

# Bi4025en Molecular Biology

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#### Lecture 12

Basic principals of genetic engineering



## **Genetic engineering – definition**

- Definition by International Union of Pure and Applied Chemistry (IUPAC):
- Process of inserting new genetic information into existing cells in order to modify a specific organism for the purpose of changing its characteristics
- Also Known as Recombinant DNA technology, gene modification, and gene therapy.
- Genetic engineering is a term that was first introduced into our language in the 1970s to describe the emerging field of recombinant DNA technology.
- The recombinant DNA technology started with pretty simple things cloning very small pieces of DNA and growing them in bacteria.
- In a broad view the genetic engineering, is more defined as an entire field of recombinant DNA technology, genomics, and genetics.



## **Gene – Genetic engineering**

- Genetic engineering is targeted construction of cells/organisms carrying such gene and protein variants and a combination of genes that do not exist in the nature.
- Methods to propagate new molecule of DNA in new variant of the protein:
  - Joining DNA fragments isolated from different species, genera, families, etc...
  - Creation of completely new DNA molecules.
  - Transfer of modified molecules into cells and organisms.
  - Direct mutational interventions in the genome of cells and organisms.



## **Gene – Genetic engineering**

- DNA manipulation in vitro, DNA amplification, gene cloning and their targeted modifications, transfer to cells / organisms (heterologous expression).
- Targeted changes in genetic information may also be carried out in vivo (genome editing, chromosome engineering).



## **Key discoveries**

- Restriction endonucleases and other enzymes for DNA manipulation:
  - DNA cutting in a precisely defined location.
  - Ligation of two foreign DNA (originating from different organisms).
  - DNA synthesis in a test tube ("artificial bacteria").
- DNA sequencing:
  - Determination of the molecular structure of the gene and regulators sequences.
- Gene cloning:
  - Use of the universal genetic code and the method of expression of genetic information.
  - Introduction of the gene into the unrelated organisms (overcoming interspecies barriers).
  - Gene amplification to the unlimited quantities.
  - Targeted introduction of mutations into the gene (function improvement).
  - Study of the manifestation of altered genes (mutation x function).



## **Key discoveries**

- 1965 Discovery of plasmids.
- 1970 isolation of the first restriction enzyme.
- 1972 Preparation of the first recombinant DNA molecules in vitro.
- 1973 the beginning of gene cloning.
- 1975 The Asilomar Conference on recombinant DNA.
- 1977 First recombinant DNA molecules carrying mammalian genes.
- 1977 DNA sequencing.
- 1978 preparation of human insulin in bacteria (since 1982 produced commercially).



## **Key discoveries**

- 1980 First attempts of gene therapy.
- 1983 Discovery and introduction of PCR, an alternative to gene cloning in vectors.
- 1997 animal cloning, Polly the sheep (injection of genomic DNA into the nucleus the human blood clotting factor, exogen present, but not expressed).
- after 2000 genome editing recombinant DNA nucleases meganucleases recognizing long target sequences, CRISPR/Cas.



## Use of genetic engineering

- Basic Research:
- Study of the <u>structure</u> and <u>function</u> of genes and genomes.
- Elucidation of basic processes of gene expression recombinant DNA, reporter genes, etc.
- Characterization of regulatory mechanisms of gene expression.



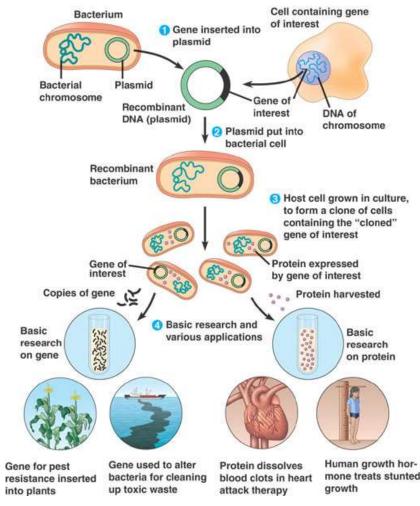
## Use of genetic engineering

- Applied Research:
- Production of medical, agriculture and industry relevant substances:
  - Introduction of foreign genes into unrelated organisms and obtaining products in large quantities - overcoming reproductive barriers.
- Preparation of substances with new properties:
  - Modification of existing genes or creating new ones enzymes, antibodies, vaccines, etc.
  - Improving the properties of therapeutic substances.
- Alteration and improvement of the characteristics of organisms:
  - Preparation of microorganisms for biotechnology (e.g. Enhanced production).
  - o Increased yields of cultural plants and livestock performance (resistance to diseases, pests or external influences, production of foreign substances in the bodies of plants and animals).
- Gene therapy treatment of genetic diseases.



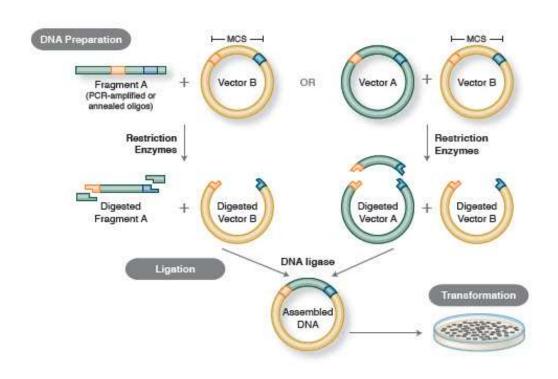
## Particular steps of genetic engineering

- 1) Insertion of the gene int the plasmid – Gene cloning.
- 2) Transfer of vector in the given organism – bacteria, plant, yeast, eukaryotic cells, etc. By transformation/transfection, electroporation, transduction, infection, microinjection, shooting gene.
- 3) Expanding of bacteria.
- 4) Basic or Applied Research.





## Gene cloning

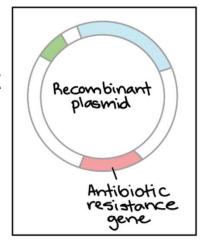


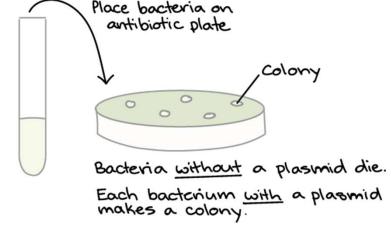
- Restriction sites at the DNA and plasmids are digested by the corresponding restriction enzymes (RE).
- The cleaved DNA is then ligated to a plasmid vector possessing compatible ends.
- DNA fragments can also be moved from one vector into another by digesting with REs and ligating with compatible ends of the target vector.
- Assembled construct is then transformed into Escherichia coli (E. coli).



#### Plasmid selection

- A plasmid typically contains an antibiotic resistance gene, which allows bacteria to survive in the presence of a specific antibiotic.
- Thus, bacteria that took up the plasmid can be selected on nutrient plates containing the antibiotic.
- Bacteria without a plasmid will die, while bacteria carrying a plasmid can live and reproduce.
- Each surviving bacterium will give rise to a small, dot-like group, or colony, of identical bacteria that all carry the same plasmid.

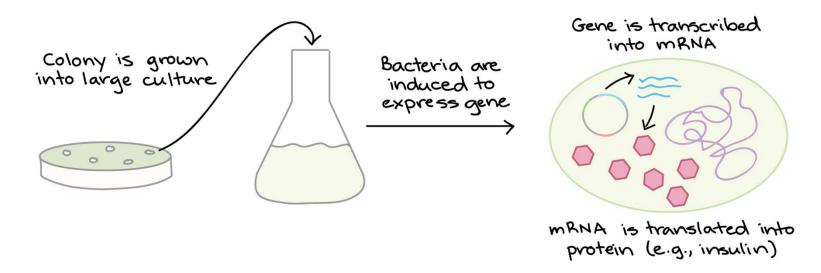






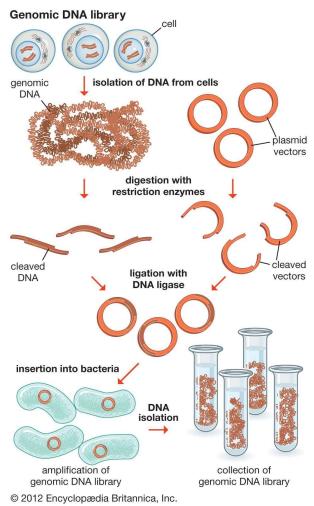
## **Production of cloned gene**

- Once the bacterial colony with the right plasmid is verified.
- After specific induction the large culture of plasmid is then grown.
- The bacteria serve as miniature "factories" churning out large amounts of protein.





## **Cloning of genomic library**



- A genomic DNA library is a collection of DNA fragments that make up the full-length genome of an organism.
- A genomic library is created by isolating DNA from cells and then amplifying it using DNA cloning technology.



## Mutagenesis in vitro

"Site - Directed Mutagenesis".

#### Aims:

- Analysis of the relationship between structure and function of nucleic acids.
- Clarification of gene function and regulatory regions.
- Targeted changes in amino acids in proteins.
- Controlled evolution (bioinformatics tools, domain modeling, gene preparation, cloning) and function test) - protein preparation with new properties (protein engineering).
- Preparation of transgenic organisms

#### Procedure:

- DNA isolation change in DNA in vitro transfer to the cell evaluation of the effect.
- Types of mutations: substitution, deletion, insertions.



## Mutagenesis in vitro

- Random
- Handling of restriction points.
- Chemical mutagenesis
- Incorporation of erroneous bases.
- Gene search, or functional areas on DNA.

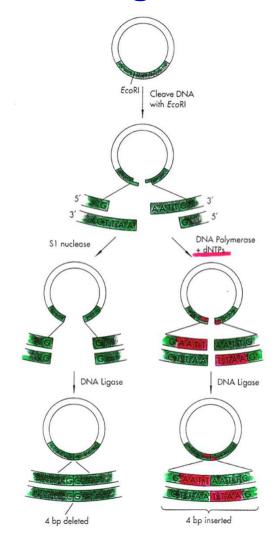
- Targeted
- Oligonucleotides mutagenesis (placement of mutations in a specific gene site).
- Cassete mutagenesis (gene synthesis).
- Substitution of bases or codons targeted changes in the structure of proteins.



## Restriction enzyme mediated mutagenesis in vitro

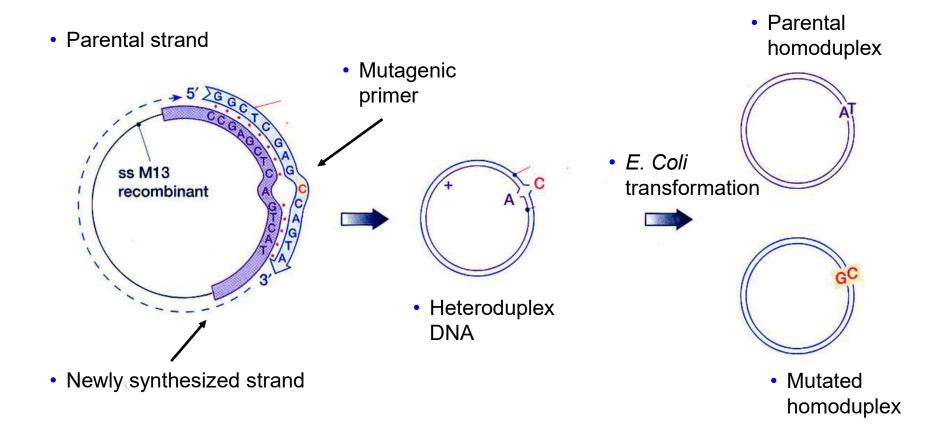
- Overhanging ends are:
- removed by Si nuclease.
- synthesized by DNA-polymerase also just limited synthesis by chosen nucleotides.

 Generation of deletions or insertions within the genome.



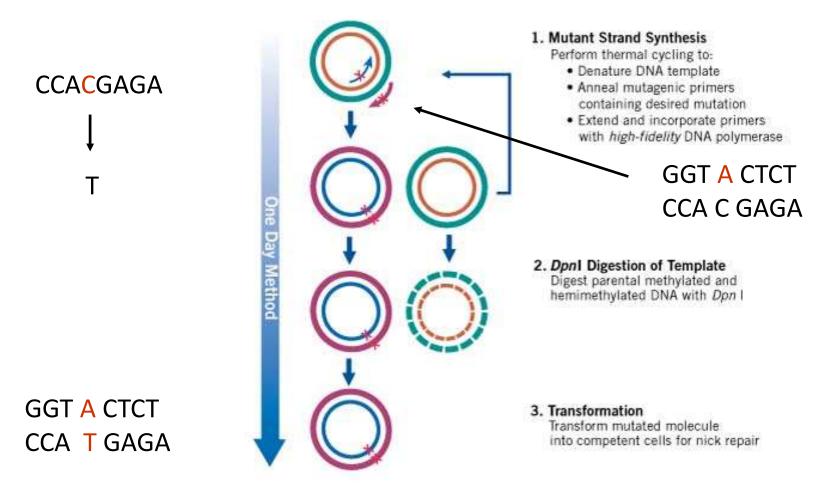


## Oligonucleotide mediated mutagenesis





## Oligonucleotide mediated mutagenesis

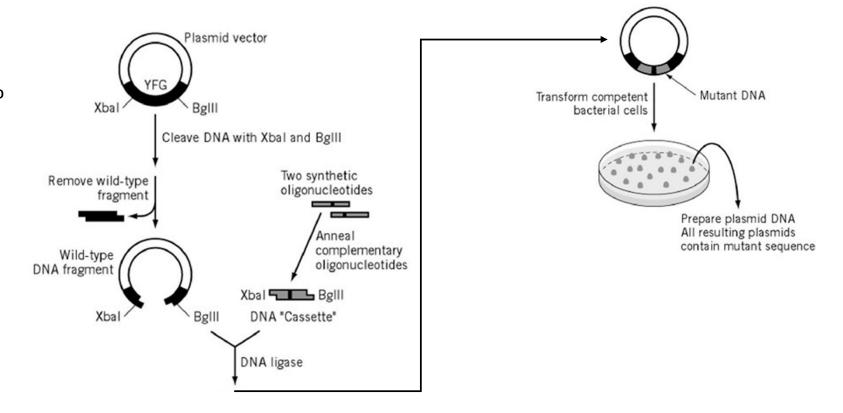


Primer **GGT A CTCT** 



## **Cassette mutagenesis**

- Generally nearly 100% of the mutant clones.
- The limitation is presence of unique, conveniently spaced restriction endonuclease recognition sites.





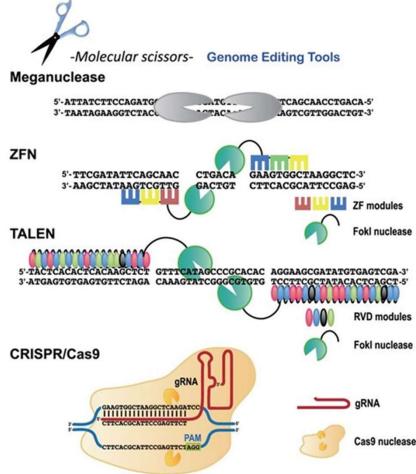
## In vivo mutagenesis - gene editing

- Genome editing with engineered nucleases.
- Since 2000.
- Targeted changes to any gene in any organism in vivo.
- Artificially prepared nuclease introduces insertions, deletions, or replacement of the existing sequence at selected location within the genome.
- Nucleases generates double-stranded breaks at designated points in the genome, thereby stimulating natural repair mechanisms in cells based on:
  - Homology-directed recombination HDR HR
  - Non-homologous end-joining NHEJ.



In vivo mutagenesis – Endonucleases

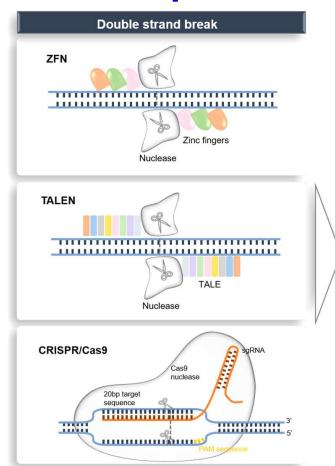
- Meganucleases
- Zinc-finger nucleases ZFN
- Transcription Activator-Like
  Effector Nuclease TALEN
- Clustered Regulatory Interspaced Short Palindromic repeats -CRISPR

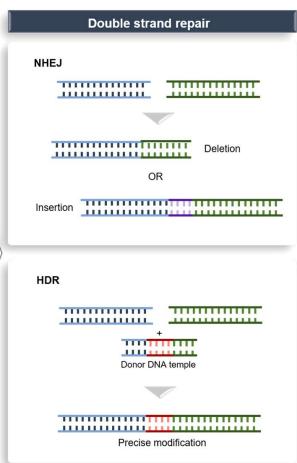




## In vivo mutagenesis – Repair Mechanism

- Double strand break can be repaired by:
- NHEJ insertions, deletions.
- HDR integration of new sequences.







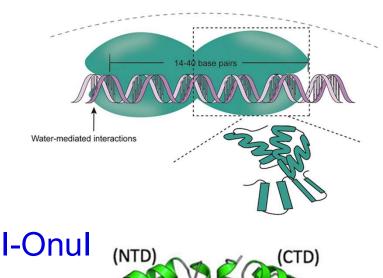
## Meganucleases

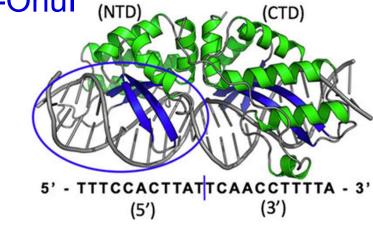
- There are present in archaebacteria, bacteria, yeast, fungi and plants.
- Meganucleases identify approximately 14-40 base pairs in the target sequences.
- 5 families of meganucleases are known. The most widespread of them "LAGLIDADG family".
- Meganuclease I Sce I recognizes the 18 nucleotide sequence "TAGGGATAACAGGGTAAT", cleaves 4 nucleotides and leaves an end with overlap.
- Complications the 18 nucleotide sequence recognized by the I-Scel meganuclease would hypothetically require a genome 20 times larger than the human genome for the sequence to occur randomly in the genome.
- The off-target activities induced by meganucleases are affected by the structure of meganucleases, and the delivery methods.



## Meganucleases

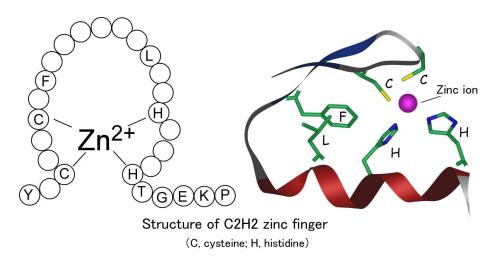
- Meganucelases form dimer at the target site.
- Each monomer can form αββαββα fold, with fourstranded antiparallel β-sheets to recognize and combine with target sequence.
- I-Onul meganuclease 290 residues and is bound to a 22 base pair DNA target site.
- The N-and C-terminal domains of the endonuclease recognize and interact with the 5' and 3' half-sites of the DNA target site.
- The interface between the target 5' half-site and the protein N-terminal domain is indicated by the oval.







## Zinc-finger nucleases – DNA binding domain

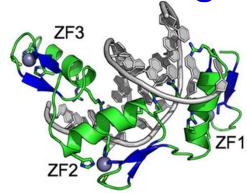


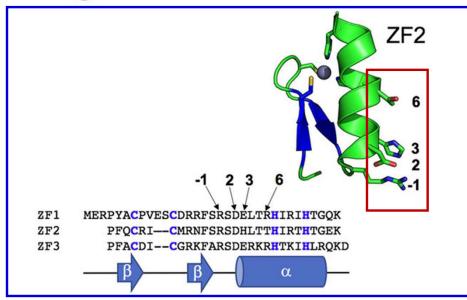
- The C2H2 zinc finger protein (ZFP) is one of the most common DNA binding motifs in multicellular organisms. The zinc binding coordinated by the amino acids Cysteine or Histidine.
- It is capable of binding DNA/RNA/Proteins.
- Individual fingers contain about 30 amino acids and these units typically occur as tandem repeats of two or more fingers.
- Fusing the nonspecific DNA cleavage domain of the Type II restriction enzyme, FokI, to a ZFP allows the resulting zinc finger nuclease (ZFN) to cleave DNA at a sequence determined by the ZFP.



## Zinc-finger nucleases – DNA binding domain

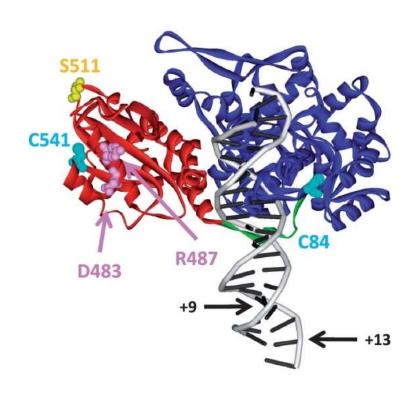
- Structure of the Zif268-DNA complex.
- Fingers are spaced at 3-bp intervals.
- The structure and primary DNA contacting residues of zinc finger 2 (ZF2) are indicated to the right blue rectangle.
- A sequence alignment of the 3 fingers of Zif268. The zinc binding Cys2-His2 motif is indicated with blue bold font; the canonical DNA-contacting residues are indicated by arrows in the red rectangle.







## Zinc-finger nucleases – Fokl endonuclease

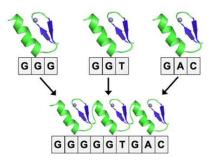


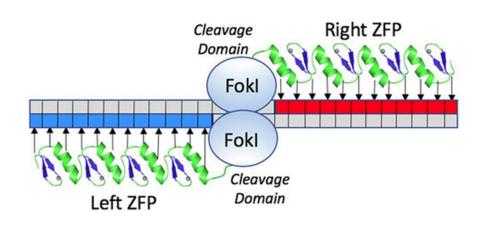
- Fokl endonuclease occurs naturally in Flavobacterium okeanokoites.
- Consists of an N-terminal DNA-binding domain distinguishing a 5' GGATG 3' binding site and a Cterminal domain capable of non-specific DNA cleavage.
- After binding to double-helix DNA, a domain capable of cutting is activated, which subsequently cuts:
  - o 9 nucleotides per 5<sup>o</sup> → 3<sup>o</sup> string
  - $\circ$  13 nucleotides per 3'  $\rightarrow$  5' chain.



## Zinc-finger nucleases

 To generate a ZFP with specificity for the sequence GGGGGTGAC, three fingers are identified that each bind a component triplet. These fingers are then linked.



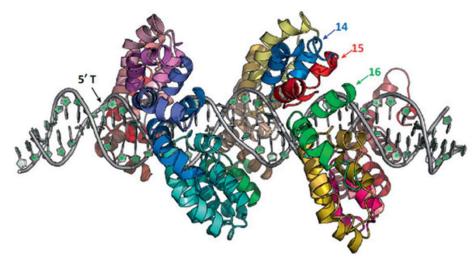


- Sketch of a pair of zinc finger nuclease (ZFN) subunits bound to two halves of a DNA target.
- Each ZFN contains the cleavage domain of Fokl linked to an array of three to six zinc fingers to specifically recognize sequences that flank the cleavage site.
- The Fokl nuclease domains transiently dimerize across those central bases and cleave each DNA strand to generate a double strand break with 5 overhangs averaging 4 bases in length.

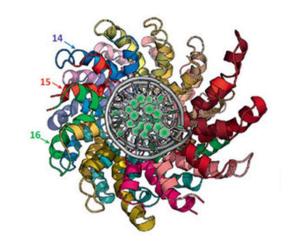


## **Transcription Activator-Like Effector Nuclease**

- TAL: Transcription Activator-Like.
- A group of proteins capable of specific binding to DNA.
- It is used by plant bacteria of the genus Xantomonas and affects the DNA of the guest/recipient.
- RVD: Repeat-variable di-residues, these are amino acids found in the TAL at position 12 & 13 that determine the specificity to the target nucleotide.

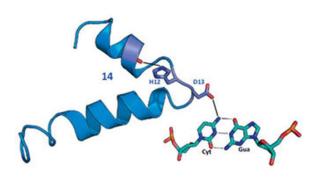


PthXo1 binding region

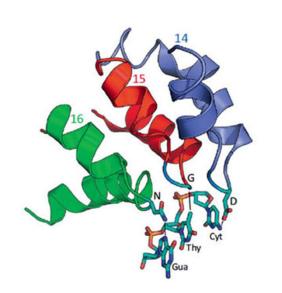




## **Transcription Activator-Like Effector Nuclease**



 The histidine at position 12 in the repeat number 14 forms a hydrogen bond to the backbone carbonyl oxygen of residue 8 in the first a-helix, while the aspartate at position 13 forms a hydrogen bond to the extracyclic amino nitrogen of the cytosine base.

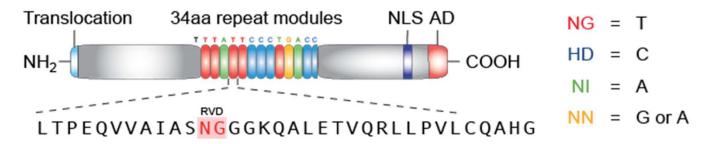


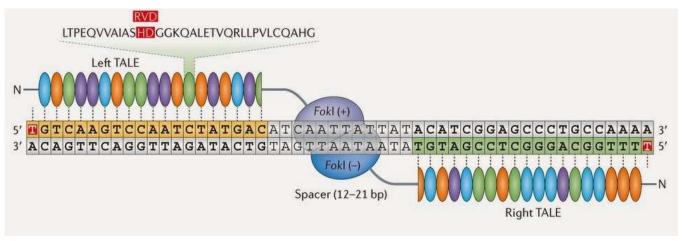
 Repeats 14, 15 and 16 interacting with the DNA, illustrating that consecutive RVDs (HD, NG and NN, respectively in these repeats) contact consecutive bases (in this case cytosine, thymine, and guanine) on the same DNA strand.



## **Transcription Activator-Like Effector Nuclease**

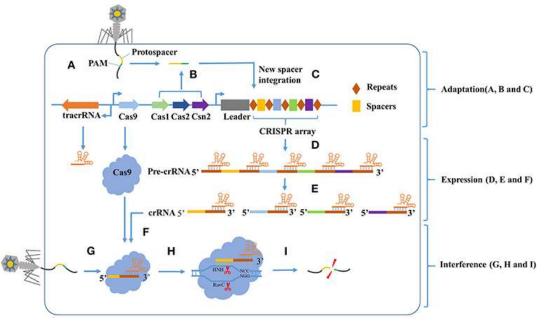
• RVD: Repeat-variable di-residues are amino acids in position 12 and 13 determine specificity to the target nucleotide - T, C, A and G.







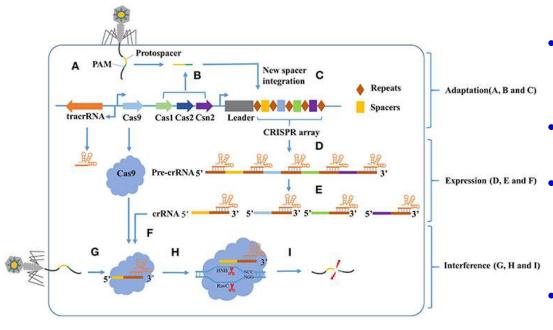
## Clustered regularly interspaced short palindromic repeats



- (A) The host is invaded by phage DNA during infection.
- (B) The invading DNA is processed by various
   Cas genes into small DNA fragments
   (protospacer). The selection of protospacer
   depends in part on the specific recognition of
   protospacer adjacent motif (PAM) present within
   the viral genome.
- (C) The small DNA fragments is then incorporated into the CRISPR locus of the bacterial genome as a new spacer, flanked by a repeat sequence.
- (D) the CRISPR locus is transcribed into a long precursor CRISPR RNA (pre-crRNA).



## Clustered regularly interspaced short palindromic repeats



- (E) The dual RNA hybrid, tracrRNA with precrRNA, is recognized and cut by RNase III.
- (F) The mature crRNA will combine to Cas9 protein and guide the DNA cleavage.
- (G) Phage invades again bacteria.
- (H) The Cas9:crRNA complex identifies a sequence that is complementary to the spacer and adjacent to the PAM.
- (I) Finally, the invading DNA is cleaved by Cas9 protein to prevent infection.

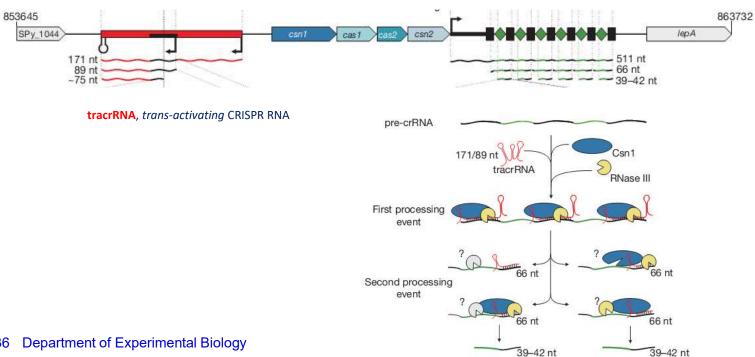


#### **Discovery of tracrRNA**

#### CRISPR RNA maturation by trans-encoded small RNA and host factor RNase III

Elitza Deltcheva<sup>1,2</sup>, Krzysztof Chylinski<sup>1,2\*</sup>, Cynthia M. Sharma<sup>3\*</sup>, Karine Gonzales<sup>2</sup>, Yanjie Chao<sup>3,4</sup>, Zaid A. Pirzada<sup>2</sup>, Maria R. Eckert<sup>2</sup>, Jörg Vogel<sup>3,4</sup> & Emmanuelle Charpentier<sup>1,2</sup>

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**Emmanuelle Charpentier: CRISPR Therapeutics** 

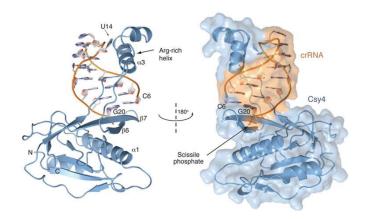


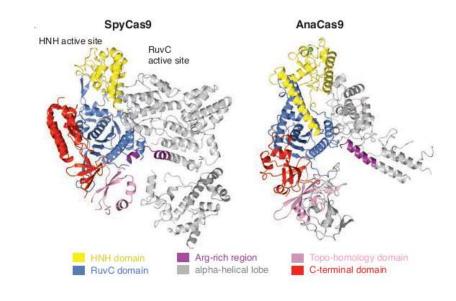


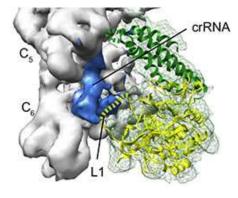
#### **Structure of Cas9**

Jennifer Doudna: Intellia Therapeutics & Caribou Biosciences











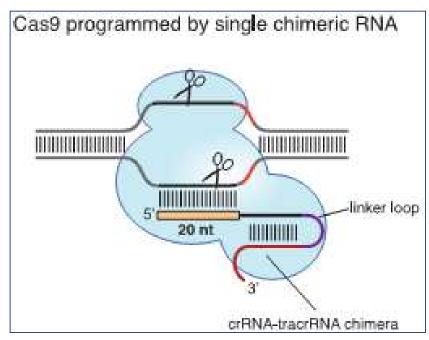
## Preparation of dual tracrRNA-crRNA into gRNA

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

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

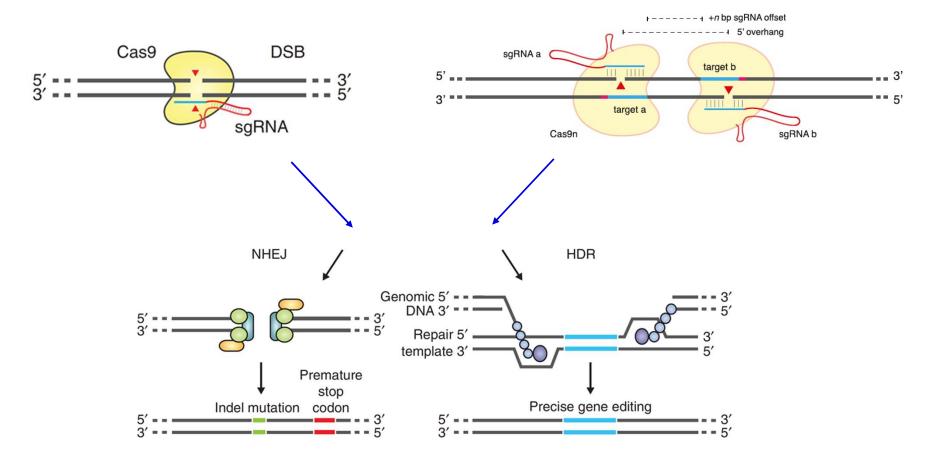
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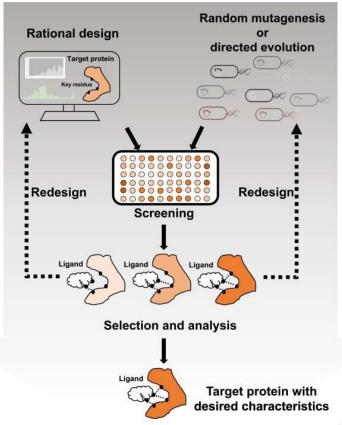
#### **CRISPR/Cas9 system**





#### **Protein engineering**

- Protein engineering is the conception and production of unnatural polypeptides, often through modification of amino acid sequences that are found in nature.
   Synthetic protein structures and functions can now be designed entirely on a computer or produced through directed evolution in the laboratory.
- Objective: to change the structure and function of proteins through recombinant DNA technology:
  - Changes in protein binding regions.
  - Thermostability.
  - Speed and substrate specificity of reactions.
  - Sensitivity to oxidation and toxic substances.

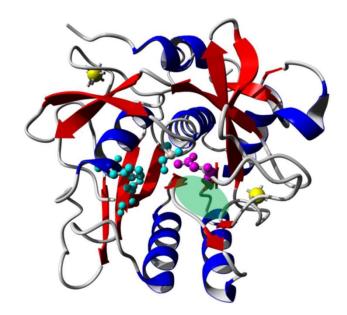






# Protein engineering - enzymes in washing powders

- Enzymes in washing powders are strongly genetically modified in order to require traits achieved by genetic modification:
  - Water softening.
  - Detergent for removing impurities (proteins, dyes, polysaccharides, etc.).
  - Efficiency at high and low temperature.
  - Whitening effect.
- Subtilisin is a protease produced by Bacillus subtilis into the extracellular space, decomposes substrates and their products use as source of nutrients.
  - Lysis a number of substrates.
  - Modified to withstand high temperatures.
  - Expansion for substrates spectrum.
  - Resistance to bleach.
  - All changes were done by successive/step by step amino acid substitutions.





# **Protein engineering – Industrial application of eznymes**

Sector	Enzymes	Applications	
Pharmaceuticals	Nitrile hydratase, transaminase, monoamine oxidase, lipase, penicillin acylase	Synthesis of intermediates for production of active pharmaceutical ingredients	
Food Processing	Trypsin, amylase, glucose isomerase, papain, pectinase	Conversion of starch to glucose, production of high fructose corn syrup, production of prebiotics, debittering of fruit juice	
Detergent	Protease, lipase, amylase, cellulase	Stain removal, removal of fats and oils, color retention,	
Biofuels	Lipase, cellulase, xylanase	Production of fatty acid methyl esters, decomposition of lignocellulotic material for bioethanol production	
Paper and Pulp	Lipase, cellulase, xylanase	Removal of lignin for improved bleaching, improvement in fiber properties	



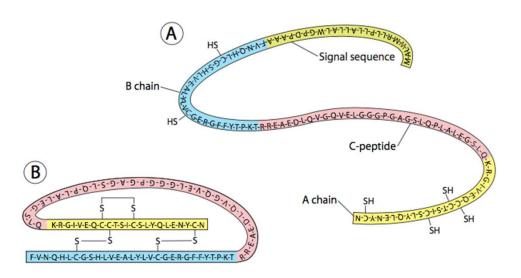
#### Insulin

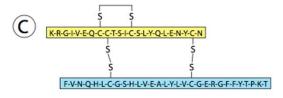
- Insulin is produced by B cells of the islets of Langerhans of the pancreas and released in the blood.
- Insulin lowers blood sugar levels (antagonizes glucagon) function).
- Used for the treatment of diabetes.
- In humans composed of two polypeptides A and B connected by two disulfide bonds.



#### **Processing of insulin in mammals**

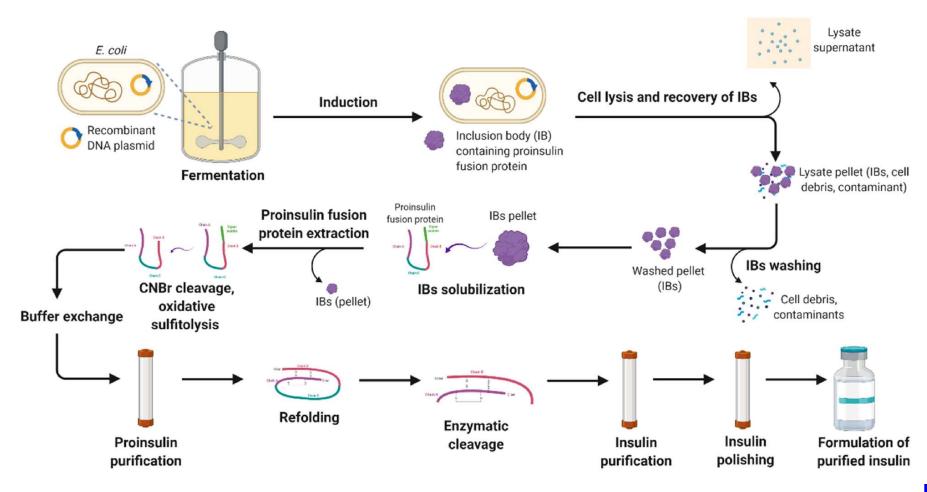
- In mammals preproinsulin (inactive as a hormone) is first translated from the insulin mRNA.
- Proteolytic processing is necessary to make biologically active insulin.
- (A) The linear protein preproinsulin contains a signal sequence, which is cleaved after the protein enters the ER, an A chain, a B chain, and a Cpeptide.
- (B) Inside the ER, the proinsulin (insulin precursor) folds and disulfide bonds form between cysteines.
- (C) Finally, two cleavages release the C peptide, which leaves the A and B chains attached by the disulfide bonds. This is now active insulin.





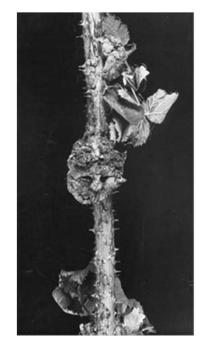


#### Industrial production of recombinant insulin





- Gene transfer from bacteria to plants occurs naturally.
- Agrobacterium tumefaciens is a soil pathogen, a gram-negative bacterium which infects many species of plants causing a disease known as "crown gall".
- It has two common species:
  - o A. umefaciens
  - o A. rhizogenes



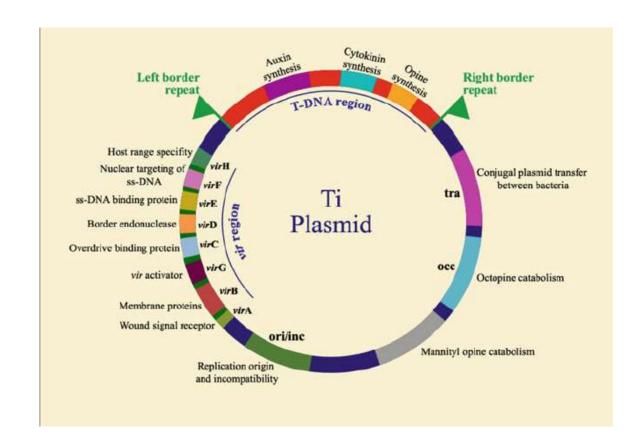




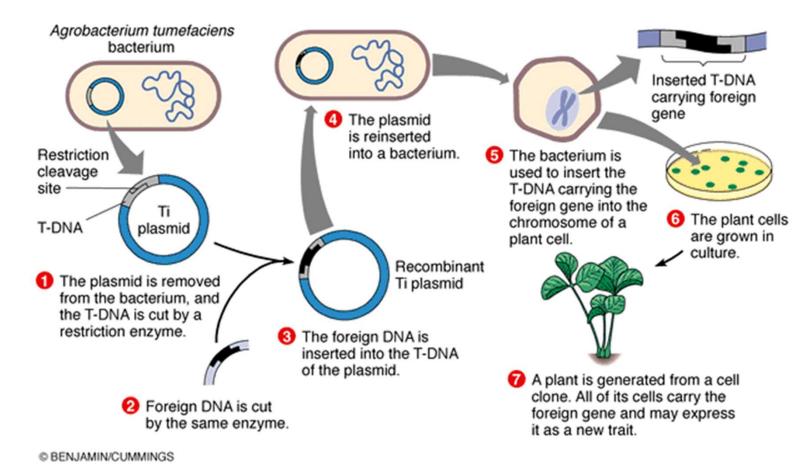
- The oncogenic activity of *Agrobacterium* is due to the presence of a large (200-kb) tumor inducing plasmid (pTi) and tumor formation results from its infection into the host plant (dicotyledonous plants).
- Upon infection of the wounded plant, the bacterium transfers a small segment of DNA, the so-called T-DNA from the Ti plasmid into the plant cell.
- The T-DNA is then translocated into the plant nucleus and stably integrated into the chromosome.
- In the Ti-plasmid itself, the T-DNA is flanked by 25-bp imperfect direct repeats known as border sequences.



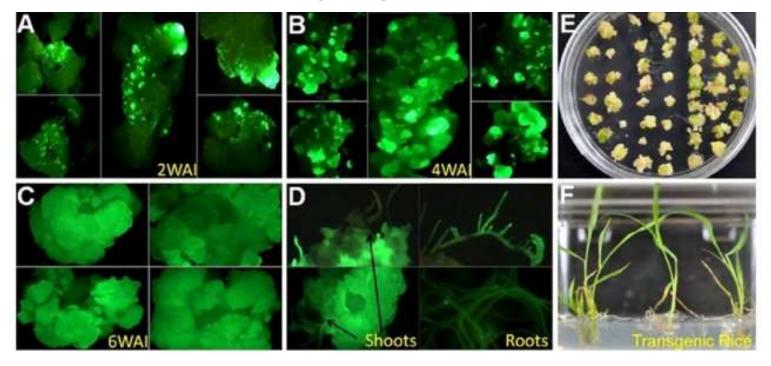
- The wild type T-DNA encodes a specific set of oncogenic (onc) genes that when it is expressed in the host cell, leads to the formation of a tumor.
- T-DNA carries genes for phytohormone (auxin and cytokinin) and opines.
- The overproduction of phytohormones at the site of infection is responsible for the proliferation of wound cells into a gall (tumor) that can harbor a population of bacteria.











- Schematic presentation of rice transformation.
- GFP expression in callus 2 weeks after infection (A), 4 weeks after infection (B), 6 weeks after infection (C), and in regenerating shoots and roots (D).



#### Use of genetic engineering in the plants

- Food and feed.
- Influencing agronomic properties:
  - Herbicide resistance.
  - Resistance to pathogens (insects, viruses, fungi, etc.).
  - Tolerance to stress (water stress drought, temperature –frost; osmotic stress – salinization of soils).
- Modification of post-harvest properties:
  - Extended shelf-life.
  - Slowing down maturation and inducing resistance to shelf storage diseases.
  - Refine nutritional value and taste.

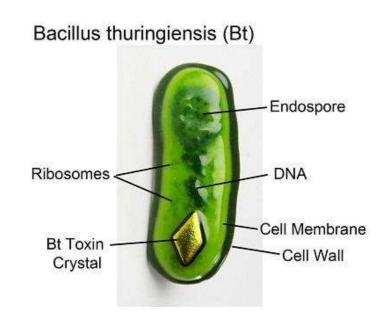


## Use of genetic engineering in the plants

- Production of secondary metabolites:
  - Pharmacological preparations.
- Technical crops:
  - Production of starch and oils for industrial use.
  - Biodegradable plastics.
- Phytoremediation:
  - OUse of green plants to remove polluting substances from the environment.



- Resistance to insect pests.
- The spores of bacterium *Bacillus thuringiensis* produce protein with toxic effects on insects Delta endotoxin =  $\delta$ -endotoxin.
- $\delta$ -endotoxin is highly poisonous to some groups of insects, but completely harmless to mammals and humans.  $\delta$ -endotoxin changes to a toxic form, the only in the intestine of insects.
- Previously bacteria *Bacillus thuringiensis* cultured in large volumes and as spraying applied to plants .
- Later the gene for δ-endotoxin transferred to plants directly transgenic potatoes, corn, rice, tobacco, tomato, broccoli, cotton.
- Indirect way cloning the gene for toxin into bacteria colonizing plants (leaves, roots)– e.g. Pseudomonas fluorescens.





#### **Transgenic BT-Maize**

- In addition, it contains 3 genes:
- Gene for plant resistance to insect pests.
  - δ-endotoxin from B. thuringiensis.
- Gene for resistance to the herbicide Basta.
- Basta is a short-lived herbicide, that is environmentally friendly.
- The gene for resistance comes from the bacterium Streptomyces.
- Gene for resistance to the antibiotic ampicillin.
  - Selection marker used for selection of transgenic plants (cells).
  - The gene is of bacterial origin.





- Herbicide resistance:
- Weeds negatively affect the yield and quality of the crop (competition for nutrients, light, pest source).
- Herbicides are chemicals that disrupt certain metabolic processes in the plant and damage it.
- Selective weed elimination is a problem, a number of herbicides are total - they also destroy cultural crops.
- Only after the introduction of herbicide insensitivity transgenes into the genome of cultural crops is achieved its selectivity.



- Resistance to the herbicide glyphosate (Roundup).
- Glyphosate non-selective, total herbicide.
- Effective on 76 of the 78 most important weeds.
- Inhibits enzymes providing the synthesis of essential amino acids.
- This enzyme is not present in animals the herbicide does not harm them.

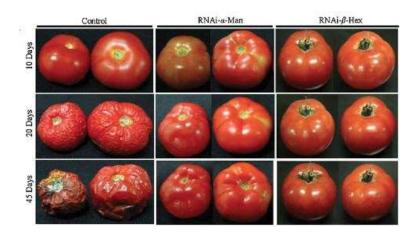
#### • Strategy:

- Introduction of a gene for the formation of the target enzyme into agricultural plants (a larger amount of enzyme will improve the resistance of plants).
- Introduction of a gene for the formation of altered more active enzyme.
- Introduction of a gene for the formation of an enzyme that inactivates the herbicide.



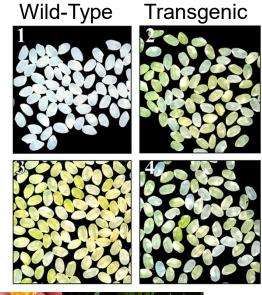


- Improvement of nutritional values of fruits and seeds or plant products used industrially.
- FlavrSavr tomato from company <u>Calgene</u>.
- The tomato was made more resistant to rotting, by adding an antisense RNA which interferes with the production of the enzyme polygalacturonase. This enzyme is normally responsible for cell walls softening during the fruit ripening.
- Unmodified tomatoes are <u>picked before fully ripened and</u> artificially ripened using ethylene gas which acts as a plant hormone - easier handling and extended shelf-life.
- FlavrSavr tomatoes could be allowed to ripen on the vine, without compromising their shelf-life.





- Improvement of nutritional values of fruits and seeds or plant products used industrially.
- Rice with provitamin A (carotene):
- Solves vitamin A deficiency in poor countries (consequence of night blindness, blindness).
- Three genes introduced into the rice genome, the expression of which leads to the formation of provitamin A.
- Further modification of rice aims to increase the iron content.



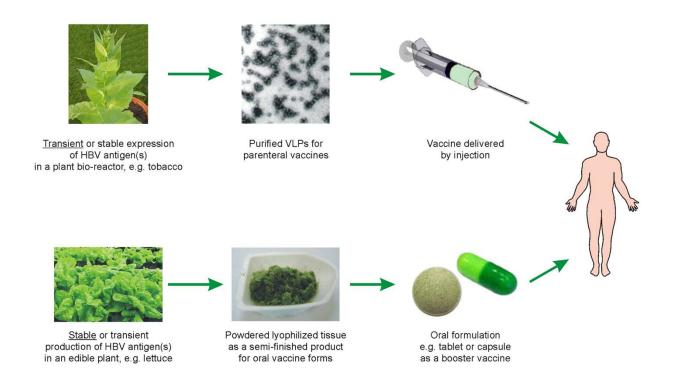




- Improvement of nutritional values of fruits and seeds or plant products used industrially.
- Brassica napus var. Napus, rapeseed seed oil containing an increased proportion of acid. Lauric (soaps and detergents).
- Rapeseed seed oil rich in myristate (cosmetics), or erucic acid (lubricants and nylon production).
- Arabidopsis and rapeseed formation of biodegradable polymers in chloroplasts usable as plastics (polyhydroxybutyrate, polyester-like polymers in cotton fibers).
- Decaffeinated coffee plant due to spontaneous mutations a rare variant found in which a key enzyme for caffeine formation is absent.



#### Transgenic plants – vaccine production



- Plants are suitable for passive immunization (eating transgenic plants containing fragments of antibodies) and active immunization (antibody induction).
- Immunization by eating raw vegetables containing antigen (vaccine), which induces the production of immunoglobulins mucosal immune system in the digestive tract.
  - Hepatitis B virus surface antigen.
  - Subunit B toxin cholera.

• Developmental status of edible vaccines in clinical trials.

Pathogen	Antigen	Host	Use	Clinical trial status
Enterotoxigenic				
	LT- B	Potato	Diarrhoea	Early phase 1
E. coli				
Enterotoxigenic				
	LT- B	Maize	Diarrhoea	Early phase 1
E. coli				
Norwalk Virus	СР	Potato	Diarrhoea	Early phase 1
Rabies Virus	GP/ NP	Spinach	Rabies	Early phase 1
HBV	HBsAg	Lettuce	Hepatitis B	Early phase 1
HBV	HBsAg	Potato	Hepatitis B	Phase 1
Vibrio cholerae	СТВ	Rice	Cholera	Phase 1
HBV	HBV	Saccharomyces cerevisiae	Chronic HBV	Phase 2



#### **Transgenic animals**

- Instrumental tool to:
- Study of the functioning of genes in the context of the whole organism.
- Improving the performance characteristics of livestock.
- Formation of foreign proteins.
- Models of study of genetic diseases.
- Searching for possibilities for gene therapy of human diseases.



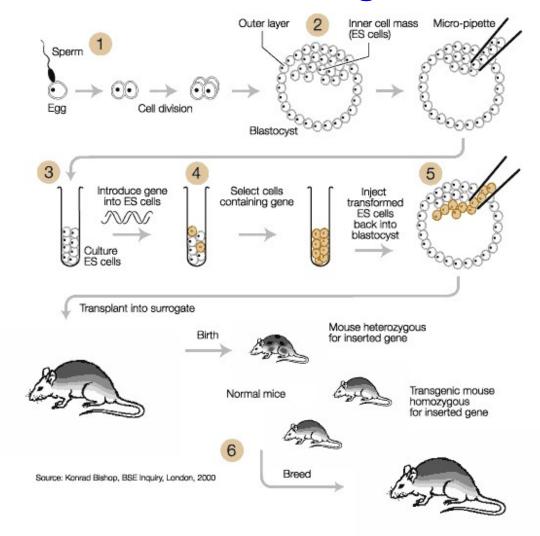
# Transgenic animals – mouse

- Mus musculus mouse
- Reproduction:
  - o 5-10 litters / year
  - o 5-10 pups / litter
  - 19-21 day gestation
  - Sexually mature at 7 weeks
  - 4-5 generations per year



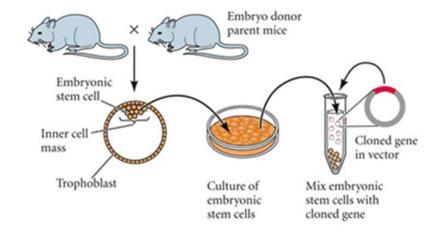


# Transgenic animals – transgenic mouse



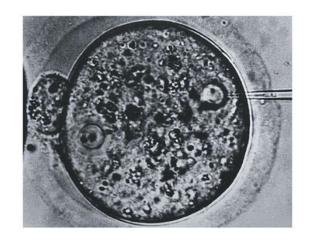


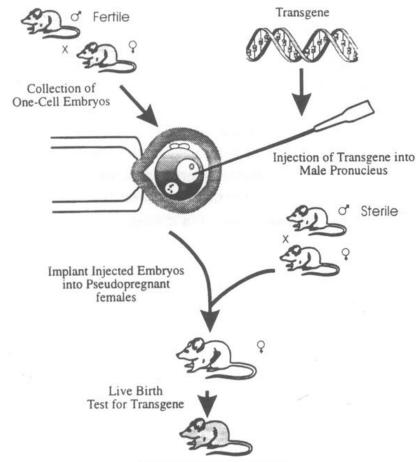
- Microinjection of cloned gene into nucleus of newly fertilized egg.
- Transfection incubate ES cells in solution that makes them take up the DNA, very inefficient need to identify cells that took up the DNA with reporter such as drug resistance.
- Electroporation a high voltage pulse "pushes" DNA into cells
- Retroviruses a more natural way or getting genes into cells.





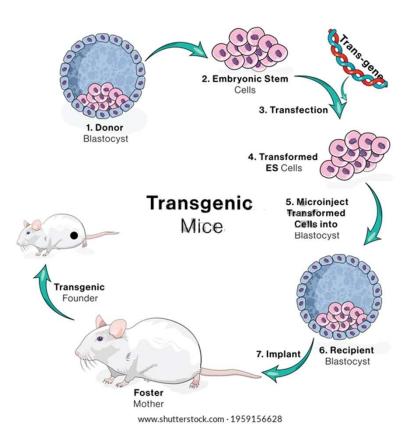
 Method used in producing transgenic mice by the microinjection of exogenous DNA into the pronuclei of fertilized eggs.





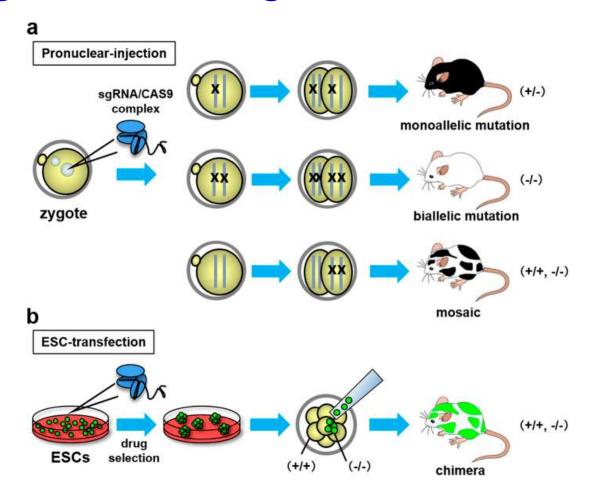
Transgenic Founder Animal





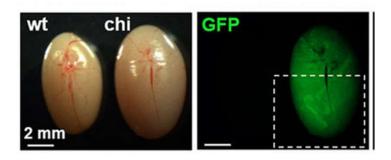
- Embryonic stem cells are cultured and subjected to DNA-mediated gene transfer or other manipulations.
- Cells with the desired genetic alteration are then inserted into blastocyst cavities, whereupon they resume normal development and produce a genetically mosaic mice, with some cells derived from the ES line, and the others from the injected blastocyst.
- Founders, usually males, are then bred to pass on the gene, provided the ES cells differentiate into sperm. The F1 hybrid then carries the new gene in all cells.



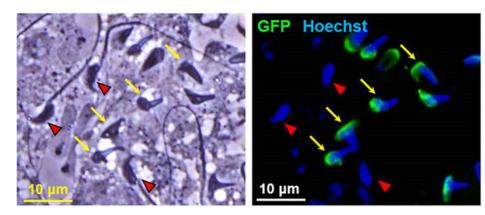




- Chimeric mice generated from ESCs carrying a biallelic mutation in *Cetn1* gene (em51/em52). Highly chimeric animals have darker coat color.
- Testes of B6D2F1 (wt) and chimeric mice (chi) were photographed under bright and fluorescent field.









#### Transgenic salmon

- AquAdvantage Salmon has been genetically engineered.
- It contains an rDNA construct that is composed of the growth hormone gene from Chinook salmon under the control of a promoter from ocean pout.







# **Transgenic animals**

Protein	Source	Against
Antithrombin III	Goat	Thrombosis
Tissue plasminogen activator	Sheep, pig	Thrombosis
á-antitrypsin	Sheep	Emphysema
Factor VIII, IX	Sheep, pig, cow	Hemophilia
á -Glucosidase	Rabbit	Pompe's disease
Fibrinogen	Cow, sheep	Wound healing
Glutarmic acid decarboxylase	Goat	Type 1 diabetes
Human serum albumin	Cow, sheep	Maintenance of blood volume
Human protein c	Goat	Thrombosis
Monoclonal antibodies	Chicken, cow, goat	Vaccine production
Pro 542	Goat	HIV
Lactoferrin	Cow	GI tract infection and infectious arthritis.



#### **Cloning of animals**

- Cloning is a controversial process that involves creating a genetically identical organism, or clone, from the genetic material of an individual.
- Clones are created by inserting a nucleus from an individual into an unfertilized egg. The egg will develop as if it had been fertilized to create a new living organism.
- In 1996, scientists were able to successfully clone the first mammal, a sheep, after 276 attempts.
- The successful clone was named Dolly. Since Dolly, scientists have successfully cloned other animals, such as, cats, cows, dogs, and rabbits.

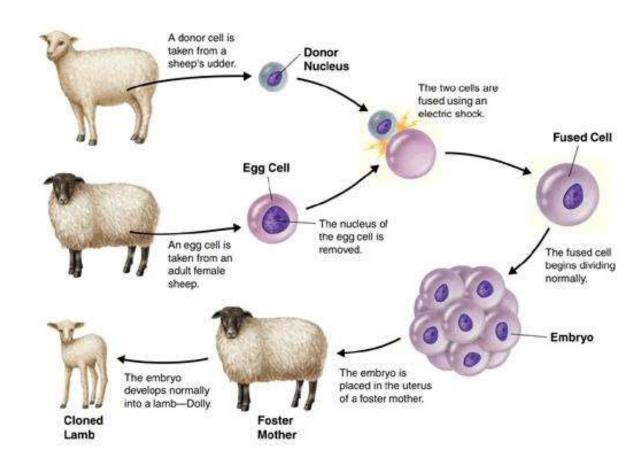


Ian Wilmut University of Edinburgh



## **Cloning of animals**





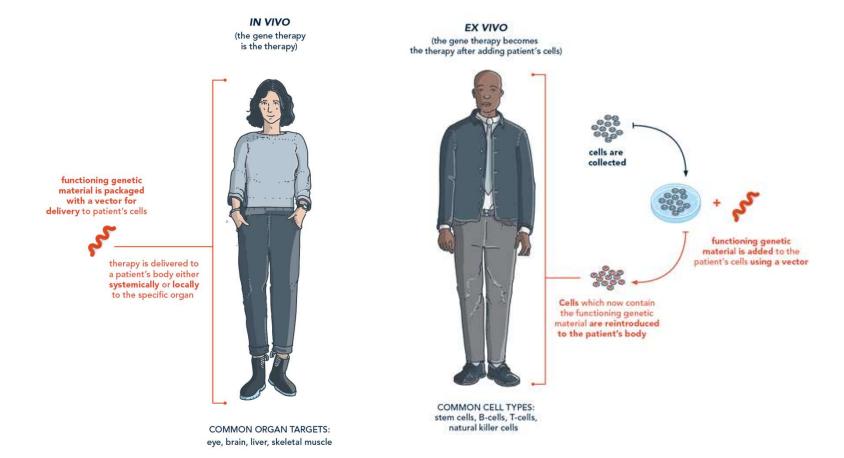


#### Gene therapy

- Gene therapy is a technique that modifies a person's genes to treat or cure disease. Gene therapies can work by several mechanisms:
  - Replacing a disease-causing gene with a healthy copy of the gene.
  - Inactivating a disease-causing gene that is not functioning properly.
  - Introducing a new or modified gene into the body to help treat a disease.
- Gene therapy products are being studied to treat diseases including cancer, genetic diseases, and infectious diseases.



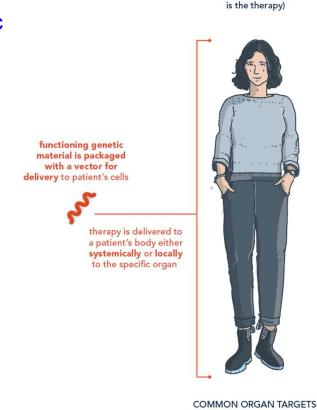
## **Road Map of Gene therapy**





### Road Map of Gene therapy

- In vivo delivery
- In vivo gene therapy refers to direct delivery of genetic material either intravenously (through an intravenous) or locally to a specific organ (eg, directly into the eye).
- In vivo gene therapy works through the help of a vector, which directly inserts functional copies of a gene into target cells to treat a mutated or missing gene.
- In vivo delivery of gene therapy has been proven in many areas of research. Some of the currently approved gene therapies deliver genetic material in vivo. Targeted in vivo gene therapy will continue to evolve as scientists continue to refine methods of gene delivery.



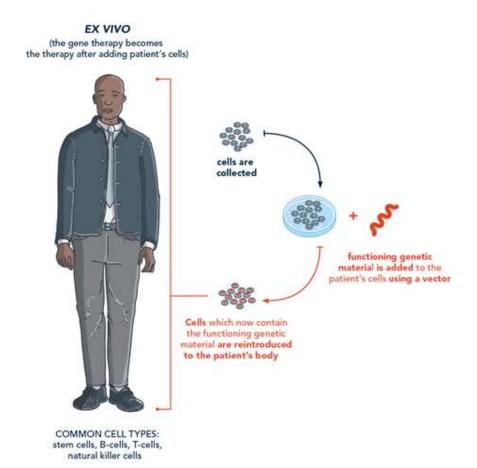
IN VIVO

(the gene therapy

eye, brain, liver, skeletal muscle

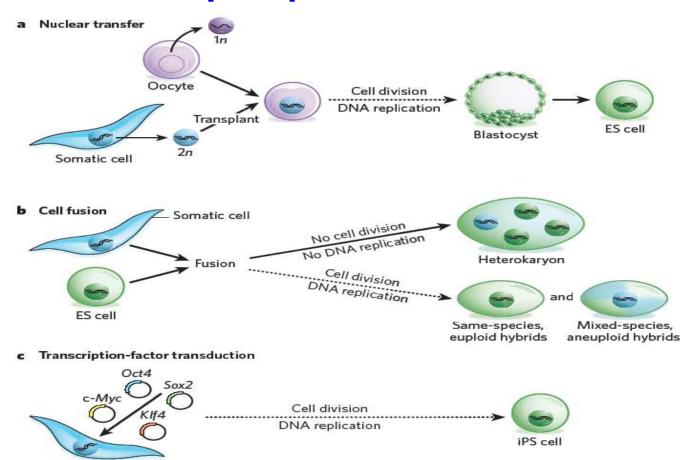


### **Road Map of Gene therapy**



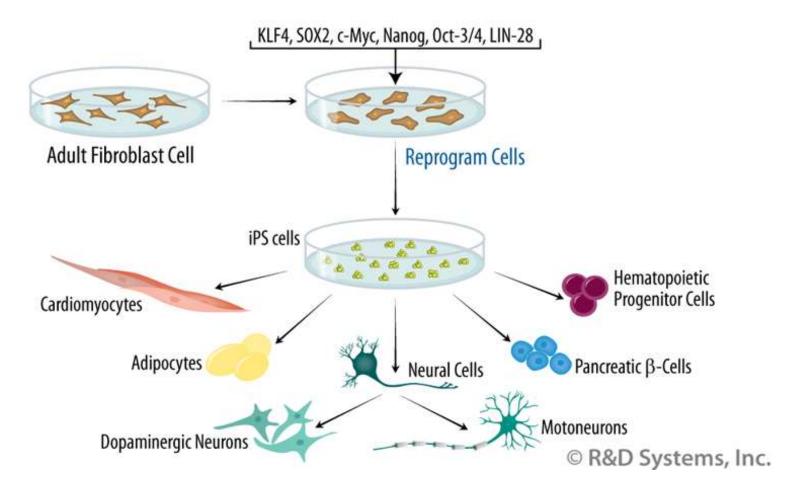
- Ex vivo delivery
- Ex vivo gene therapy refers to the process of removing specific cells from a person, genetically altering them in a laboratory, and then transplanting them back into the person.
- Ex vivo gene therapy works by genetically modifying a patient's stem cells, which then replace target cells that have a missing or malfunctioning gene.
- Today, ex vivo gene therapy is most frequently applied to hematopoietic stem cells (HSCs); this is used to treat blood and immunological diseases as well as genetic diseases that affect tissues and organs that can be reached by blood cells.

#### Induced pluripotent stem cells – iPSC





## Induced pluripotent stem cells – iPSC

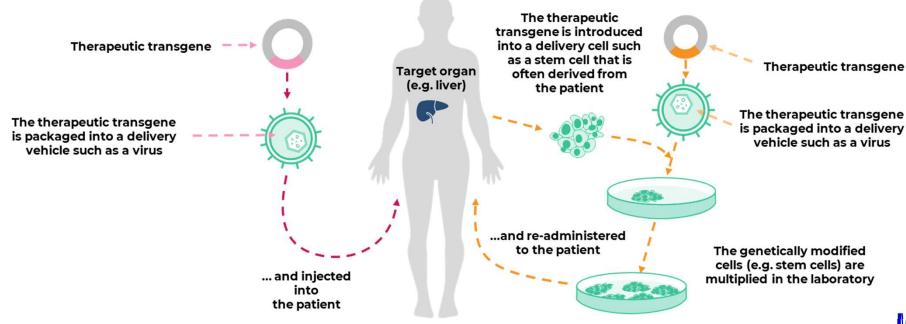




#### **Gene therapy – delivery**

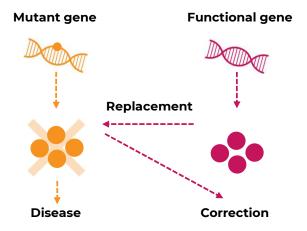
- Direct Delivery
- The therapeutic gene is transferred directly to target cells in the body.

- Cell-Based
- Cells are extracted from the patient, modified with the therapeutic gene, and injected back into the patient.

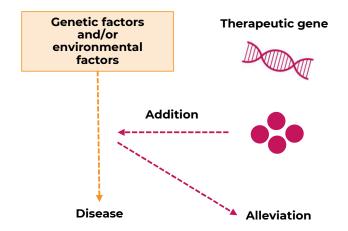


## Gene therapy – Approach I

- Gene Replacement Therapy
- For monogenic diseases; involves replacing a mutated gene that causes disease with a healthy gene.



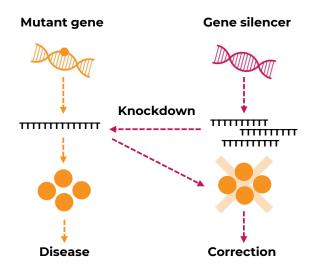
- Gene Addition
- For complex and infectious diseases; introduces a new gene into the body to help fight a disease, often to supplement a targeted therapeutic agent.



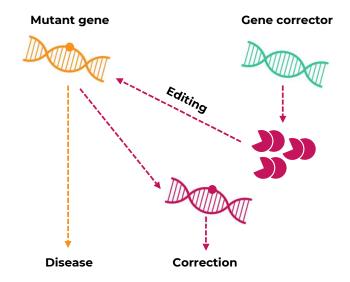


## **Gene therapy – Approach II**

- Gene Inhibition or "Knockdown"
- Inactivating a mutated gene that is overproducing its product by targeting RNA.



- Gene Editing
- Making a targeted change to the gene sequence.



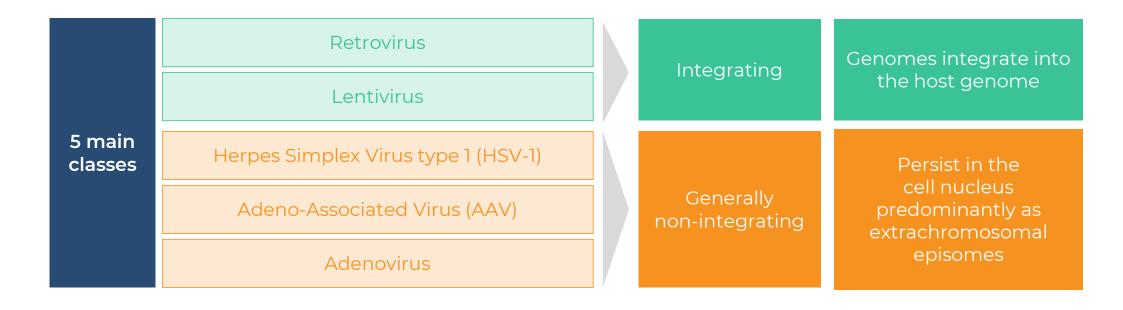


#### Gene replacement therapy

- The aim of Gene Replacement Therapy GRT is to provide sufficient gene expression in enough targeted cells to ameliorate or correct dysfunctional phenotype.
- Suitable for:
- Recessive disorders, single gene mutations, and X-linked diseases.
- Generally unsuitable for:
- Dominant disorders where an errant gene codes for destructive or interfering proteins.
- Gene therapy:
  - Either directly in vivo or through ex vivo cell therapy.
  - Vehicle/vector used to deliver therapeutic transgene.



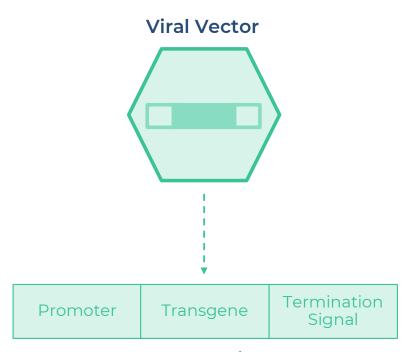
## **Gene therapy – viral vectors**





## **Gene targeted therapy – in vivo approach**

- Three key components:
- the vector, or vehicle, which is injected into the patient and by which a transgene is delivered to the targeted cells
- the transgene, which is a sequence of complementary DNA (cDNA) coding the replacement gene
- the promoter, which is the DNA sequence that acts as a "turn on" switch and modulates the expression of the transgene.
- Also typically includes, either:
- A termination signal to end gene transcription.
- Inverted terminal repeats (ITRs) at either end of the cassette to allow for synthesis of complementary DNA.



**Transgene Expression Cassette** 

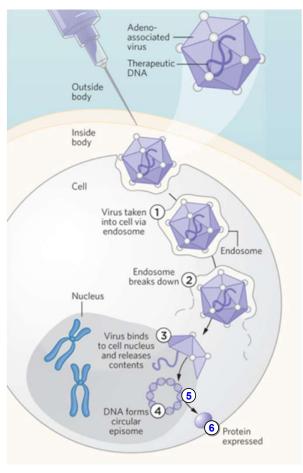


### **Gene targeted therapy – AAV**

- Adeno-associated virus AAV is a member of the parvovirus family of single-stranded small DNA viruses.
- Requires a helper virus such as <u>adenovirus</u> or <u>herpes simplex virus</u> for replication1
- Considered a Risk Group 1 microorganism infection with AAV is not associated with disease in humans.
- Has several serotypes that impact tropism (susceptible tissues), but all appear to be nonpathogenic.
- rAAV based vectors used in GRT have wild-type viral genes removed and typically exist as extrachromosomal episomes.
- AAV can efficiently infect both non-dividing and dividing cells.
- AAV is typically transmitted by respiratory and gastrointestinal routes.



## **Gene targeted therapy – AAV**



- AAVs deliver genes without integrating them into the genome:
- 1) Virus is taken into the cell via the endosome.
- 2) The endosome breaks down.
- 3) Therapeutic DNA enters cell nucleus as a double-stranded molecule ready for transcription.
- 4) Therapeutic DNA forms a circular episome.
- 5) Upon promoter activation, transcription occurs.
- 6) The resulting transcript leaves the nucleus and travels to the ribosome for translation (protein synthesis).



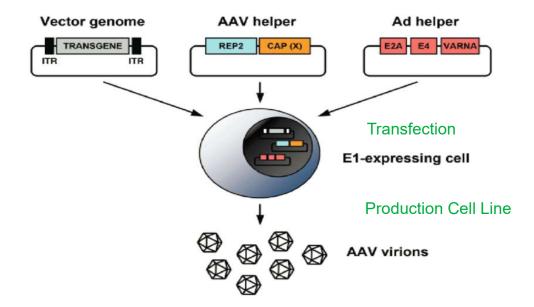
### **Gene targeted therapy – - AVV production overview**

The AAV production process is complex and sophisticated.

- 1. Transgene Development
- ----- 2. AAV GRT Production

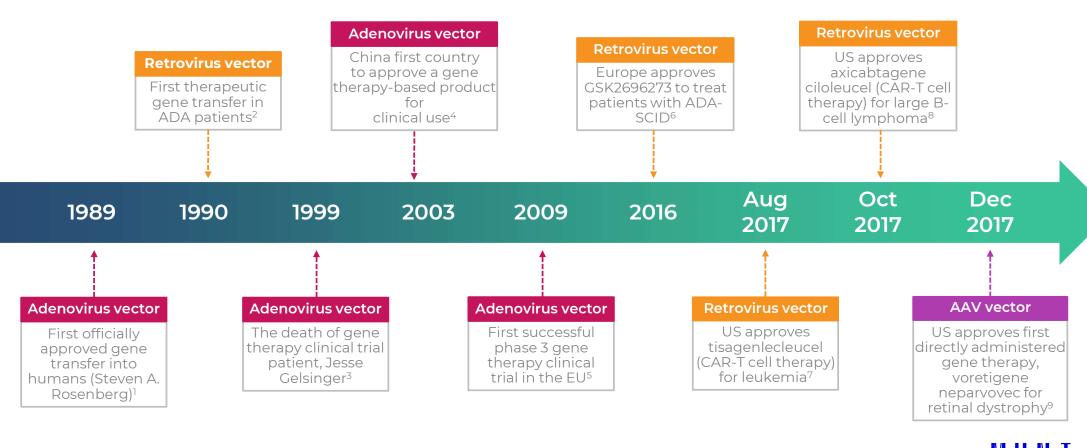
- 3. Concentration & Purification
- 4. Quality Assurance

 cDNA is reverse transcribed from human mRNA (no introns).





#### **Gene therapy – history of delivery vectors**



## **Gene therapy - Approved genes therapies**

FDA-approved therapy	Indication (approval date)	Delivery	Viral vector
Imlygic (talimogene laherparepvec)	Melanoma (2015)	In vivo	HSV-1
Kymriah (tisagenlecleucel)	B-cell precursor acute lymphoblastic leukemia (2017); diffuse large B-cell lymphoma (2018)	Ex vivo	Lentivirus
Yescarta (axicabtagene ciloleucel)	Large B-cell lymphoma (2017)	Ex vivo	Retrovirus
Luxturna (voretigene neparvovec-rzyl)	Biallelic RPE65 mutation-associated retinal dystrophy (2017)	In vivo	AAV
EMA-approved therapy	Indication (approval date)		
Imlygic	Melanoma (2015)	In vivo	HSV-1
Strimvelis	ADA-SCID (2016)	Ex vivo	Retrovirus
Zalmoxis	Adjunctive treatment in HSCT in high-risk blood cancer (2016)	Ex vivo	Retrovirus



#### THANK YOU FOR YOUR ATTENTION



# **GENE THERAPY**

