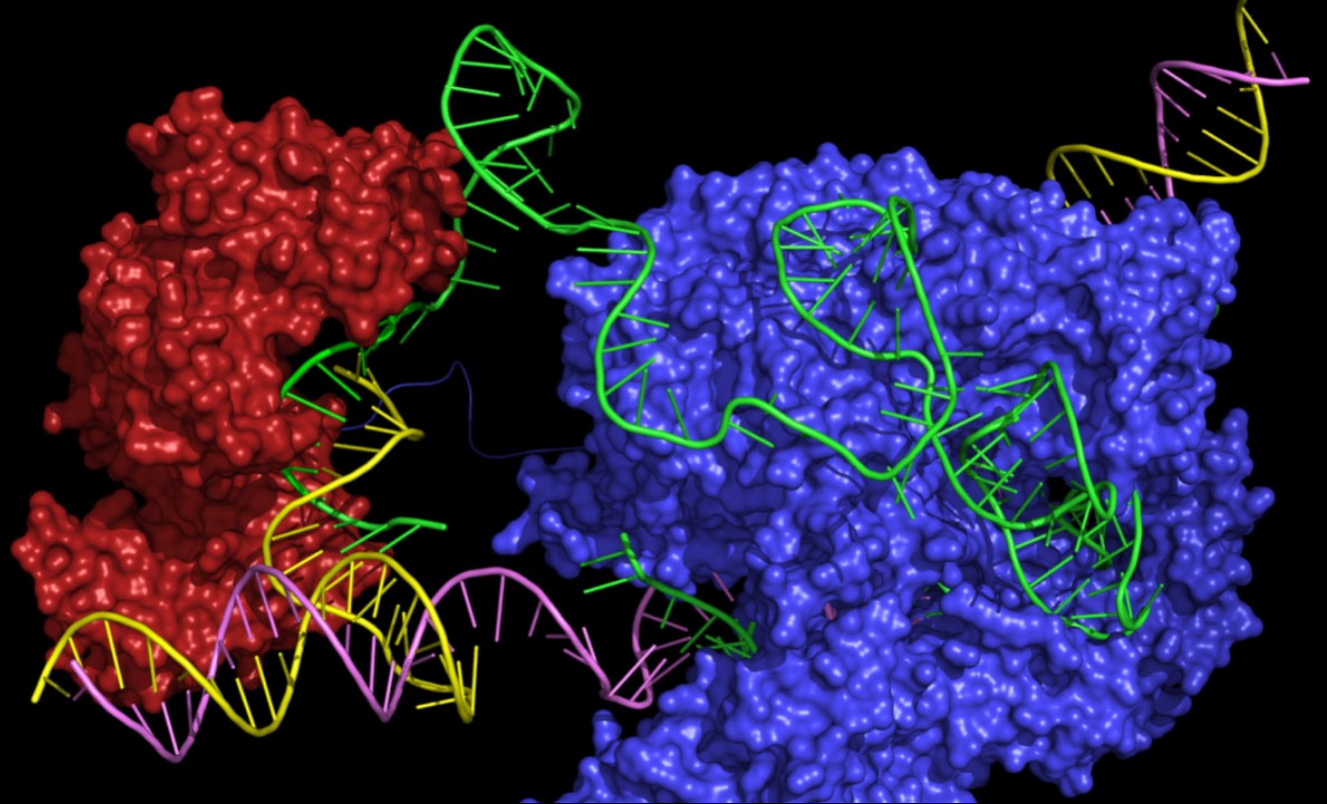
## CAS Endonuclease

- **Cas9 binding to the targeted is mediated both by the gRNA and a 3-base pair sequence known as the Protospacer Adjacent Motif or PAM.**
- For DNA to be cut by Cas9, it must contain a PAM sequence immediately downstream (3') of the site targeted by the gRNA. In the absence of either the gRNA or PAM sequence, Cas9 will neither bind nor cut the target.
- Depending on the type of editing tool and PAM sequence, the appropriate endonuclease is chosen.



Source: Wikipedia: Marius Walter.

# Other Gene Editing Techniques



# Base Editing vs CRISPR vs Prime Editing

| Feature                     | CRISPR Nuclease                        | Base Editing                | Prime editing                                        |
|-----------------------------|----------------------------------------|-----------------------------|------------------------------------------------------|
| Size                        | ~4 kilobase (kb)                       | ~5kb                        | ~6kb                                                 |
| PAM Dependence              | High                                   | High                        | Low                                                  |
| Endonuclease                | Cas                                    | d/nCas9                     | Cas                                                  |
| Breaks                      | Double strand break (DSB)              | Single strand nick          | Single strand nick                                   |
| Ideal Use                   | Very large DNA insertions or deletions | Correcting point mutations  | Multi letter mutations and somewhat longer sequences |
| Target Sequence Recognition | sgRNA, RNA-DNA interactions            | sgRNA, RNA-DNA interactions | pegRNA, RNA-DNA interactions                         |

## Definitions

- **Base editing**- CRISPR-Cas9-based genome editing technology that allows the introduction of point mutations in the DNA without generating DSBs. Two major classes of base editors have been developed: cytidine base editors or CBEs allowing C>T conversions and adenine base editors or ABEs allowing A>G conversions.
- **Prime editing**- The more versatile prime editing (PE) system contains a prime editing extended guide RNA (pegRNA)-guided reverse transcriptase instead of a deaminase. The development of PE was a breakthrough as it requires no PAM sequence adjacent to the target site and it can accomplish not only all 12 types of point mutations, but also insertions (of up to 44 bp) and deletions (of up to 80 bp), or even combination of substitutions, insertions and deletions.

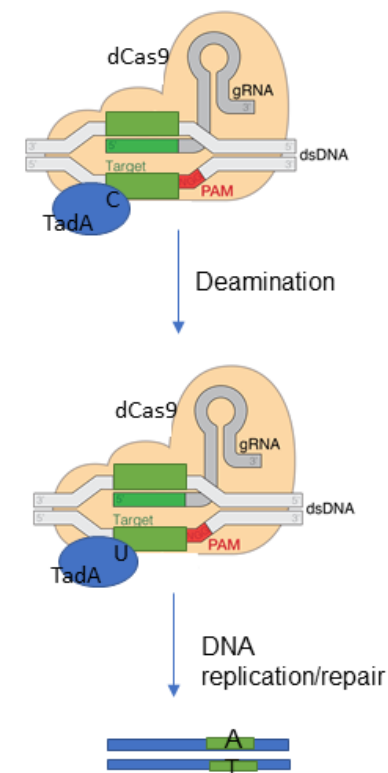
# How Does Base Editing Work?


 DNA base editors (BEs) are made of fusions between a catalytically impaired Cas nuclease and a base-modification enzyme that operates on a single strand DNA.


 Upon binding to its target in DNA, base pairing between the guide RNA and target DNA strand leads to displacement of a small segment of single-stranded DNA in an “R-loop”.


 DNA bases within this single-stranded DNA bubble are modified by the deaminase enzyme.


 To improve efficiency in eukaryotic cells, the nuclease also generates a nick in the non-edited DNA strand, inducing cells to repair the non-edited strand using the edited strand as a template.



DNA base editing

Source: MATER METHODS 2019;9:2800.

# How Does Prime Editing Work?

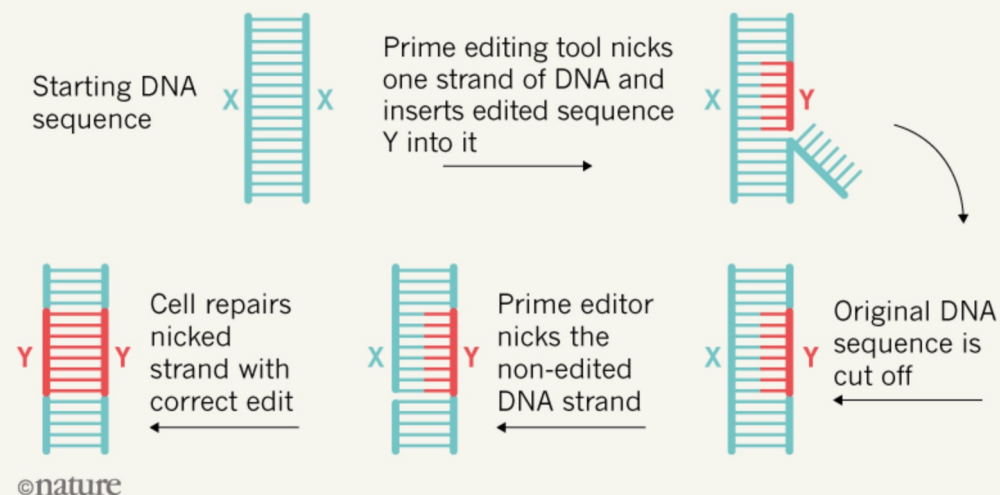
Prime Editing Works by nicking DNA at a specific point in the genome

Although it also **uses Cas9** to recognize specific DNA sequences — just like CRISPR-Cas9 does — the **Cas9 enzyme in Prime editing is modified to nick only one DNA strand.**

Then, a second enzyme called reverse transcriptase, guided by a strand of RNA, makes the edits at the sight of the cut.

## PRECISION EDITOR

Prime editing reduces the number of unintended changes to a genome by inserting the edits researchers want to make into the DNA itself. This contrasts with CRISPR-Cas9, which relies on the cell's repair system to make the changes.



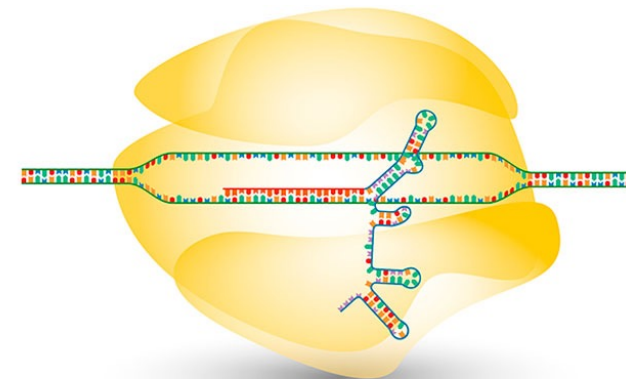
# Prime Editing Advancements

## Prime shows promise

- Prime editing has shown promise in reducing ‘off-target’ effects that are a key challenge for some applications of the standard CRISPR-Cas9 system.
- This could make prime gene therapies safer for use in people.

## Prime editing seems capable of making a wider variety of edits.

- One day, it can be used to treat the many genetic diseases that have so far stumped gene-editors.



Source: teubio.



# Where Prime Prevails

## Crispr drawbacks

- **CRISPR-Cas9 breaks both strands of DNA and then relies on the cell's own repair system** to patch the damage and make the edits, however, **that repair system is unreliable.**
- It can insert or delete DNA letters at the points where the genome was cut potentially leading to an uncontrollable mixture of edits that vary between cells.
- Although researchers **include a template** to guide how the gene is edited, the **DNA repair system in most cells is far more likely to make small, random insertions or deletions rather than adding a specific DNA sequence to the genome.**
- That makes it difficult — and in some cases, nearly impossible — for researchers to use CRISPR-Cas9 to overwrite one piece of DNA with a sequence of their choosing.

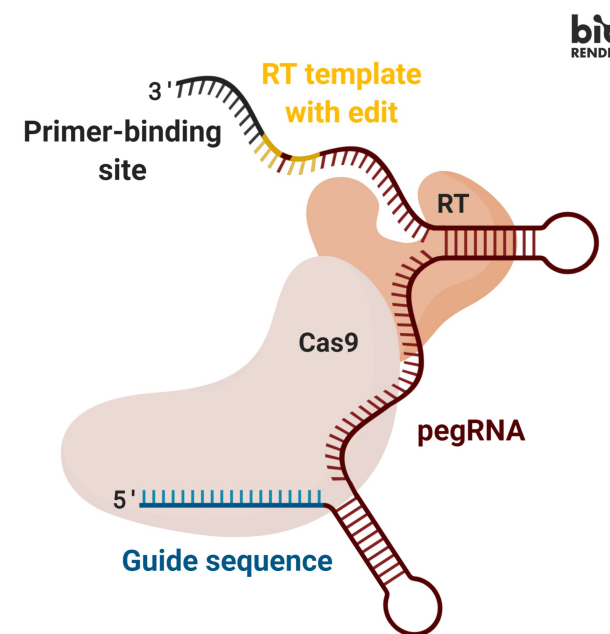


Source: Lily Padula / The New York Times.

# Where CRISPR Prevails

## Prime editing drawbacks

- “Prime editing may not be able to make the very large DNA insertions or deletions that CRISPR–Cas9 is capable of.
- Prime is unlikely to completely replace the well-established editing tool,” says molecular biologist Erik Sontheimer at the University of Massachusetts Medical School in Worcester.
- This is due to edits being encoded in a strand of RNA. The longer that strand gets, the more likely it is to be damaged by enzymes in a cell.



Source: Wikipedia: Ldinatto.

# CRISPR Achievements and Developments



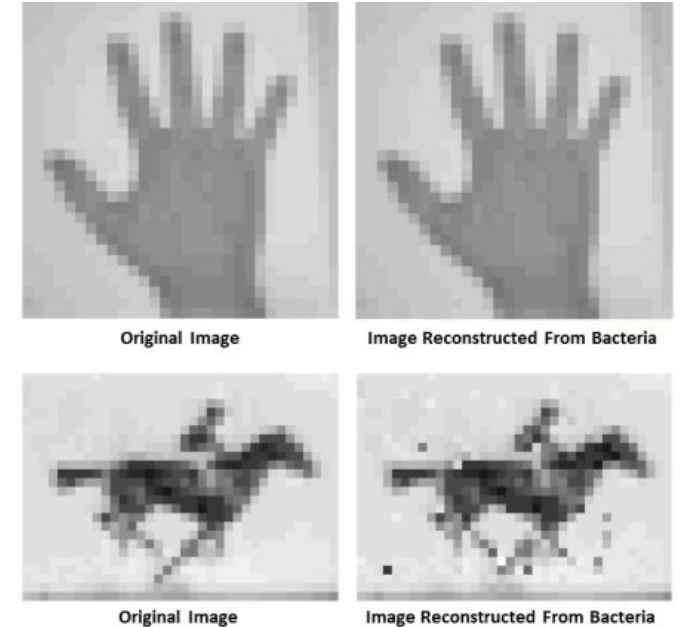


## Present: What Has Been Done?

### CRISPR progress

- Aside from being a noble prize-winning tool, CRISPR has many achievements.
- Scientists inserted a short-animated image in the E. coli DNA, which could eventually be used as a tiny hard drive.
- Gene editing techniques have shown potential to fight superbugs.
- Crispr has shown potential in extracting HIV from a living organism
- In 2018, the first gene edited babies announced by a Chinese Scientist who was convicted for this in 2019.

Images stored in bacterial DNA



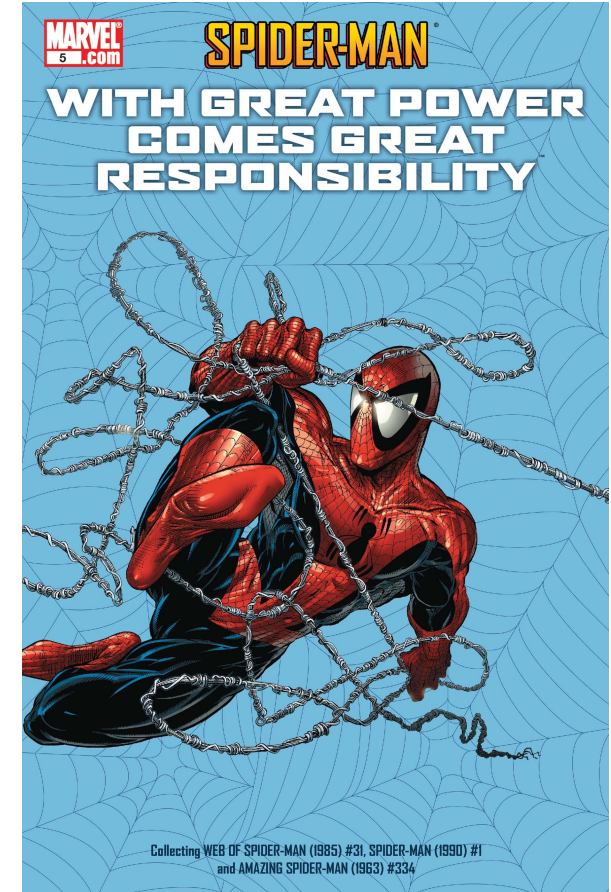
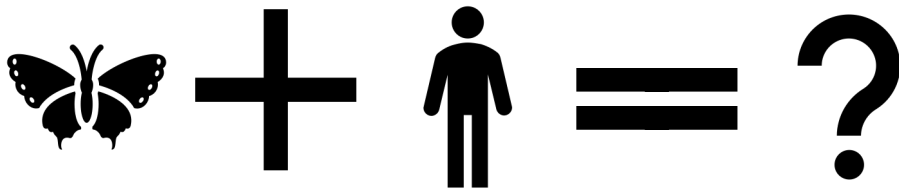
Source: Seth Shipman, Harvard Medical School, Boston.



## Future: What is Possible?

### Genetic mutations that change traits

- Steven Pete of Washington state carries a genetic mutation in the SCN9A gene, which causes insensitivity to pain. Unfortunately, Steven has broken more bones than he can count, and he once chewed off part of his tongue without realizing it.
- Scientists have experimented with ACTN3 gene mutations in mice and have found increased endurance.



Source: Marvel.

# Ethics, Issues, & Limitations



# Ethics & Issues

## Genetic manipulations can be passed on.

- While we think we know what we're doing it's possible that we can miss something, or our technology might not catch the changes that were made.

## Germline cell and embryo genome editing bring up several ethical challenges.

- One of which is whether it would be permissible to use this technology to enhance normal human traits (such as height or intelligence).

## Germline cell and embryo editing is banned.

- Based on ethical and safety concerns, germline cell and embryo genome editing are currently illegal in the United States and many other countries.

## How to guide the field.

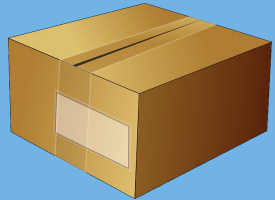
- Adopting a moratorium on heritable genome editing has become an idea to guide science.



Source: CompTIA.



# Limitations



## Delivery

- It's difficult to deliver the CRISPR/Cas material to mature cells in large numbers, which remains a problem for many clinical applications.

## Efficiency

- CRISPR is not 100% efficient, so even the cells that take in CRISPR/Cas may not experience genetic edits.



## Accuracy

- It's not 100% accurate, and “off-target” edits, while rare, may have severe consequences, particularly in clinical applications.



# Notable Players



## General Market Information

**In 2021 the Gene Editing market surpassed USD 5.4 billion.**

**Gene Editing Market to hit USD 19.9 billion by 2030, says Global Market Insights Inc.**

**The 2022-2030 compounded annual growth rate is 15.5%.**



# Caribou Biosciences, Inc.(NasdaqGS: CRBU)

- **Mission**

- Develop innovative, transformative therapies for patients with devastating diseases through novel genome editing.

- **chRDNA Platform**

- chRDNA is a proprietary CRISPR platform with significant advantages over 1st gen CRISPR-Cas9.

## chRDNA: a proprietary CRISPR platform with significant advantages over 1st gen CRISPR-Cas9

Significantly improved genome-editing specificity

Substantially fewer off-target events compared to first generation CRISPR-Cas9

High efficiency gene knockouts and insertions

Enables robust multiplex editing with high genomic integrity

Versatility across a broad range of cell types

Sophisticated genome editing across many cell types including immune cells and stem cells

Simple chemical synthesis

chRDNA guides are manufactured via chemical synthesis using readily available technologies

# CRISPR Therapeutics AG (NasdaqGM: CRSP)

- **About**

- CRISPR Therapeutics is a gene editing company focused on translating revolutionary CRISPR/Cas9 technology into transformative therapies.

- **Focus**

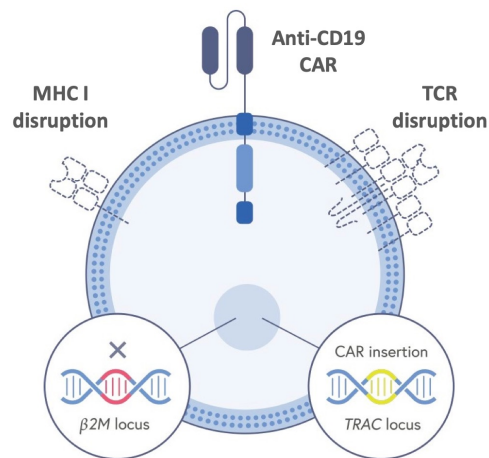
- hemoglobinopathies, immuno-oncology, regenerative medicine, and in vivo approaches

- **CRISPR enables regenerative medicine**

- CRISPR gene editing and pluripotent stem cell technology enable a new class of cell replacement therapies.

**Multiplex CRISPR gene editing in one step designed to:**

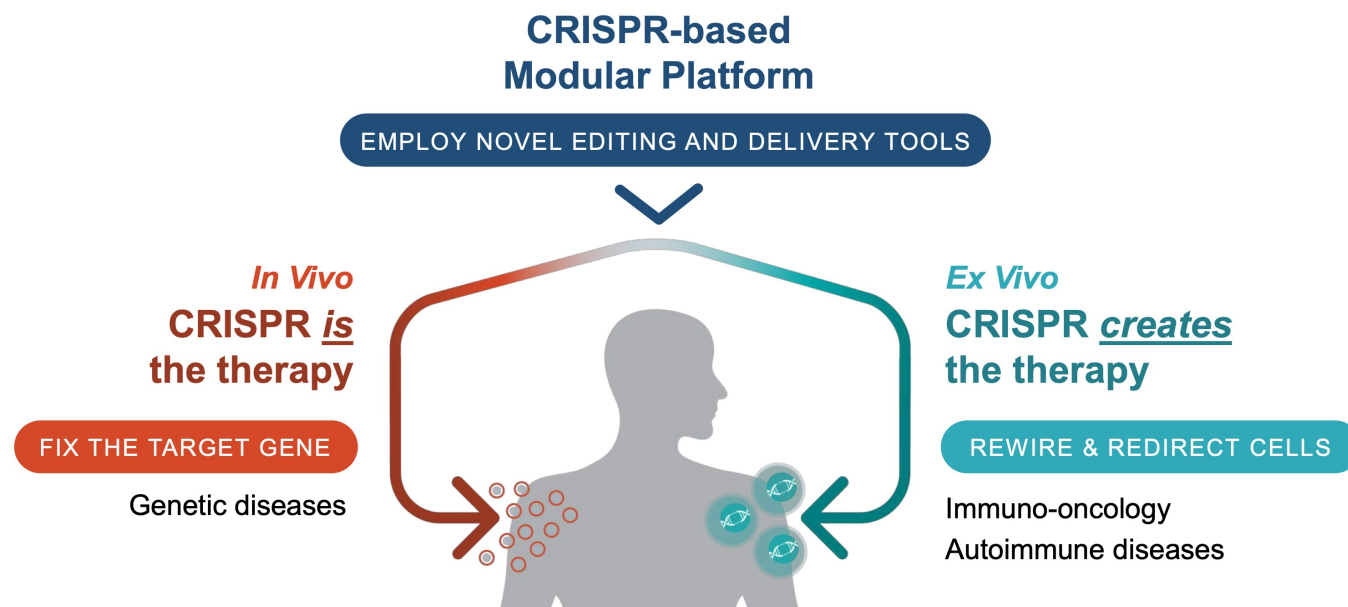
- **Improve persistence in the allo setting** via  $\beta 2M$  knock-out to eliminate MHC I expression
- **Avoid need** for more toxic lymphodepletion regimens



- **Prevent GvHD** via TCR disruption
- **Improve consistency and safety by precise insertion** of CAR construct into *TRAC* locus without using lentivirus or retrovirus

# Intellia Therapeutics, Inc. (NasdaqGM: NTLA)

- **In vivo Crispr therapy**
  - Potentially curative in one dose, capable of delivering to multiple tissue types for various therapeutic applications, permanent gain of function with targeted gene insertion, and systemic non-viral delivery of CRISPR/Cas9 provides transient expression and potential safety advantages
- **Modular Delivery Platform Enables Rapid and Reproducible Path to Clinical Development**
- **Proprietary Engineering Platform to Power Next-Generation Engineered Cell Therapies**





# Editas Medicine, Inc. (NasdaqGM: EDIT)

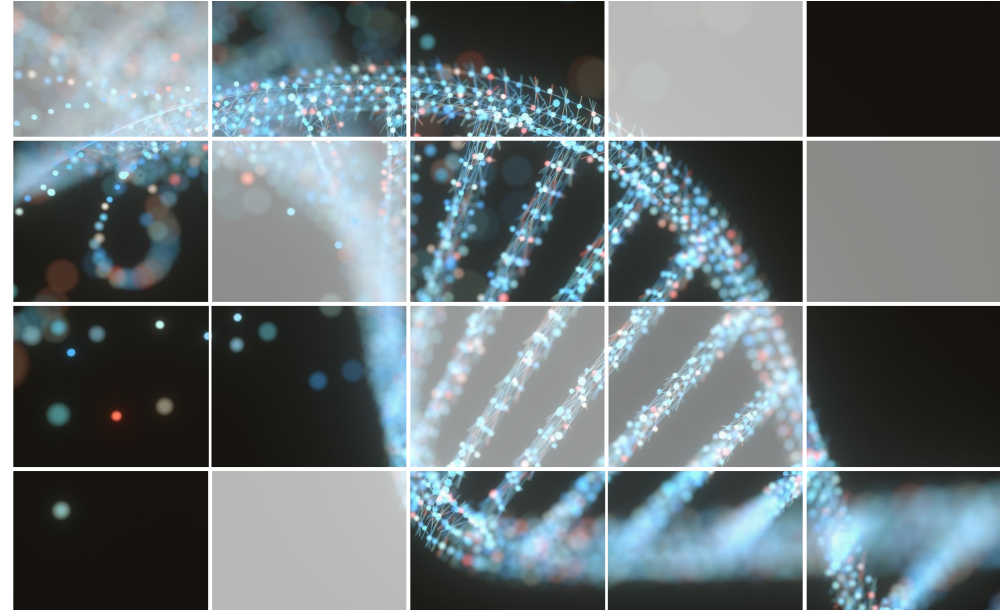
- **Intellectual Property Portfolio in CRISPR Gene Editing**

- Over 220 issued patents, over 800 applications pending
- Multiple species and CRISPR forms to address widest range of diseases
- Exclusive foundational IP for CRISPR/ Cas9 and Cas12a (Cpf1) editing in human therapeutic
- Global coverage including US, Europe, Japan, Australia, Canada, China





# Thanks!



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