

# Mutations & Gene Editing (HL)

## GENE KNOCKOUT

Rendering a gene unusable to see the effect it has on an organism, and hence its function.

## Human Genome Project

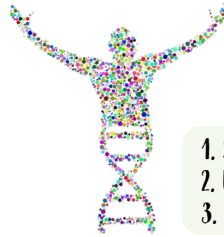
A project that aimed to find the base sequence (order) of our genome.

Gene knockout is used to help understand the purpose of each gene.

### ANALOGY



By removing the wheels, we can see what their function was by analyzing the leftover "frame".



### Test on humans

1. Expensive
2. Unethical
3. Dangerous

Knockout organisms

1. Share many of the same genes
2. Easy to maintain and control
3. Life cycle is much shorter

### Test on Mice

OR model organism\*



\* One that is used in place of another organism for ethical and practical purposes.

## Purposes of gene knockout:

1

### Studying gene function

By knocking out genes, we can figure out what their purpose is, helping us understand how we function

2

### Modeling human disease

By replicating disease in knockout mice, it helps researchers understand disease mechanism and test potential treatments.

3


### Developmental biology


Knocking out genes involved in embryonic development can help scientists understand birth defect and developmental disorders.

A technique called **CRISPR-Cas9** was developed which allows for genes to be edited; whereby mutated genes can be removed and/or replaced with an unmutated/new gene:

## CRISPR-Cas9 Gene Editing - 2012



Guide RNA (gRNA) 

Cas9 

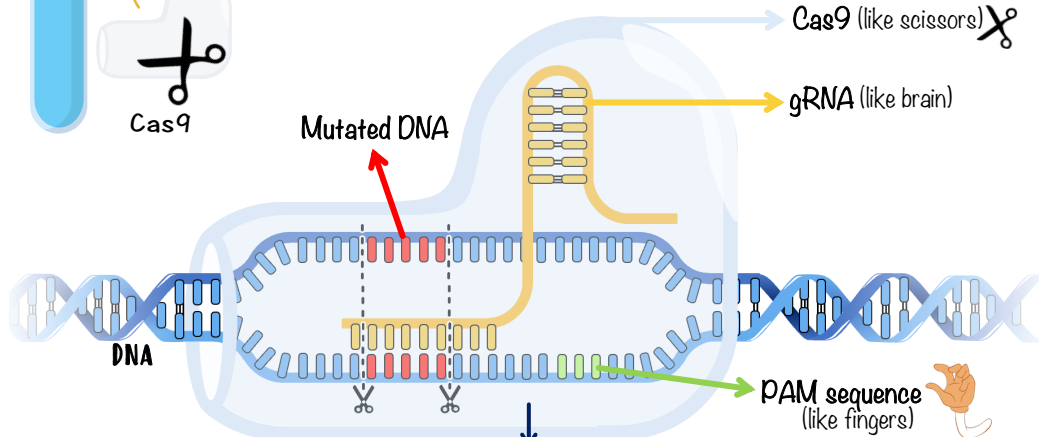
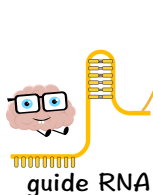
PAM sequence 

- 1 Mutated DNA is added to a guide DNA which was created to match the mutated DNA.

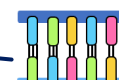
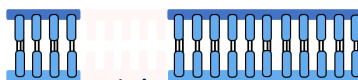
- 2 Cas9 (scissors) are also added to the mix.

- 3 The GUIDE RNA (gRNA) and the CAS9 PROTEIN attach to each other and this mix is injected into cells.

- 4 The guide RNA identifies the mutated DNA segment, and Cas9 uses a PAM SEQUENCE (protospacer adjacent motif) found near the mutated DNA to anchor to the DNA.



- 5 Cas9 cleaves the DNA double strand, removing the mutated segment of DNA.

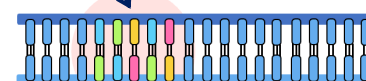


Unmutated gene/  
New gene

- 6 Mutated gene is removed or attempts at repair lead to errors leading to the inactivation or silencing of the gene.



- 7 Mutated gene can otherwise be replaced with a functional (not mutated) version of the gene OR a new gene.



# Mutations & Gene Editing (HL)

## ADVANTAGES of Gene Editing

### 1. DELETE HARMFUL GENE

Useful in single gene disorders - Huntington's or sickle cell disease

Even delete a full chromosome - Down syndrome



### 2. OPTIMIZE / AMPLIFY FUNCTION



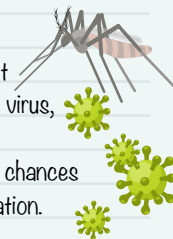
Agriculture - more food and more resistance, make food more nourishing or livestock more productive or resistant to disease.

Example - One CRISPR-Cas9 project, produced tomato plants that yielded twice as much as non-modified plants.

### 3. DISEASE TRANSMISSION

Human health - modifying mosquitoes so that they can no longer transmit the dengue fever virus, malaria parasite or west Nile virus.

Gene drive - a mechanism that increases the chances of a gene being passed on to the next generation.



## DISADVANTAGES of Gene Editing

### 1. GERM CELL EFFECTS ( FUTURE EFFECT)

If CRISPR-Cas9 is used in somatic cells, only the individual's genome is modified.

If it is germ cells, the individual is modified and can pass on the modification to future generations.



### 2. OFF TARGET EFFECTS

Another concern with gene modification is off-target effects. This is when the editing technology accidentally changes a part of the genome that was not intended to be modified.



### 3. TIME



Even when possible, in the laboratory, it doesn't mean it can be used yet: still needs testing.

Rules of safety and testing need to be followed.

## CONSERVED AND HIGHLY CONSERVED SEQUENCES

**Conserved sequences:** genetic sequences found in DNA or RNA that show **MINIMAL** mutations over time (in a species or population). Genomic\* & bioinformatics\*.

**Highly conserved:** those that show **NO** or almost no changes.

\* **Genomics** - science of gene sequencing \* **Bioinformatics** - Using computer programs to analyze the enormous data sets.

## What is the mechanism?

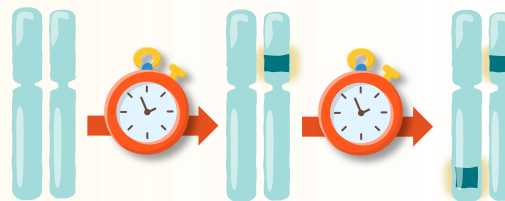
### 1. SLOWER MUTATION RATES

Key sequences (high risk) of the genome has a **MORE ACTIVE** DNA repair system. Repair system is less active in low risk zones (non-coding regions, silences genes).

Zones with low mutation rate  $\neq$  Less mutations

They are just corrected better than other zones [highly transcribed genes show lower mutation rates than less expressed ones.]

**Mutation rate** - refers to how many changes there are in a genetic sequence over time. It can be expressed as the number of base pairs changing in a single gene at each generation (or each cell division).



### 2. FUNCTIONAL REQUIREMENTS

Sequences which are critically **REQUIRED** for the proper **FUNCTIONING** of the cell (see right) are more highly conserved.

**NATURAL SELECTION** conserves such sequences by necessity (prevents mutated sequences from being passed on to the next generation as offspring wouldn't survive with the mutation).

#### PURIFYING SELECTION ( OR NEGATIVE SELECTION)

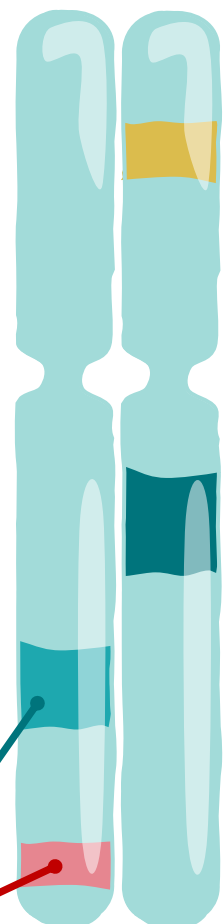
Phenomenon of eliminating harmful variation of genes.



Proteins used in DNA replication, transcription and translation: Helicase, non-coding RNA such as tRNA, and ribosomes.



Proteins used in cellular respiration such as cytochrome c and ferredoxin etc.



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