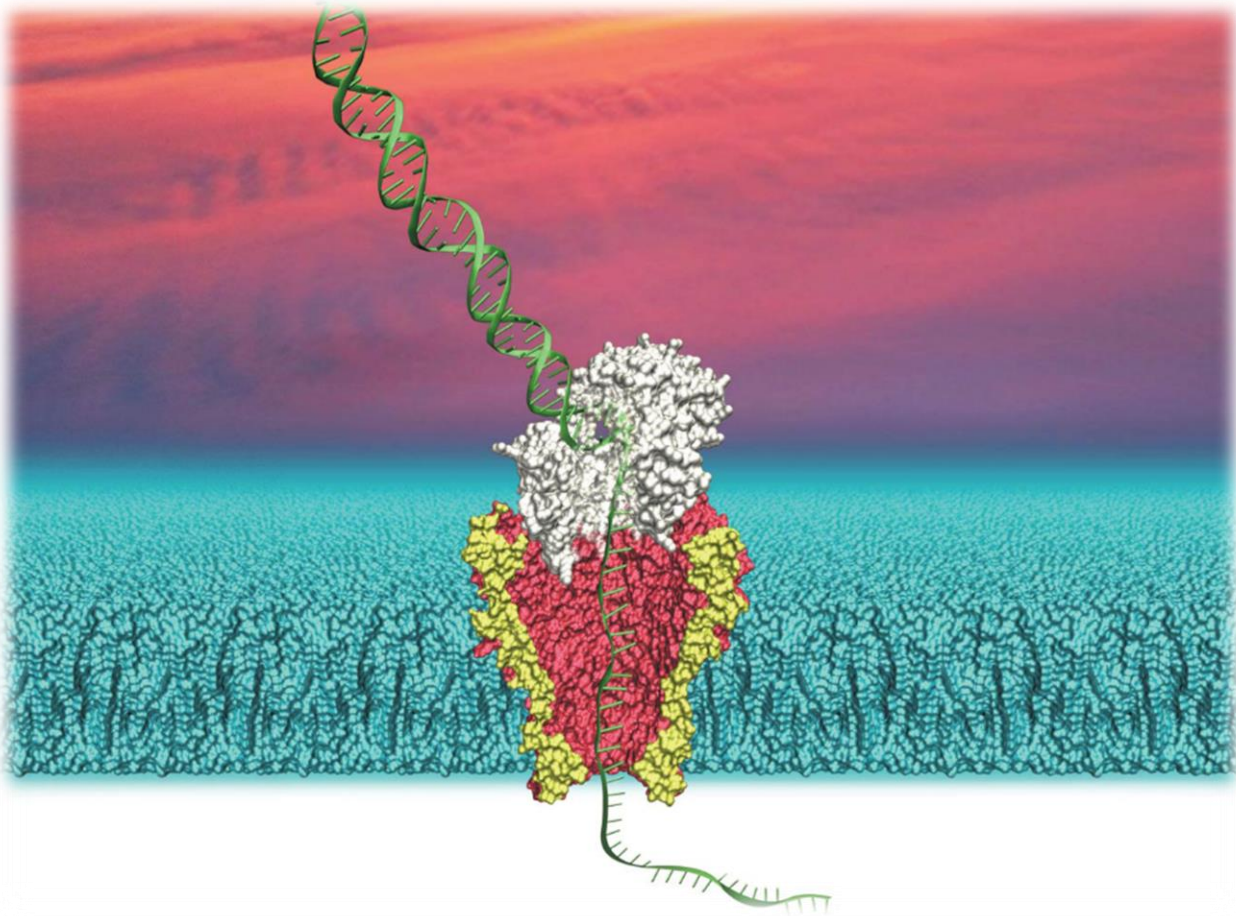




Chapter 19 DNA Technology

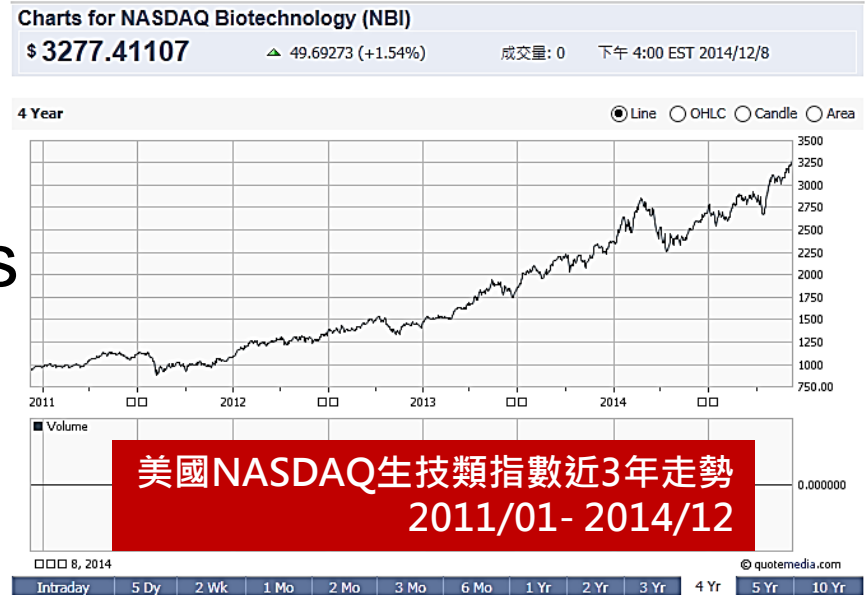


Overview: The DNA Toolbox

- One of the greatest achievements of modern science has been the **sequencing of the human genome (2003)**--- depended on advances in **recombinant DNA technology**

 - In **recombinant DNA**, nucleotide sequences from two different sources, often two species, are combined *in vitro* into the same DNA molecule

 - Large-scale sequencing of the genomes of various species was under way since then
-

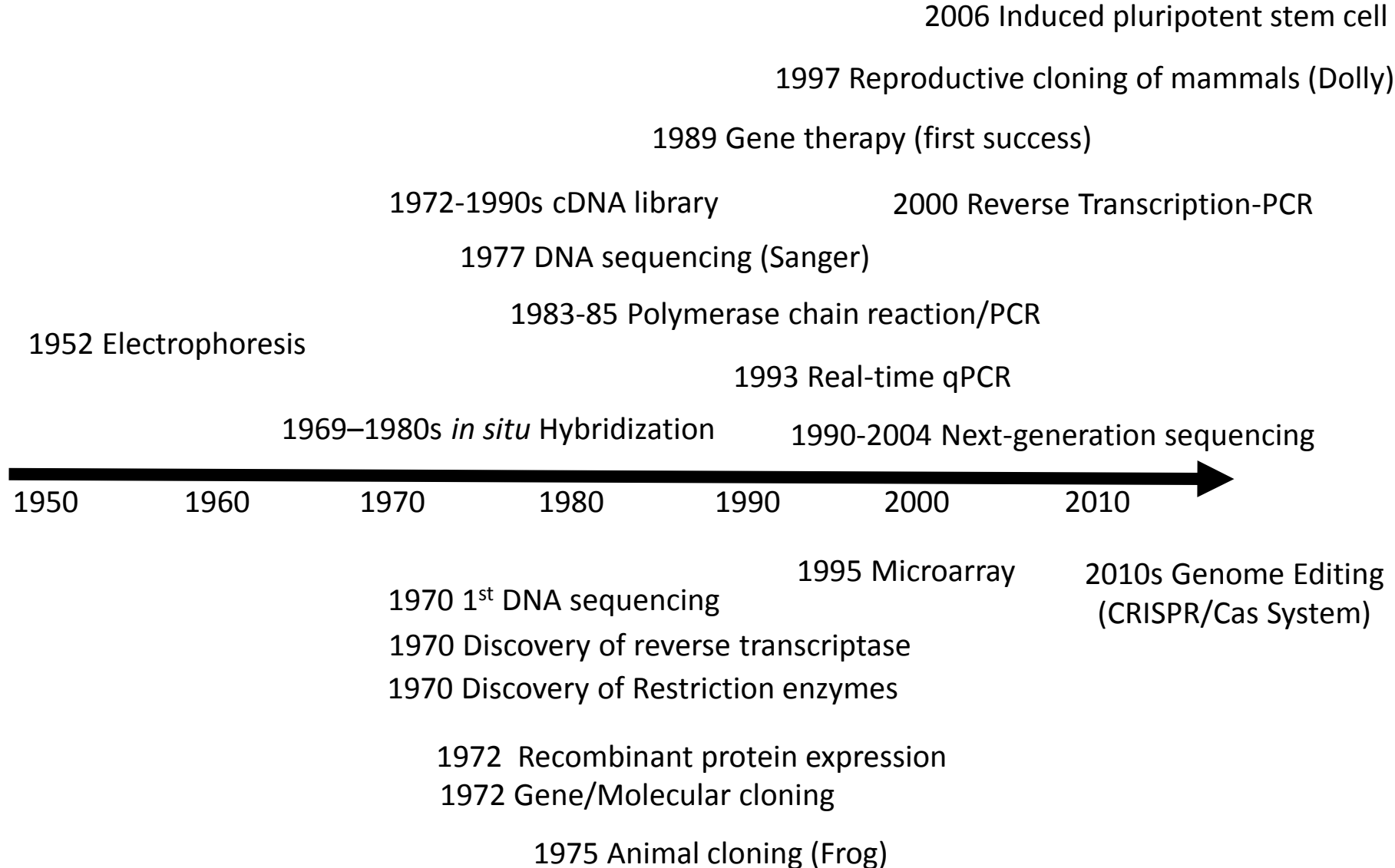
DNA technology & Biotechnology

- **Biotechnology** is the manipulation of organisms or their components to make useful products



The applications of **DNA technology** affect everything from agriculture, to criminal law, to medical research. Thus, DNA technology has revolutionized biotechnology

Milestones in DNA Technology



Concept 19.1: DNA sequencing and DNA cloning are valuable tools for genetic engineering and biological inquiry

- A gene's complete nucleotide sequence can be determined using a process called **DNA sequencing** 定序
 - The first automated procedure was based on a technique called dideoxy or chain termination sequencing, developed by Frederick Sanger
 - In the last 15 years, “next-generation sequencing” techniques have been developed that are much faster
-

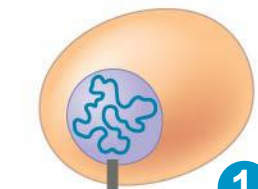
Next-Generation Sequencing (NGS)

次世代定序

- “Next-generation sequencing” techniques use a single template strand that is immobilized and amplified to produce an enormous number of identical fragments
- Thousands or hundreds of thousands of fragments (400–1,000 nucleotides long) are **sequenced in parallel**
- This is considered as a type of **high-throughput technology**
高通量

Next-Generation Sequencing (NGS) 次世代定序

Technique



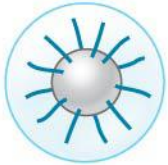
1 Genomic DNA is fragmented.



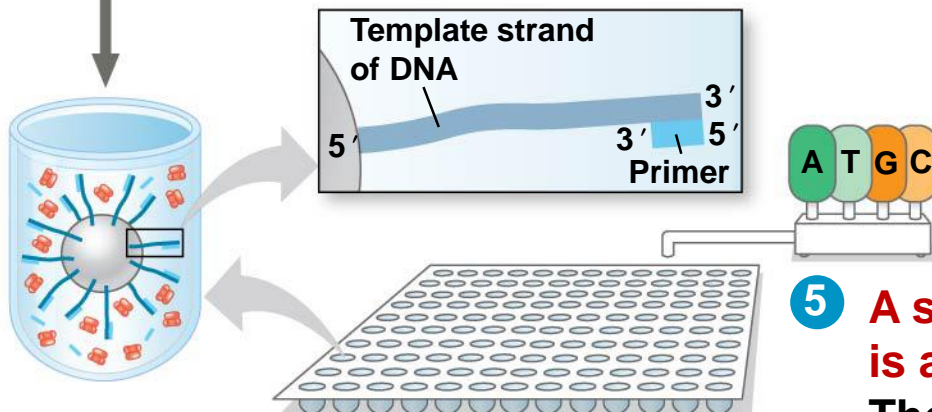
2 Each fragment is isolated with a bead.



3 Using PCR, 10^6 copies of each fragment are made, each attached to the bead by 5' end.



4 The bead is placed into a well with DNA polymerases and primers.



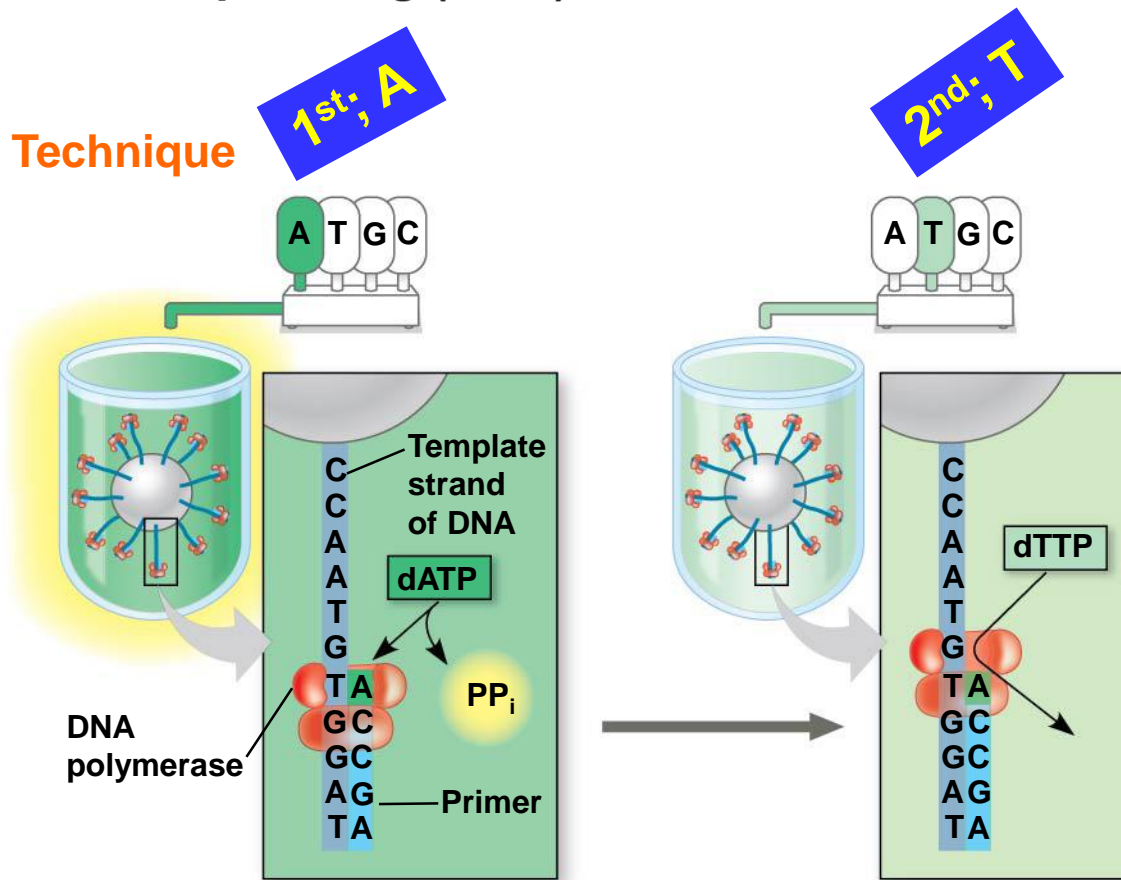
一次反應只用一種 去氧核苷酸
dATP, dGTP, dTTP, dCTP (dNTP)
沖洗後，才換下一種; 4個輪換

5 A solution of each of the four nucleotides is added to all wells and then washed off. The entire process is then repeated.

Next-Generation Sequencing (NGS)

次世代定序

Technique



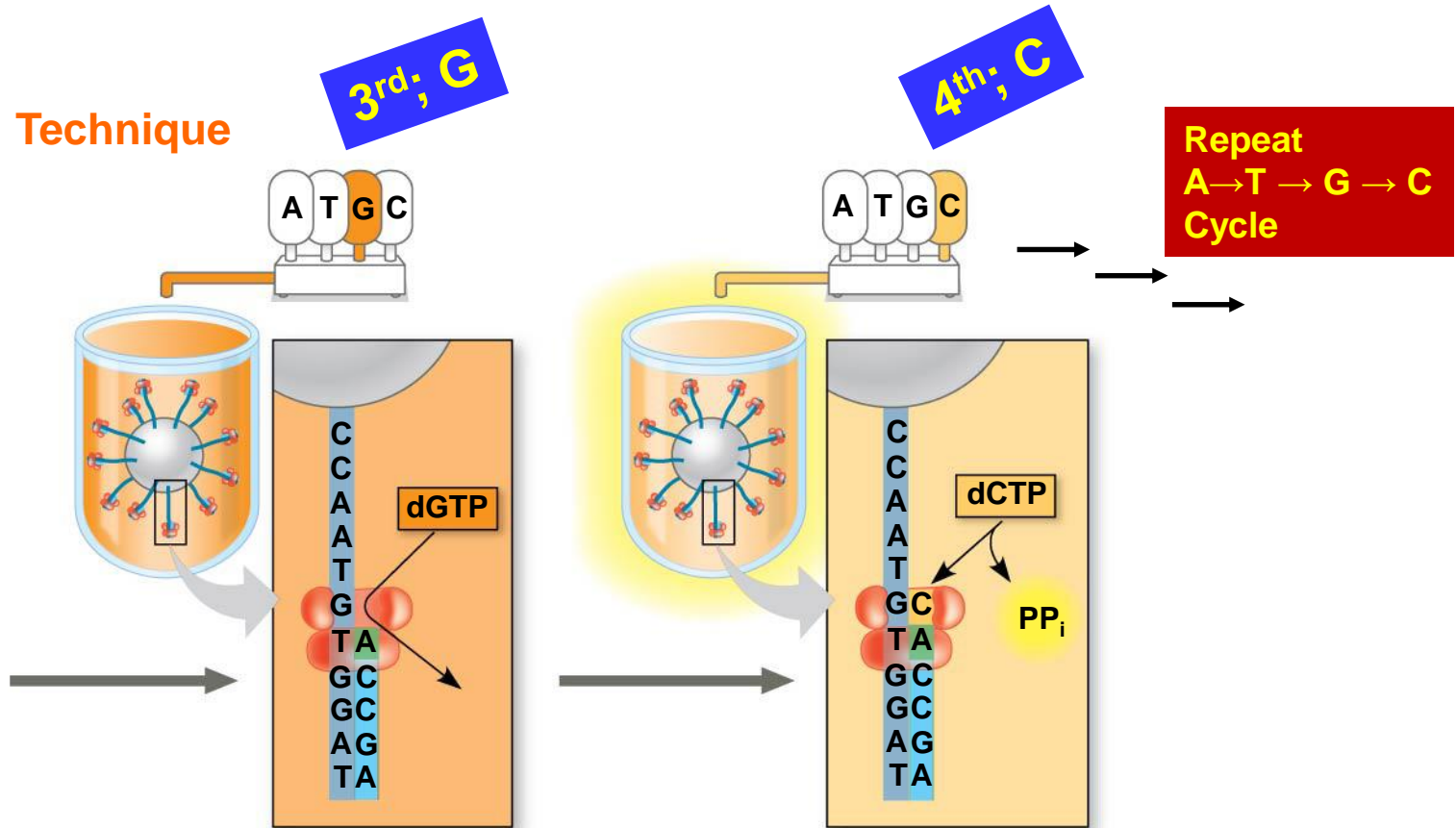
⑥ If a nucleotide is joined to a growing strand, **PP_i is released, causing a flash of light that is recorded.**

⑦ If a nucleotide is not complementary to the next template base, no PP_i is released, and no flash of light is recorded.

No match (no DNA elongation), No signal

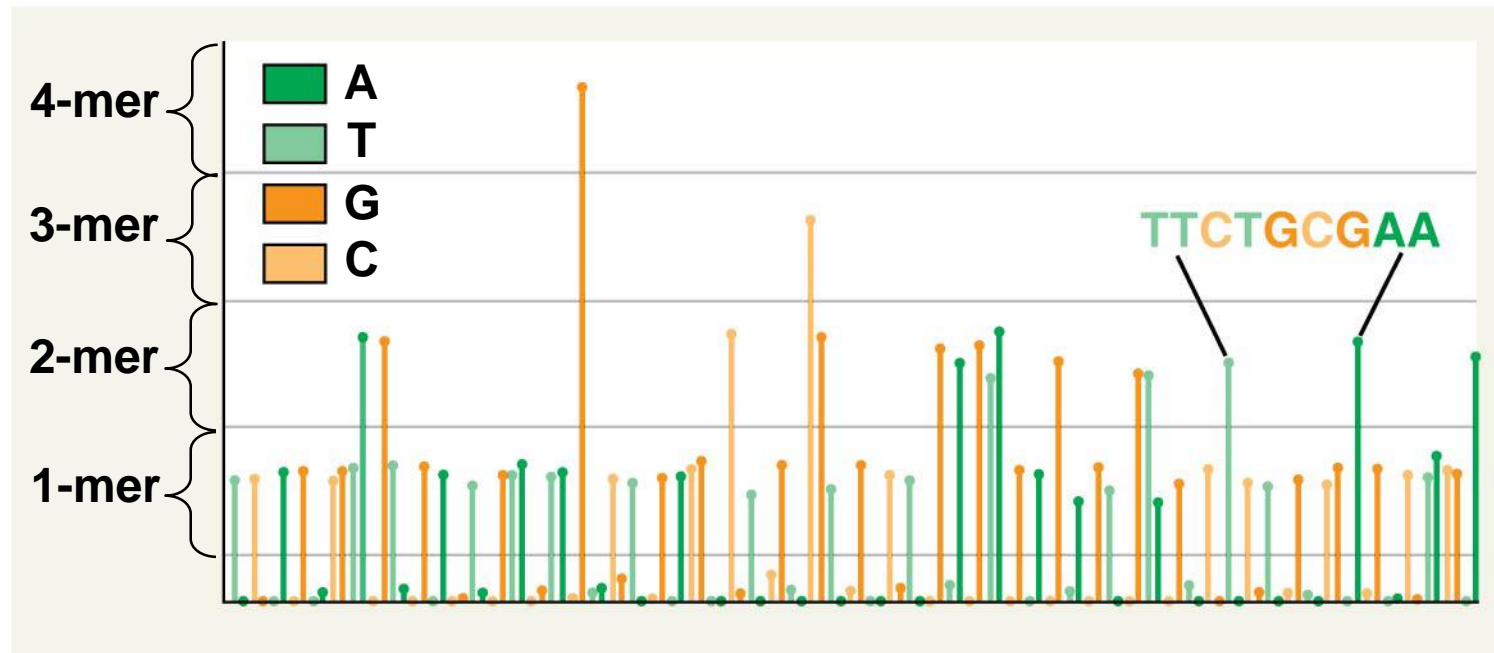
Next-Generation Sequencing (NGS)

次世代定序



- 8 The process is repeated until every fragment has a complete complementary strand. The pattern of flashes reveals the sequence.

Results

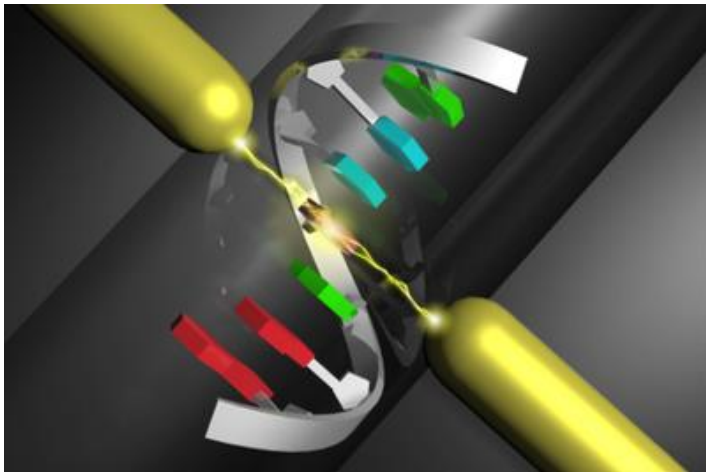


dATP, dGTP, dTTP, dCTP 4個依設定輪換，
有反應 (complementary base pairing)，
即記錄藉以完成定序

And the sequence technology keeps advancing....

- In “**third- and fourth-generation sequencing,**” the techniques used are even faster and less expensive than the previous, while providing spatial information

請自主學習



Real-time sequencing of single DNA (or RNA) molecules

Reference:

- Schadt EE et al. **A window into third-generation sequencing.** Hum. Mol. Genet. (2010)
- 19(R2):R227
- Liang J et al. **Single-cell sequencing technologies: current and future.** J Genet Genomics. (2014)

DNA Cloning

- **How to do DNA cloning:**
 - (1) 目標 : DNA of interest
 - (2) 載體 : Plasmids (cloning vector):
replication origin, multiple cloning sites,
selection marker (for example: ampicillin resistance gene; LacZ gene : β -galactosidase digests X-gal into blue product)
 - (3) 宿主 : Host: i.e. bacteria (E. Coli), yeast
 - Cloned genes are useful for making copies of a particular gene and producing a protein product
-

Figure 19.5a

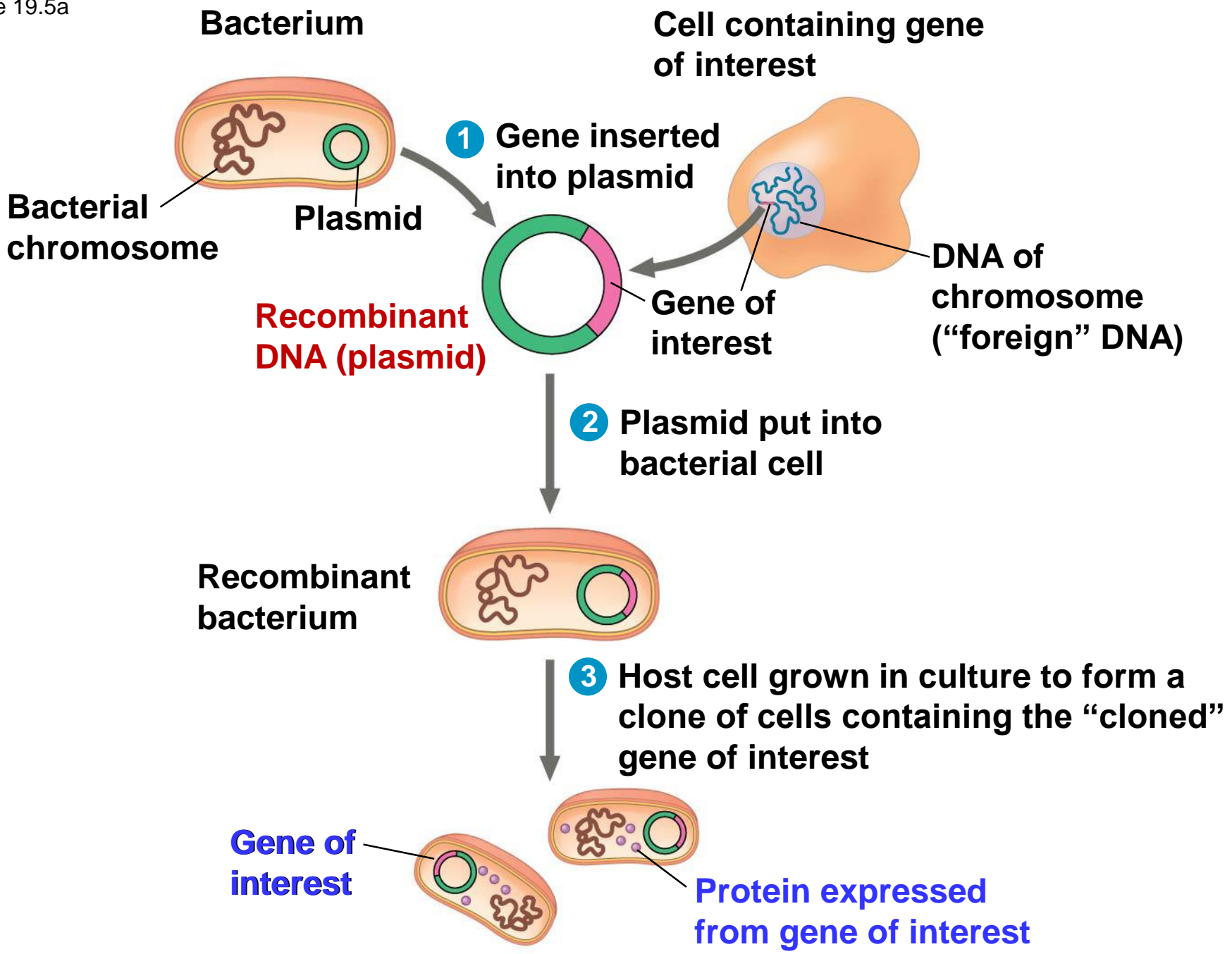
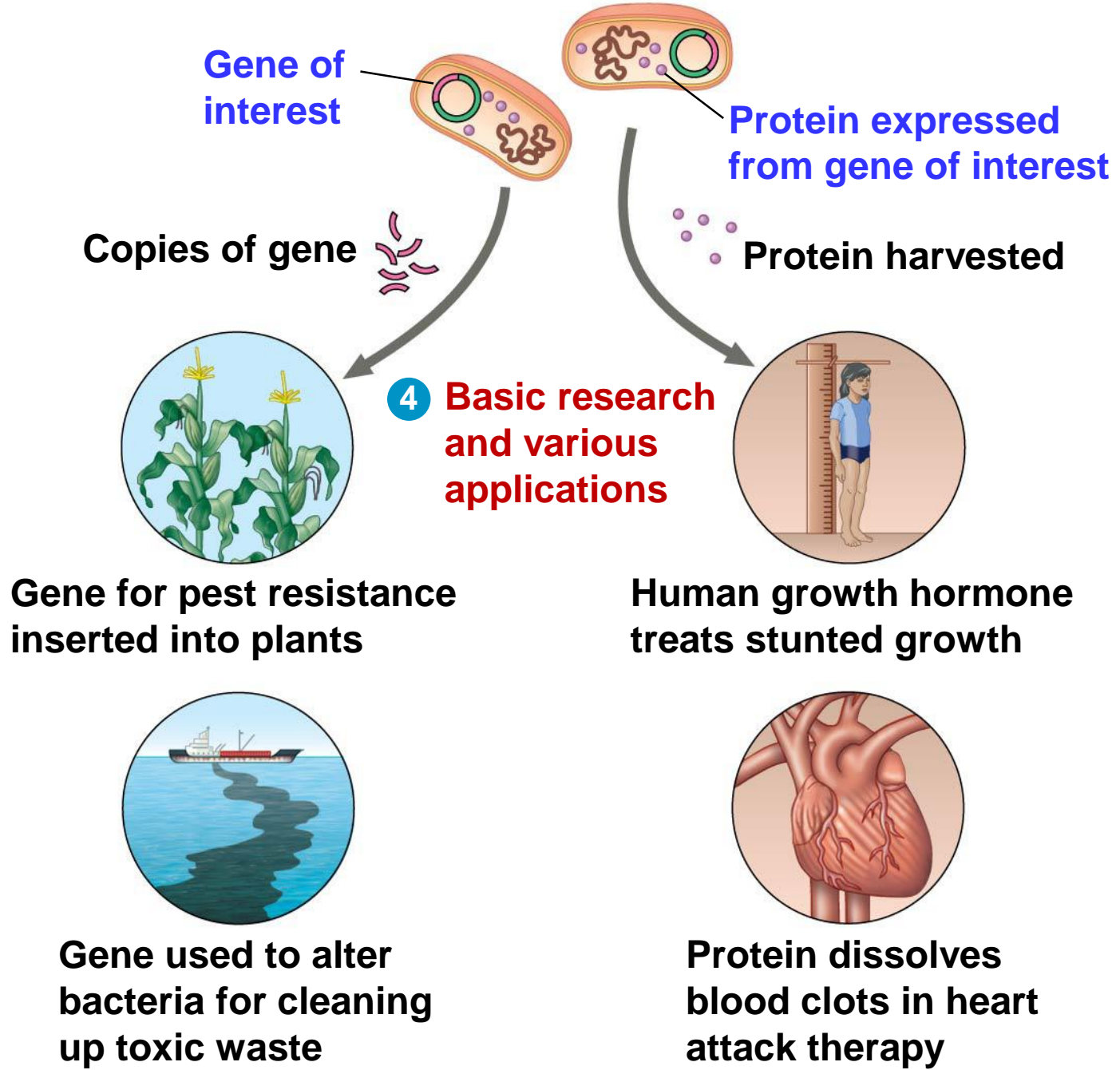
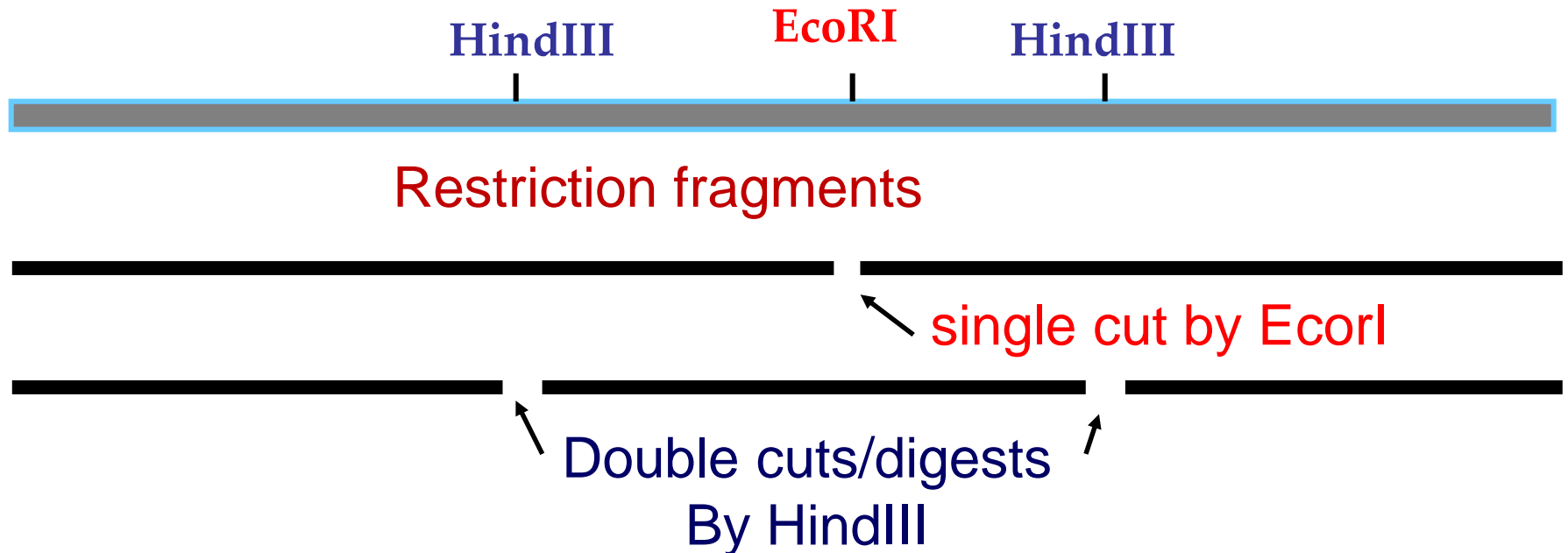


Figure 19.5b



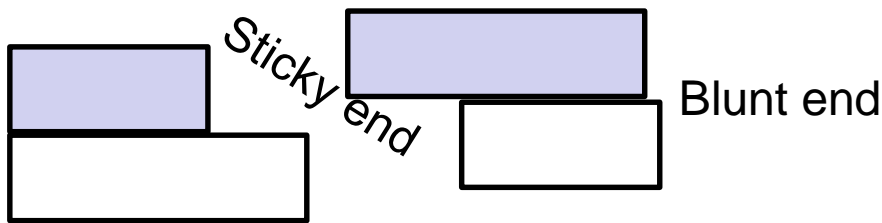
How to Make Recombinant DNA: **Restriction Enzymes**

(1) Bacterial **restriction enzymes** -- cut DNA molecules at a limited number of **specific** DNA sequences (4-8 nucleotides), called **restriction sites** – GAATTC (EcoRI),



Using Restriction Enzymes to Make Recombinant DNA

- The most useful restriction enzymes cut DNA in a **staggered way**, producing fragments with “**sticky ends**” that bond with complementary sticky ends of other fragments



PLAY

Animation: Restriction Enzymes

DNA ligase

- **DNA ligase** is an enzyme that seals the bonds between restriction fragments

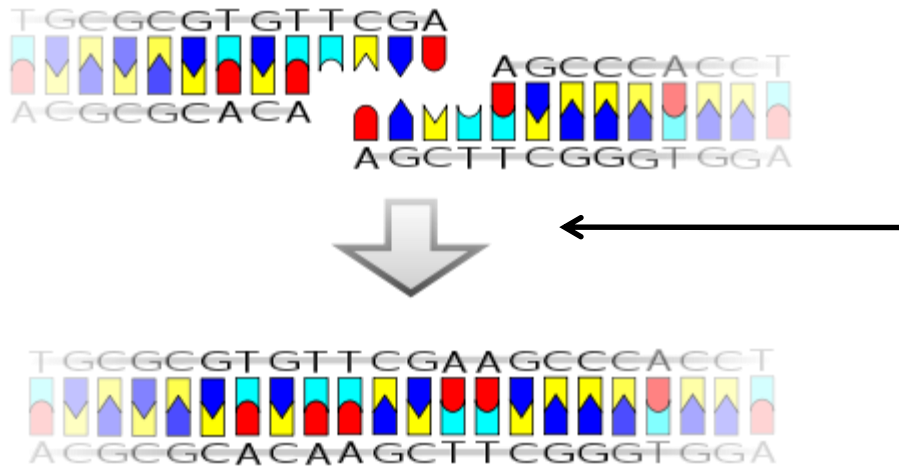
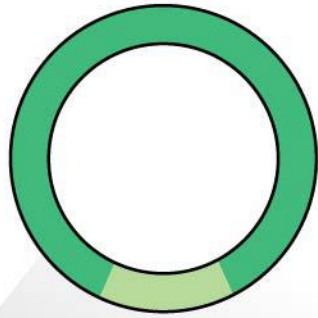


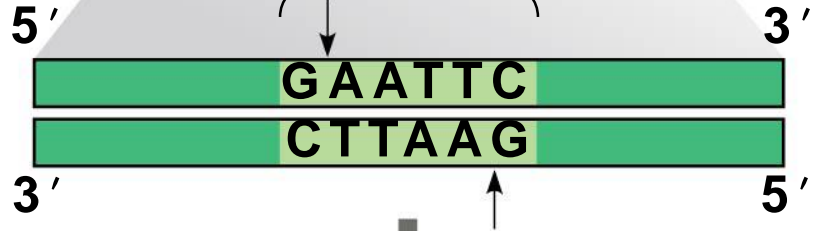
Figure 19.6a

**Bacterial
plasmid**



Restriction site

DNA



1 **Restriction enzyme** cuts the sugar-phosphate backbones at each arrow.

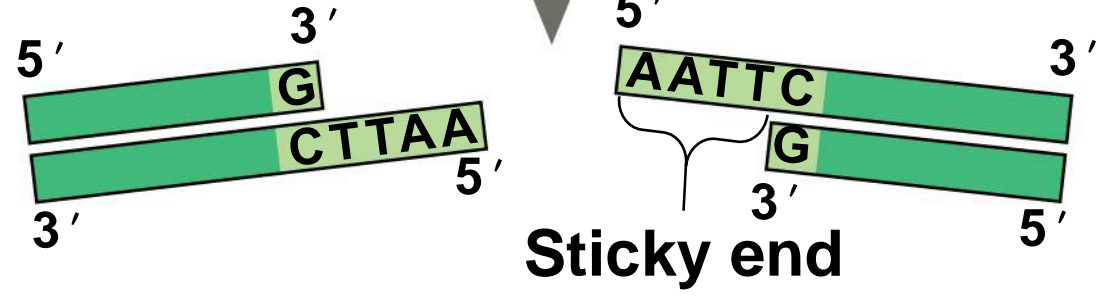
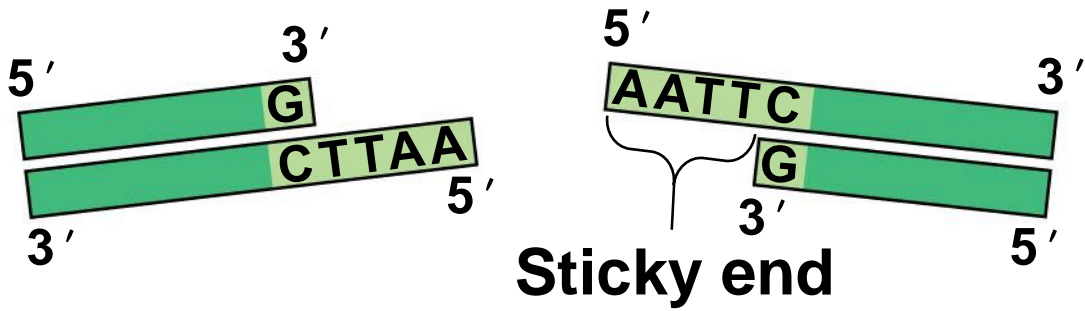
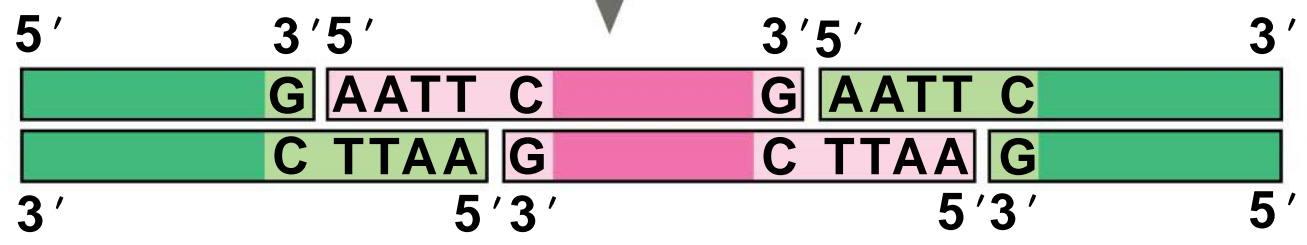
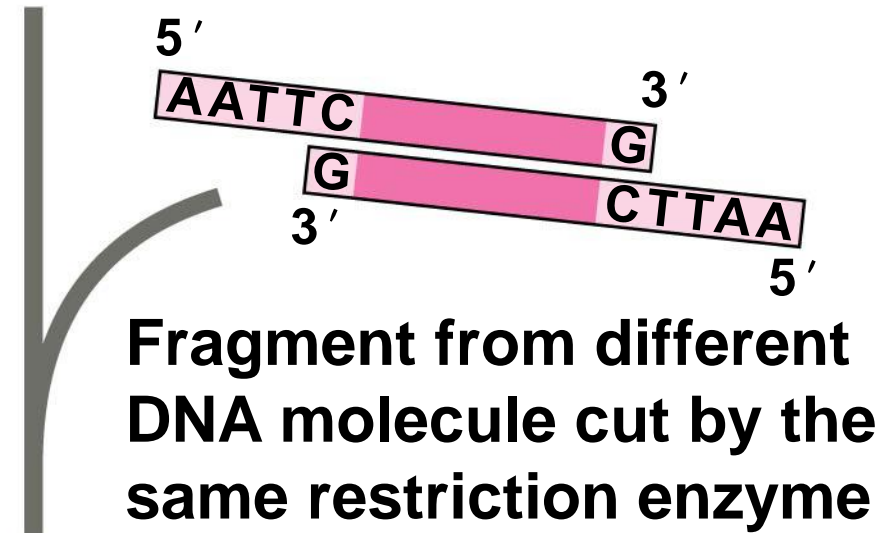


Figure 19.6b

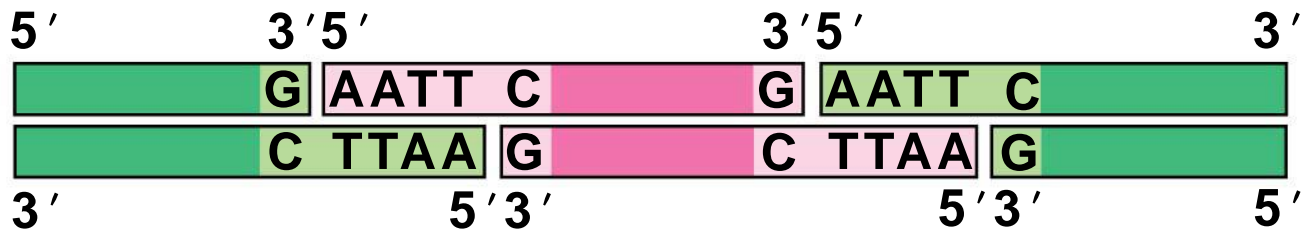


2 Base pairing of sticky ends produces various combinations.



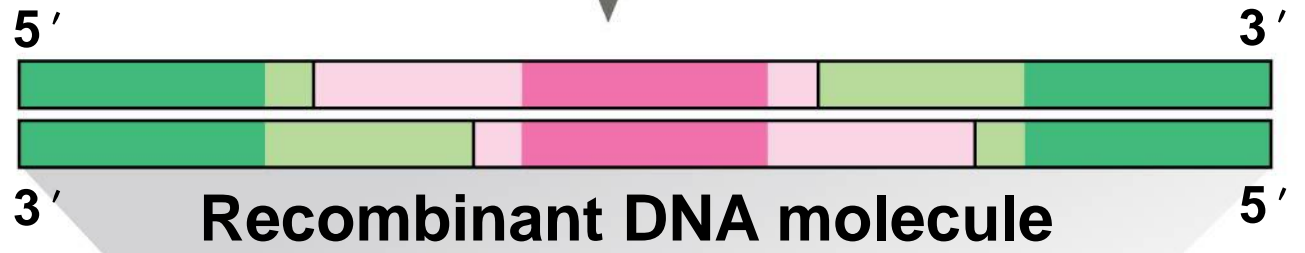
One possible combination

Figure 19.6c



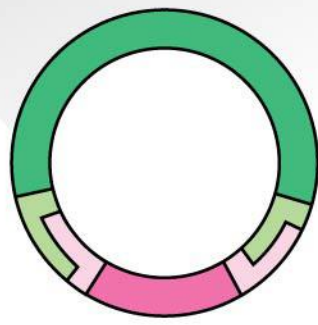
One possible combination

3 DNA ligase seals the strands



Recombinant DNA molecule

Recombinant plasmid



DNA assay: DNA Gel Electrophoresis 膠體電泳

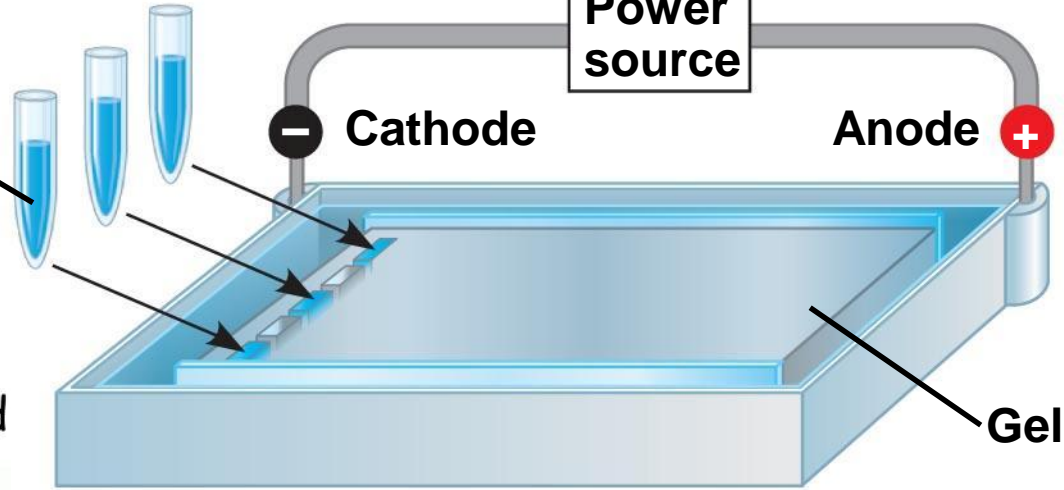
- One indirect method of rapidly analyzing and comparing DNA molecules is **gel electrophoresis**
 - This technique uses a gel as a **molecular sieve** (分子篩) to separate nucleic acids or proteins by size
 - A **electrical current** is applied that causes charged molecules to move through the gel
 - Molecules are **sorted into “bands” by their size**
-

Fig. 20-9a

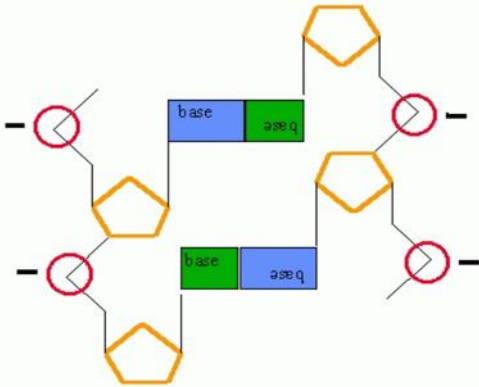
Gel electrophoresis

TECHNIQUE

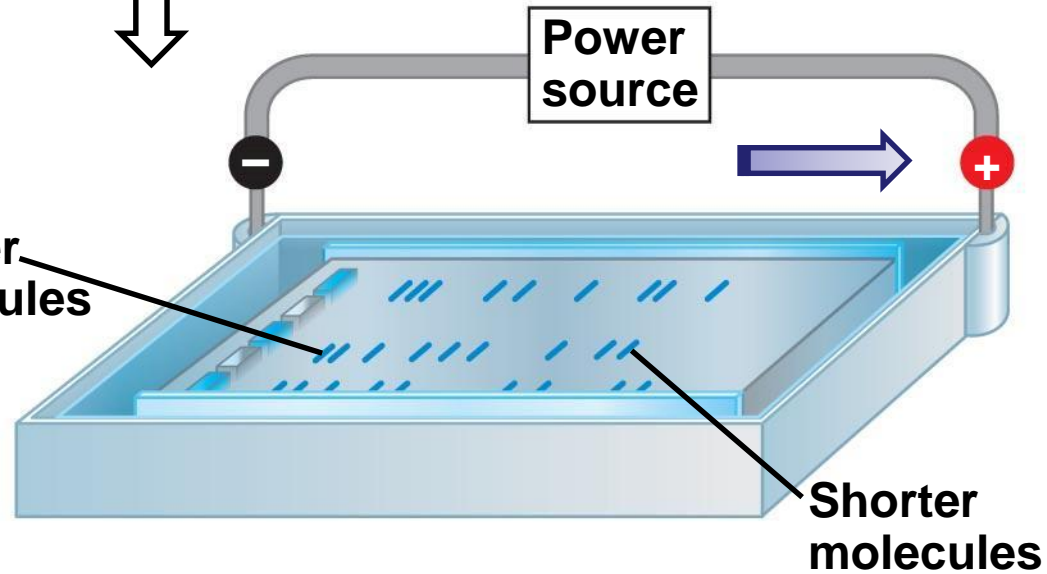
Mixture of DNA molecules of different sizes



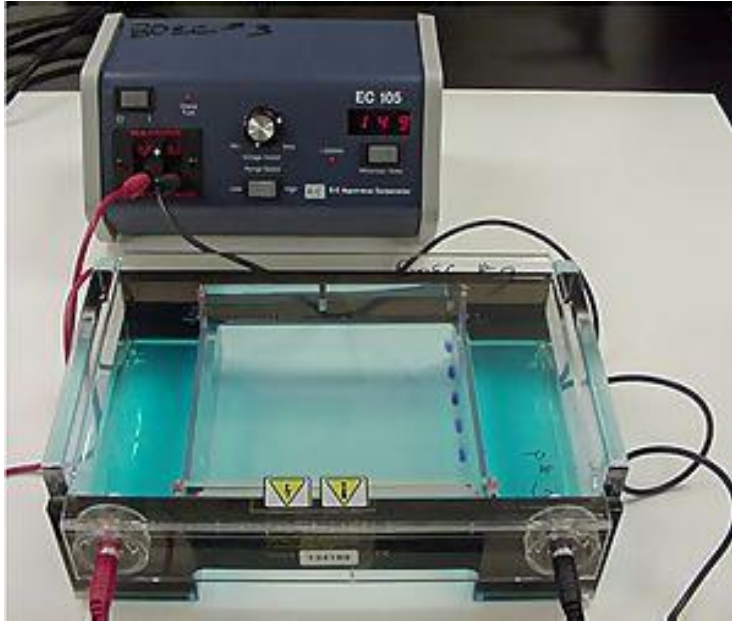
DNA negatively charged



Longer molecules



Gel electrophoresis



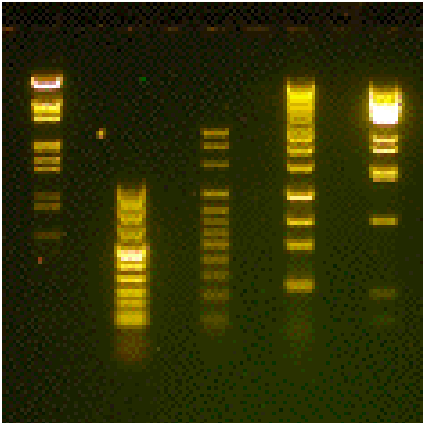
Visualization by **EtBr staining**
(old method)

Ethidium Bromide is now be replaced by safer fluorescent staining reagents, such SYBR gold.

Longer
DNA sequence

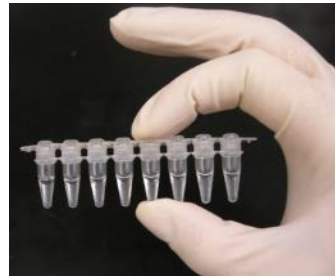


Shorter
DNA sequence



Amplifying DNA *in Vitro*: The Polymerase Chain Reaction (PCR)

- The **polymerase chain reaction, PCR**, can produce many copies of a specific target segment of DNA
- A three-step cycle — **heating, cooling, and replication** — brings about a chain reaction that produces an exponentially growing population of identical DNA molecules



Significance of PCR in artificial gene replication

With **polymerase chain reaction (PCR)**, any specific segment of the target sequence within a DNA sample can be **copied many times (amplified)** completely *in vitro*.

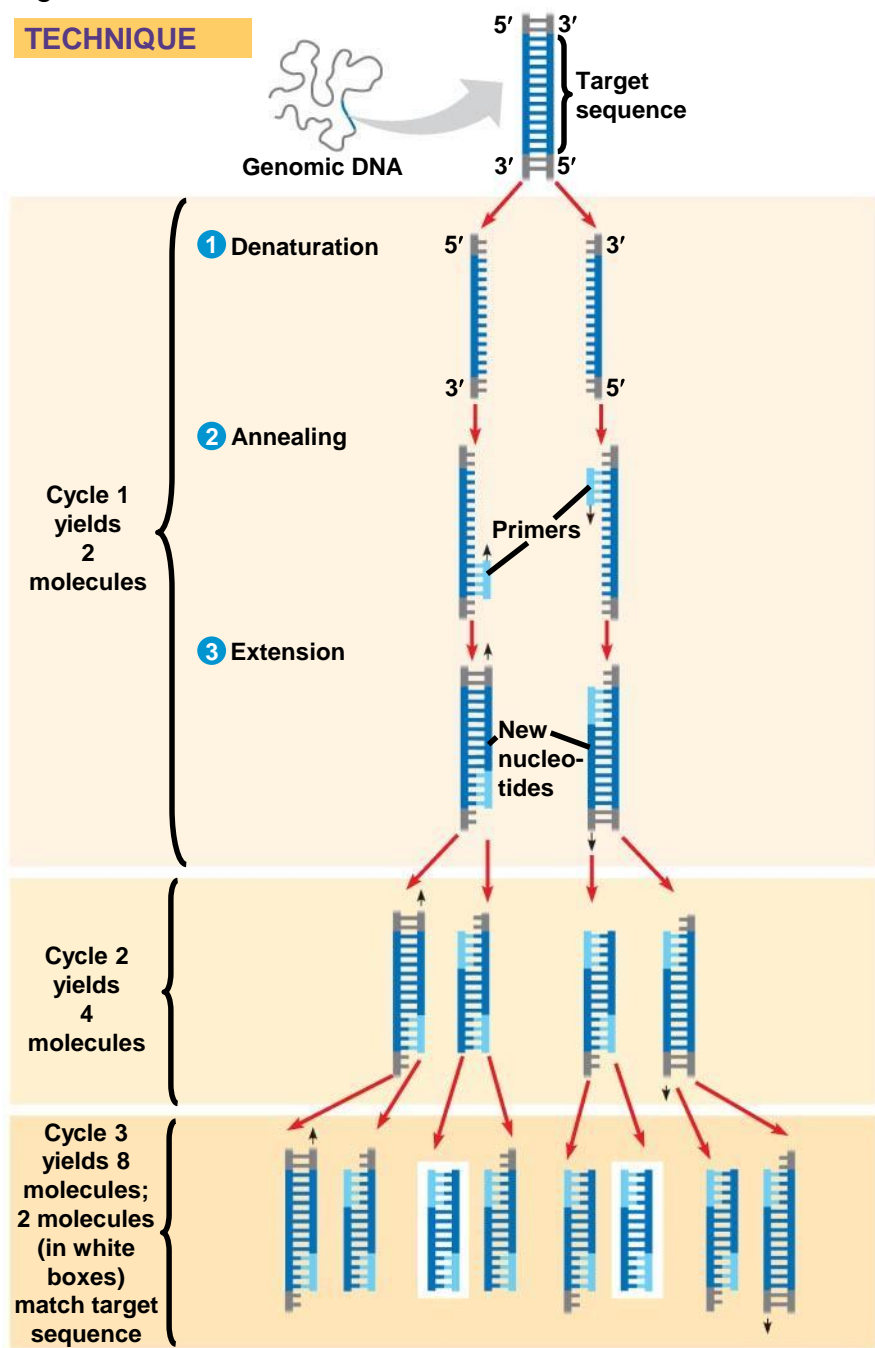
Key components in a PCR reaction:

--Uses **primers** (with unique restriction sites to allow the product to be cloned into plasmid vectors) that bracket the desired sequence

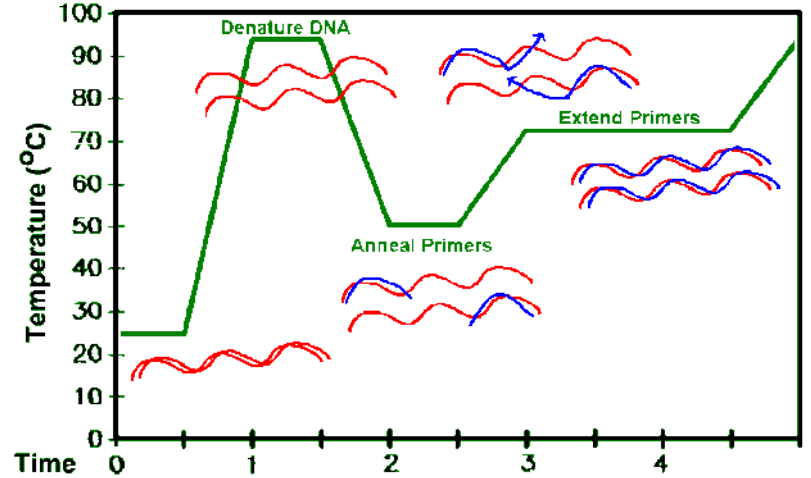
--Uses a **heat-resistant DNA polymerase** (active at high temperature) isolated from prokaryotes living in hot spring: extend primers in 5' → 3' direction

Fig.19-8

TECHNIQUE



Only tiny amount of DNA template (target sequence) is required



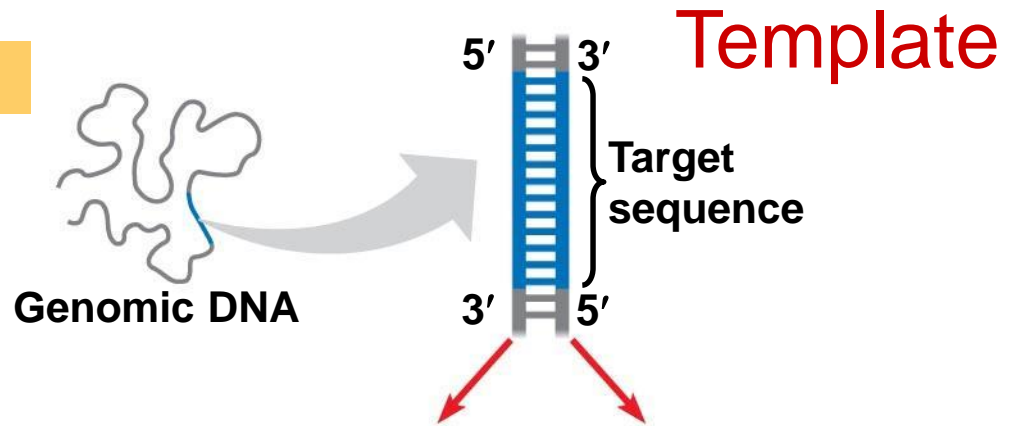
http://www.mun.ca/biology/scarr/PCR_simplified.html



Nobel prize in chemistry 1993
Kary B. Mullis
“for his invention of the polymerase chain reaction (PCR) method”

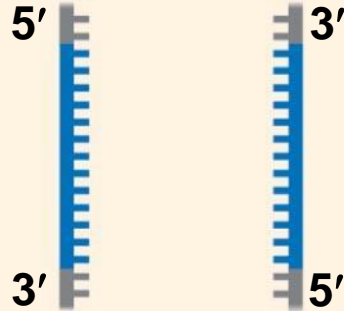
The polymerase chain reaction (PCR)

TECHNIQUE



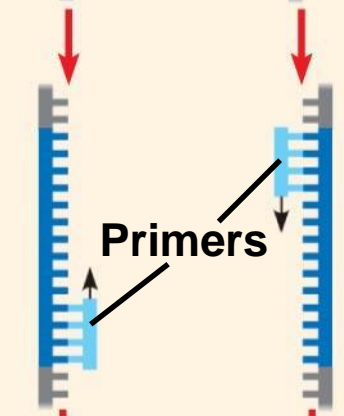
1 Denaturation

$\sim 94^{\circ}\text{C}$



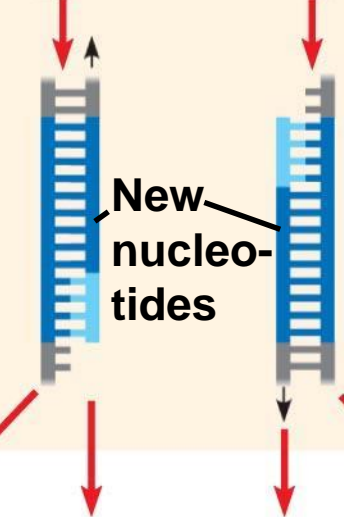
2 Annealing

$\sim 55^{\circ}\text{C}$

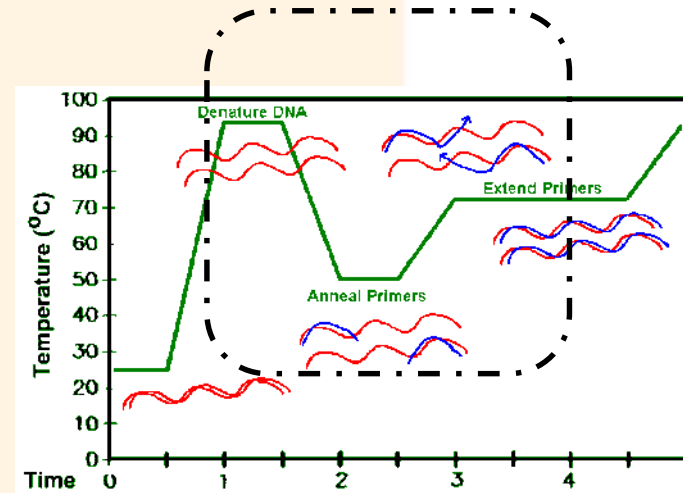


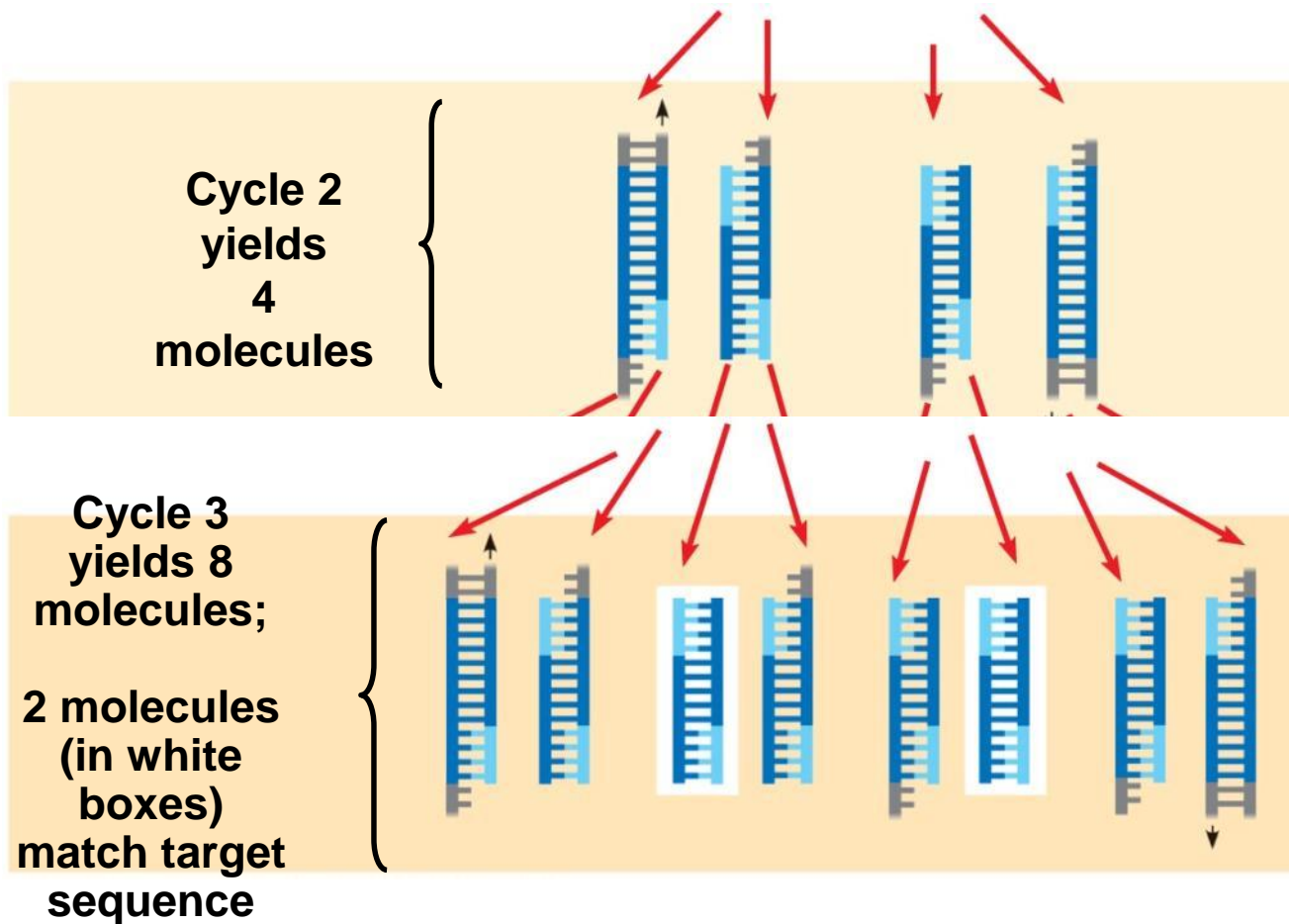
3 Extension

$\sim 72^{\circ}\text{C}$



Cycle 1
yields
2
molecules



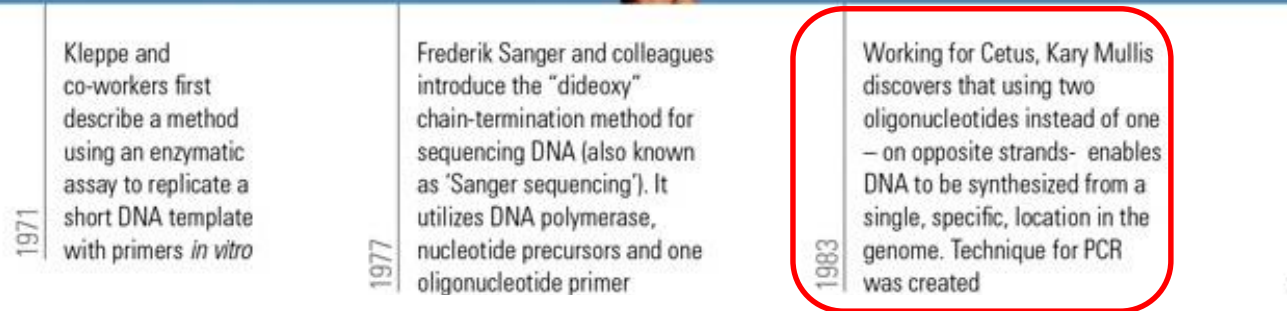
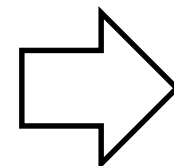
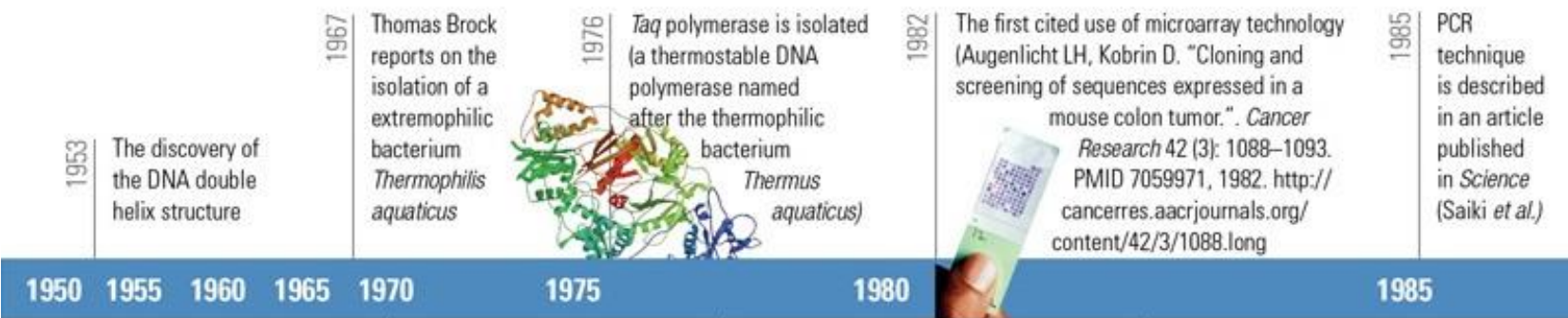


<u>Cycles</u>	<u>Copies</u>
1	2
2	4
4	16
10	1,024
15	32,768
20	1,048,576
25	33,554,432
30	1,073,741,824

The polymerase chain reaction (PCR)

to generate a large number of copies of specific DNA sequence

PCR through the ages



補充

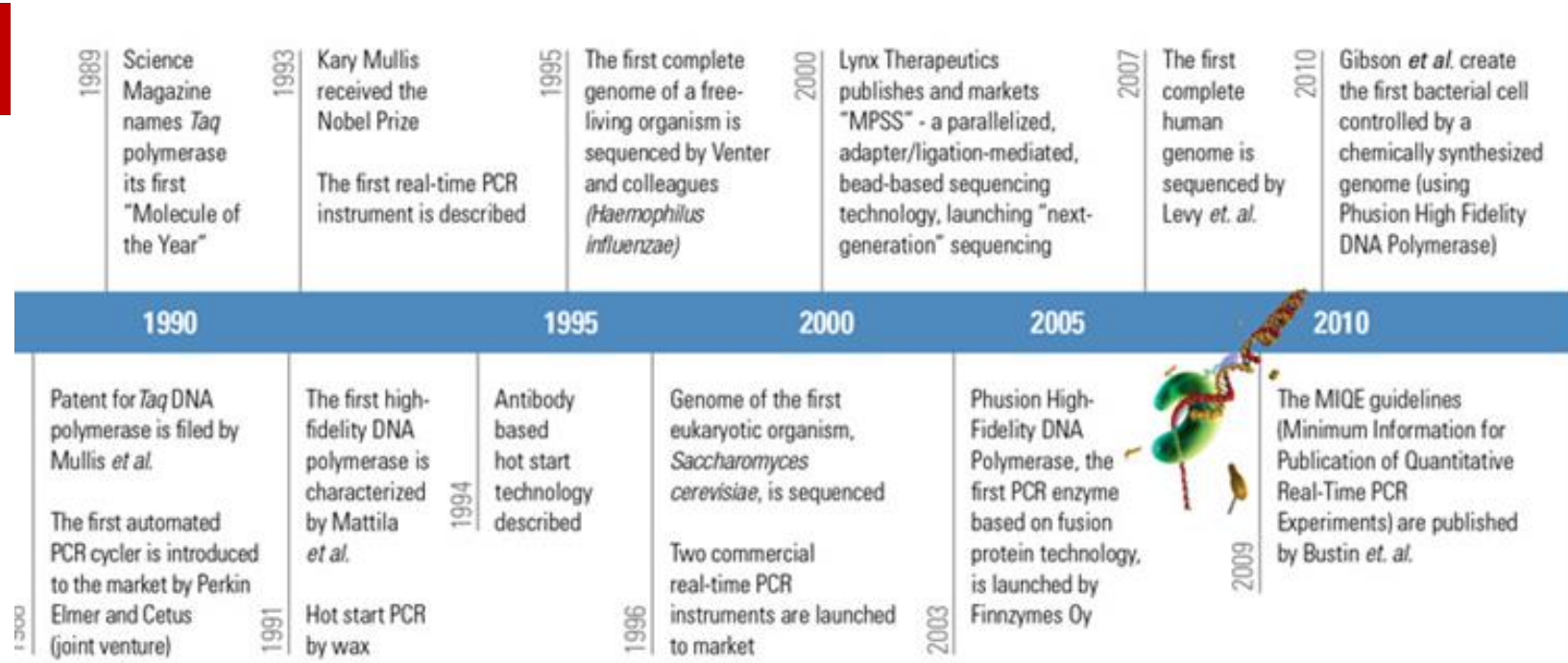
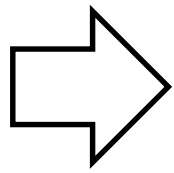
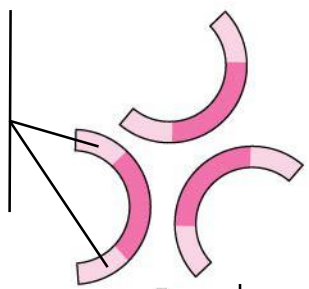


Figure 19.9

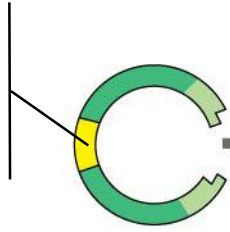
Gene cloning

DNA fragments obtained by PCR with restriction sites matching those in the cloning vector



Cut with same restriction enzyme used on cloning vector

A gene that makes bacterial cells resistant to an antibiotic is present on the plasmid.



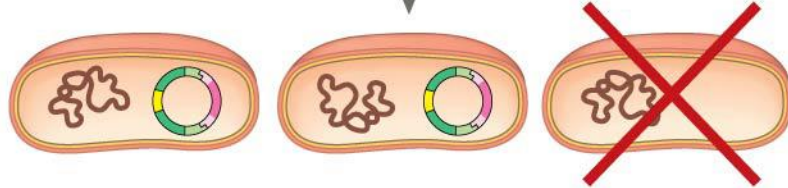
Cloning vector (bacterial plasmid)

Mix and ligate



Recombinant DNA plasmid (with acquired resistance to antibiotic)

(selection by antibiotic) Only cells that take up a plasmid will survive

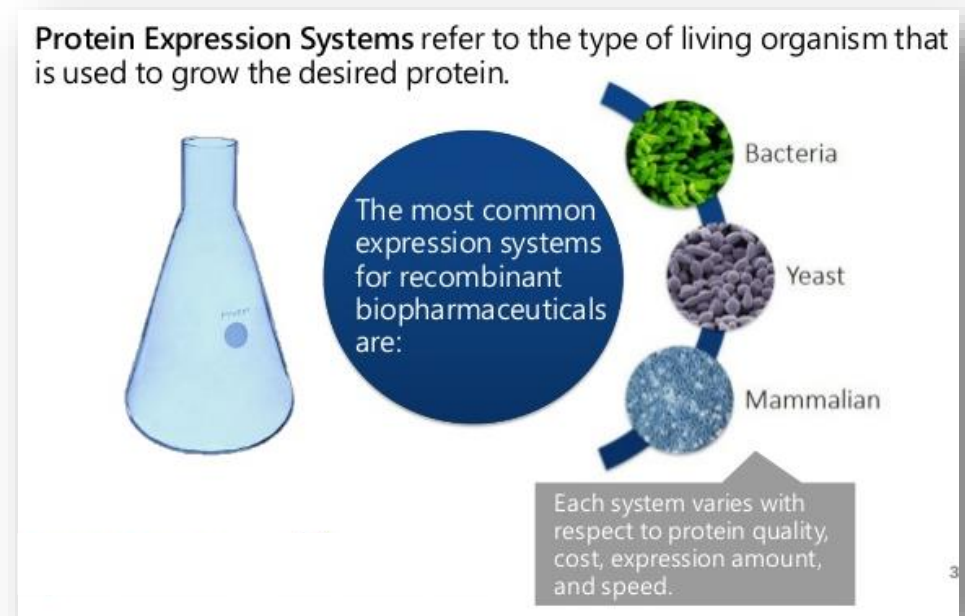


DNA technology allows us to study the sequence, expression, and function of a gene

- DNA cloning allows researchers to
 - **Compare genes and alleles between individuals** (in term of sequence)
 - Locate **gene expression** in a body
 - Determine **the role of a gene** in an organism
-

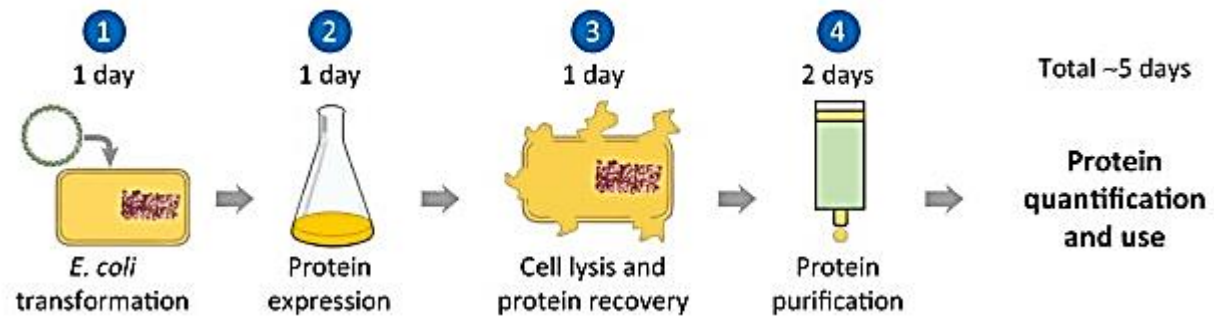
Expressing Cloned Eukaryotic Genes

- After a gene has been cloned, its **protein product** can be produced in larger amounts for research
- Cloned genes can be expressed as protein in either **bacterial** or **eukaryotic** cells



Bacterial Expression Systems

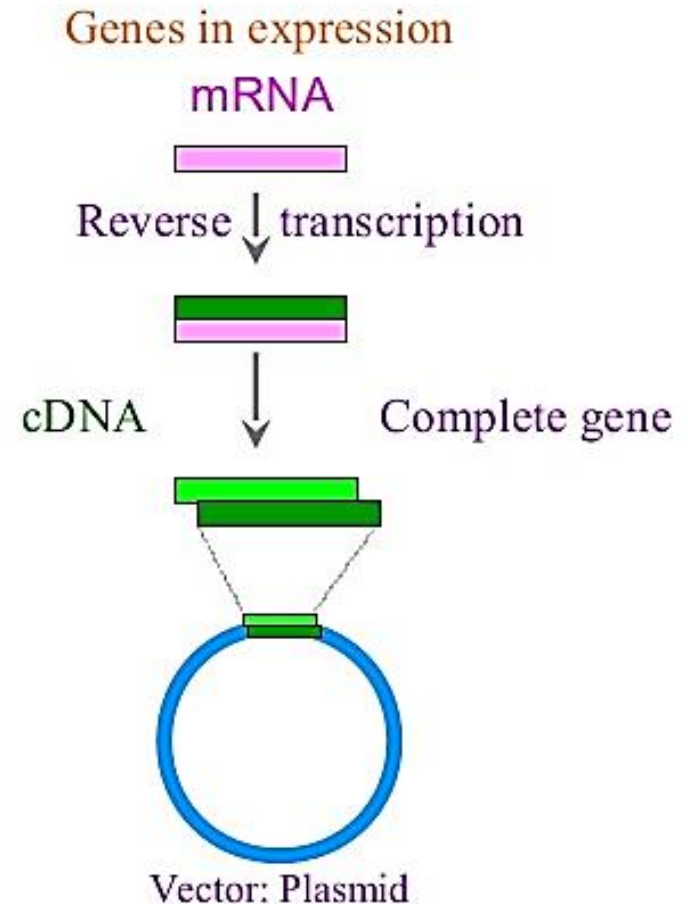
- Several **technical difficulties** hinder expression of cloned eukaryotic genes in bacterial host cells, such as no expression.
- To overcome **differences in promoters and other DNA control sequences**, scientists usually employ an **expression vector**, a cloning vector that contains a highly active bacterial promoter



Bacterial Expression Systems

- Most popular methods to deliver plasmids into *E.coli* – **heat shock** (42°C for 45 seconds)
 - **Technical difficulties** hinder expression of cloned eukaryotic genes in bacterial host cells: **no expression, not soluble, not proper folded,** etc.
-

- Another difficulty with eukaryotic gene expression in bacteria is the presence of introns in most eukaryotic genes
- Researchers can avoid this problem by using **cDNA**, **complementary to the mRNA**, which contains only exons



Read more on RT-PCR

Eukaryotic DNA Cloning and Expression Systems

- Molecular biologists can avoid **eukaryote-bacterial incompatibility issues** by using eukaryotic cells, such as **yeasts, as hosts for cloning and expressing genes**
- Even yeasts may not possess the proteins required to modify expressed mammalian proteins properly; which means **the post-translational modifications** that many proteins require, i.e. **glyco-/lipo-proteins**
- In such cases, **cultured mammalian or insect cells** may be used to express and study proteins

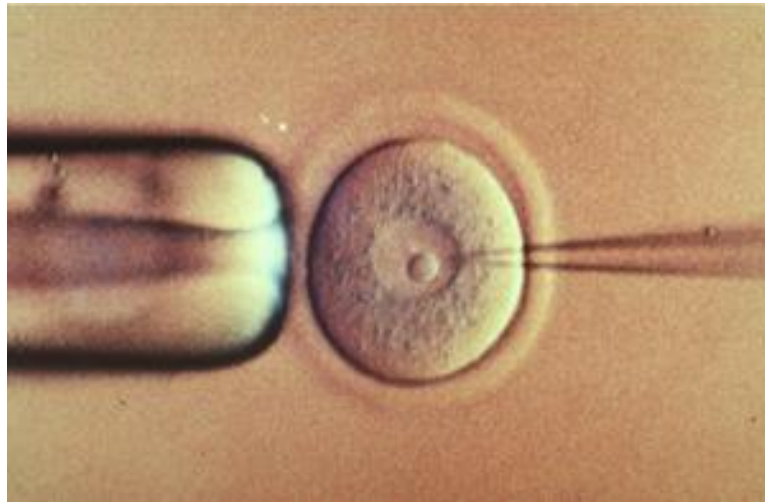
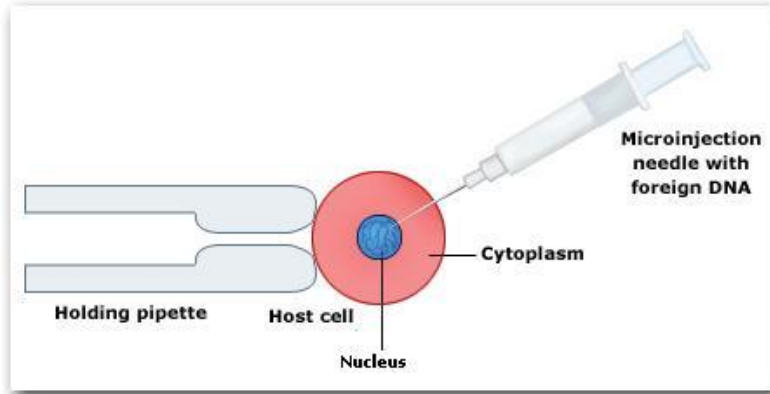
Delivery gene/plasmid into the eukaryotic cells

- One method of introducing recombinant DNA into eukaryotic cells is **electroporation**, applying a **brief electrical pulse** to create **temporary holes** in **plasma membranes**



- Alternatively, scientists can **inject DNA** into cells using microscopically thin needles (**microinjection**)
 - Once inside the cell, the DNA is incorporated into the cell's DNA by natural genetic recombination
-

Microinjection 顯微注射 for DNA Transfer



DNA microinjection

Microinjector



Cross-Species Gene Expression and Evolutionary Ancestry

- The remarkable ability of bacteria to express some eukaryotic proteins underscores the shared evolutionary ancestry of living species
- For example, *Pax-6* is a gene that directs formation of a **vertebrate eye**; the same gene in flies directs the formation of an **insect eye** (which is quite different from the vertebrate eye)
- **The *Pax-6* genes in flies and vertebrates can substitute for each other**

Analyzing Gene Expression



-  ***In situ* hybridization**  uses fluorescent dyes attached to probes to identify the location of specific mRNAs in place in the intact organism
 - *In situ* 在原位 hybridization (ISH) 偵測完整個體上標的基因的真實表現狀況
- Probes can be used to identify **where** **or when** a gene is transcribed in an organism
-

Figure 19.10a

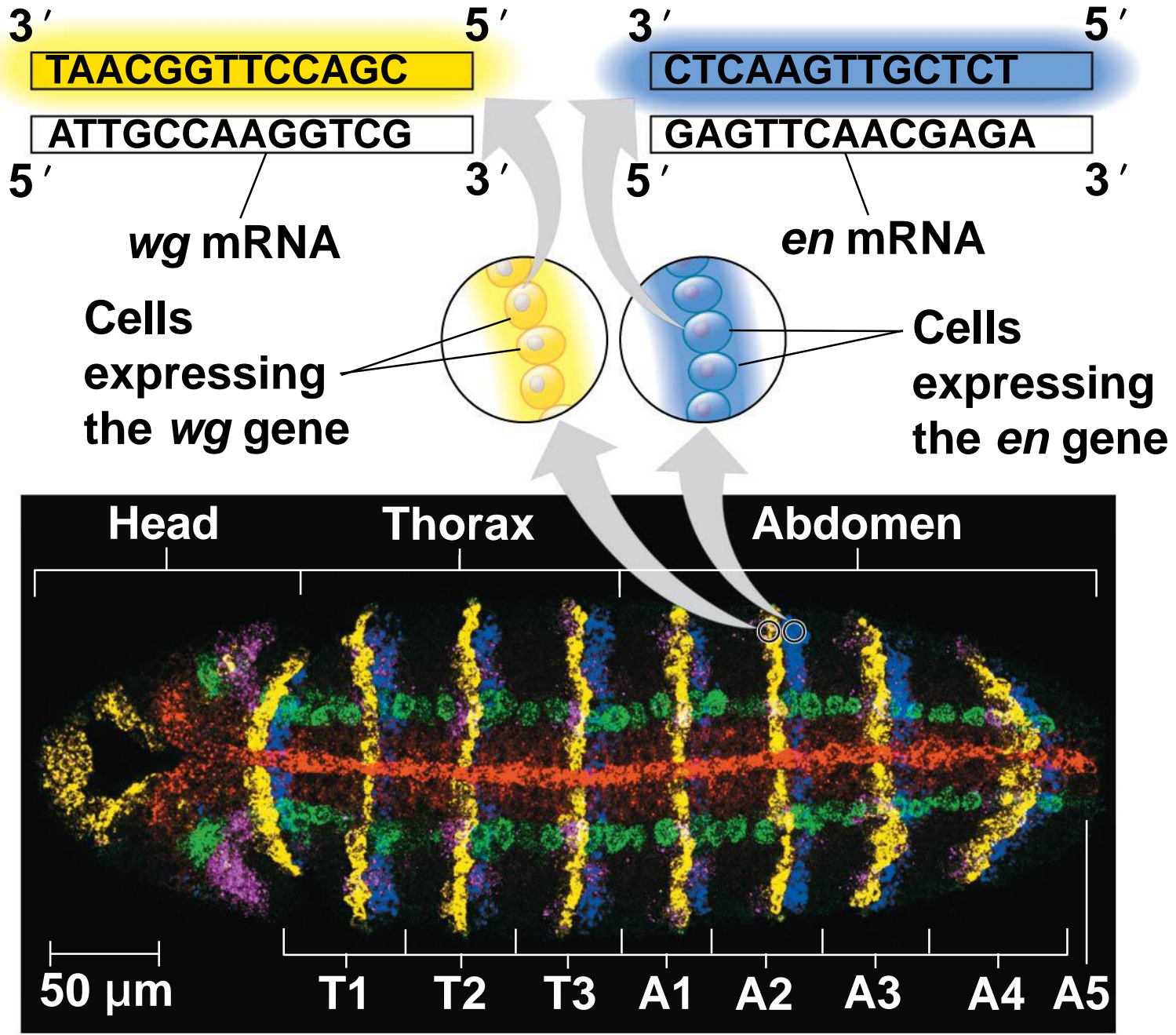
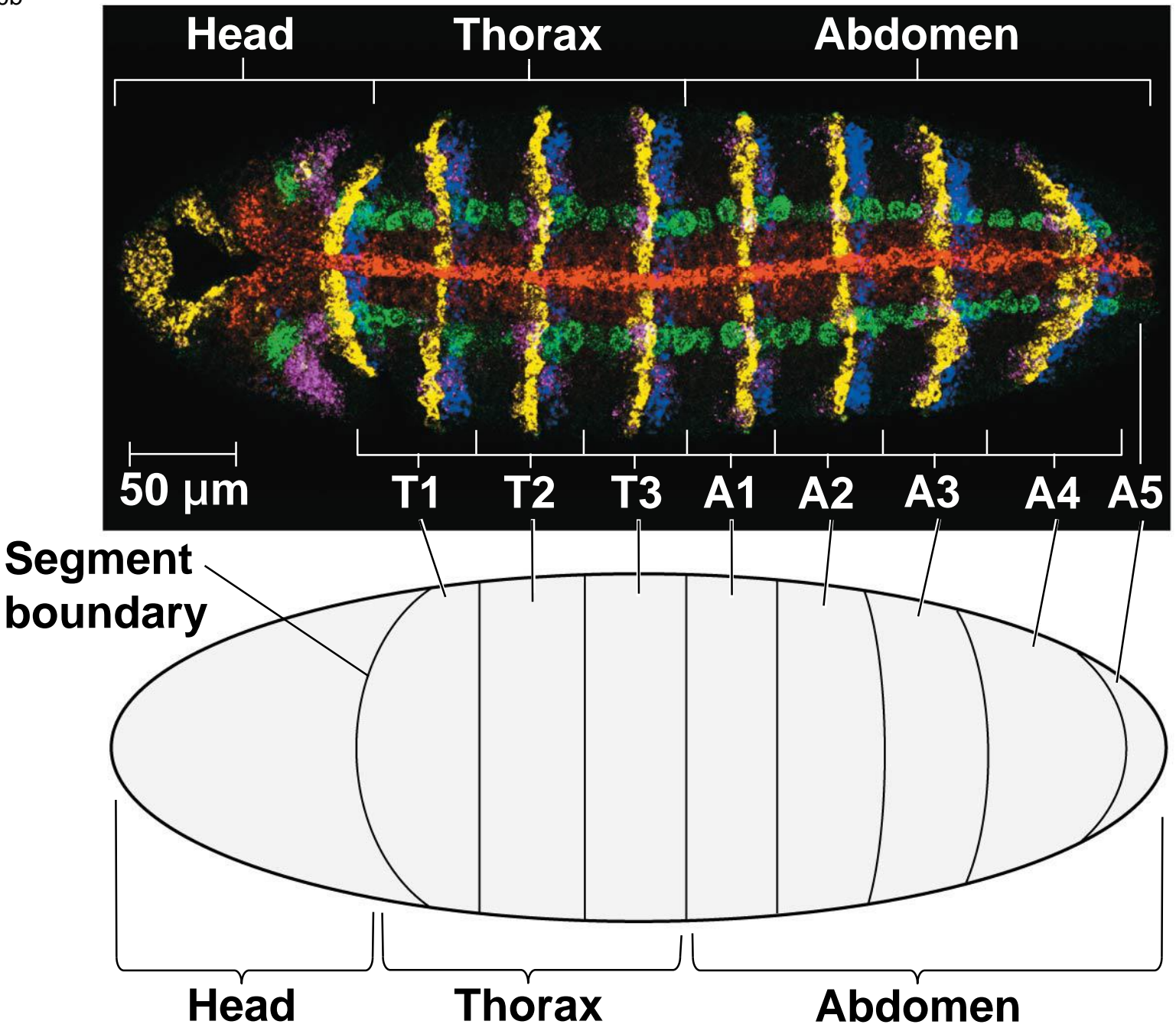


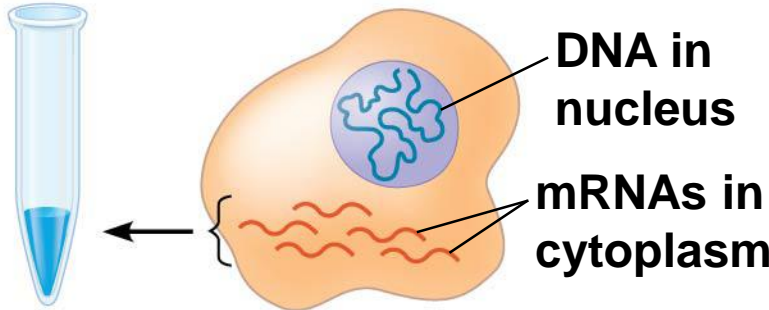
Figure 19.10b



Reverse transcriptase-polymerase chain reaction (RT-PCR)

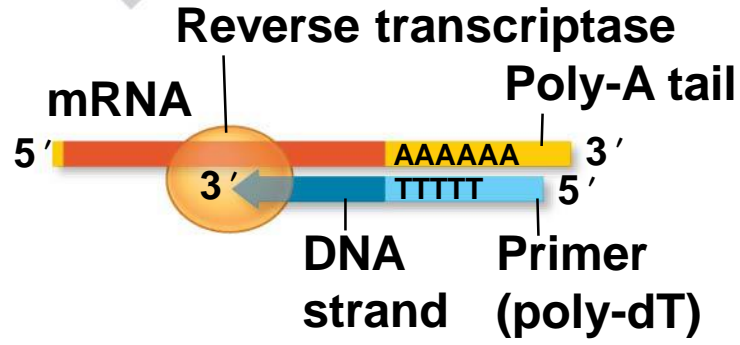
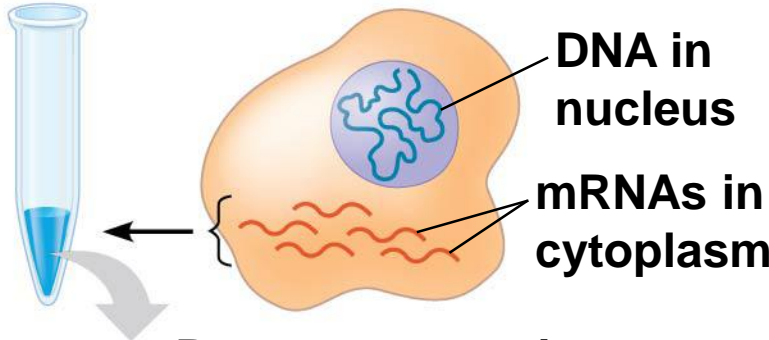
- **Reverse transcriptase-polymerase chain reaction (RT-PCR)** is useful for comparing **amounts of specific mRNAs** in several samples at the same time
- Reverse transcriptase is **added to mRNA to make complementary DNA (cDNA)**, which serves as a template for PCR amplification of the gene of interest
- The products are run on a gel and the mRNA of interest is identified

Figure 19.11-1



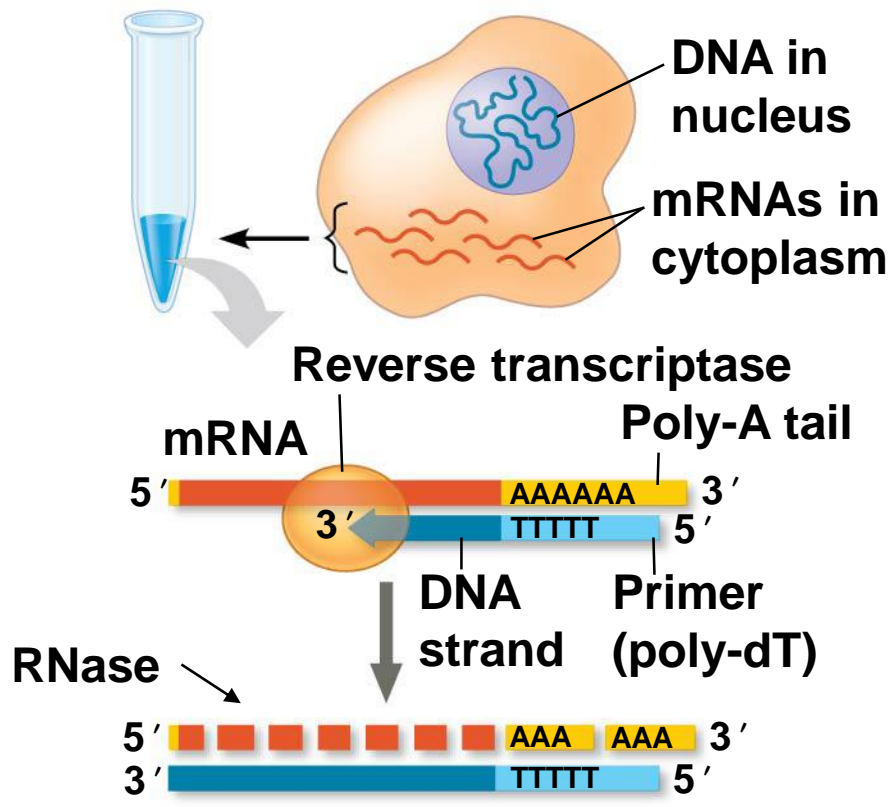
Making complementary DNA (cDNA)

Figure 19.11-2



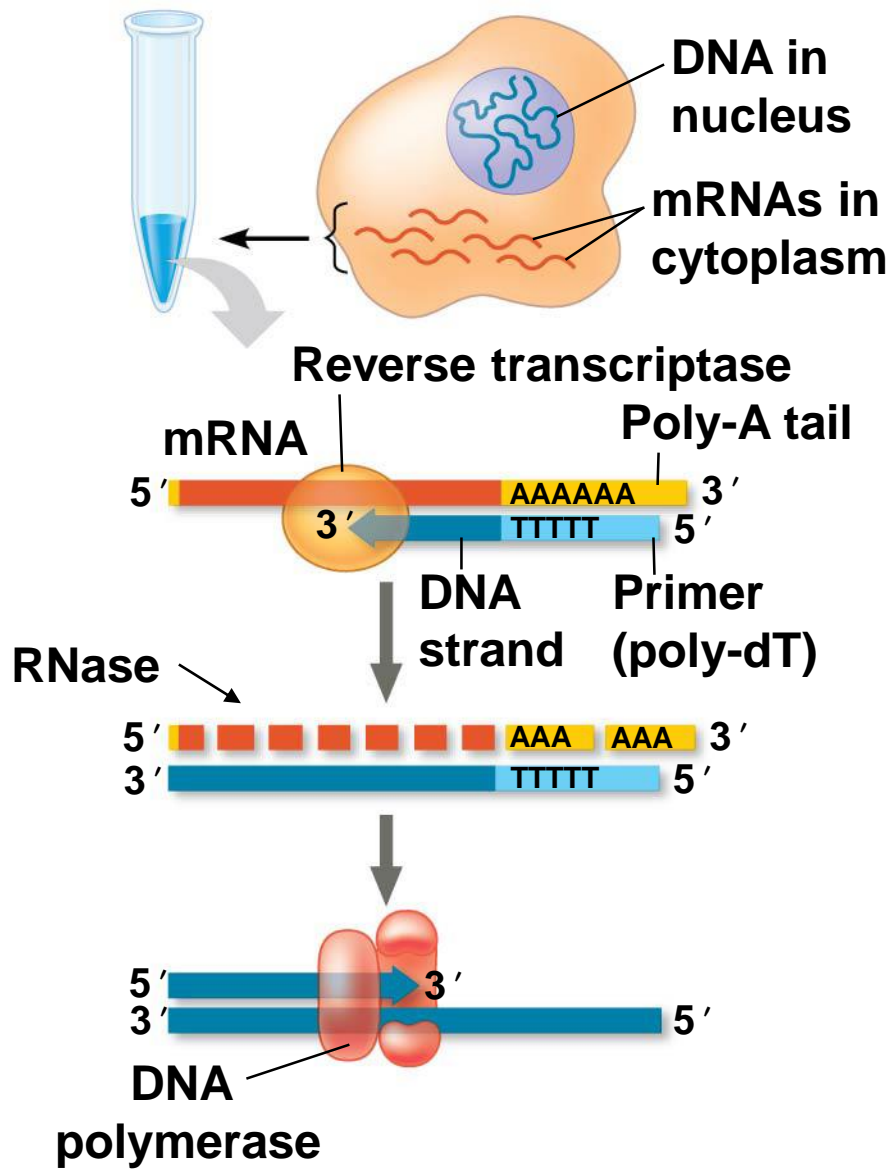
Making complementary DNA (cDNA)

Figure 19.11-3



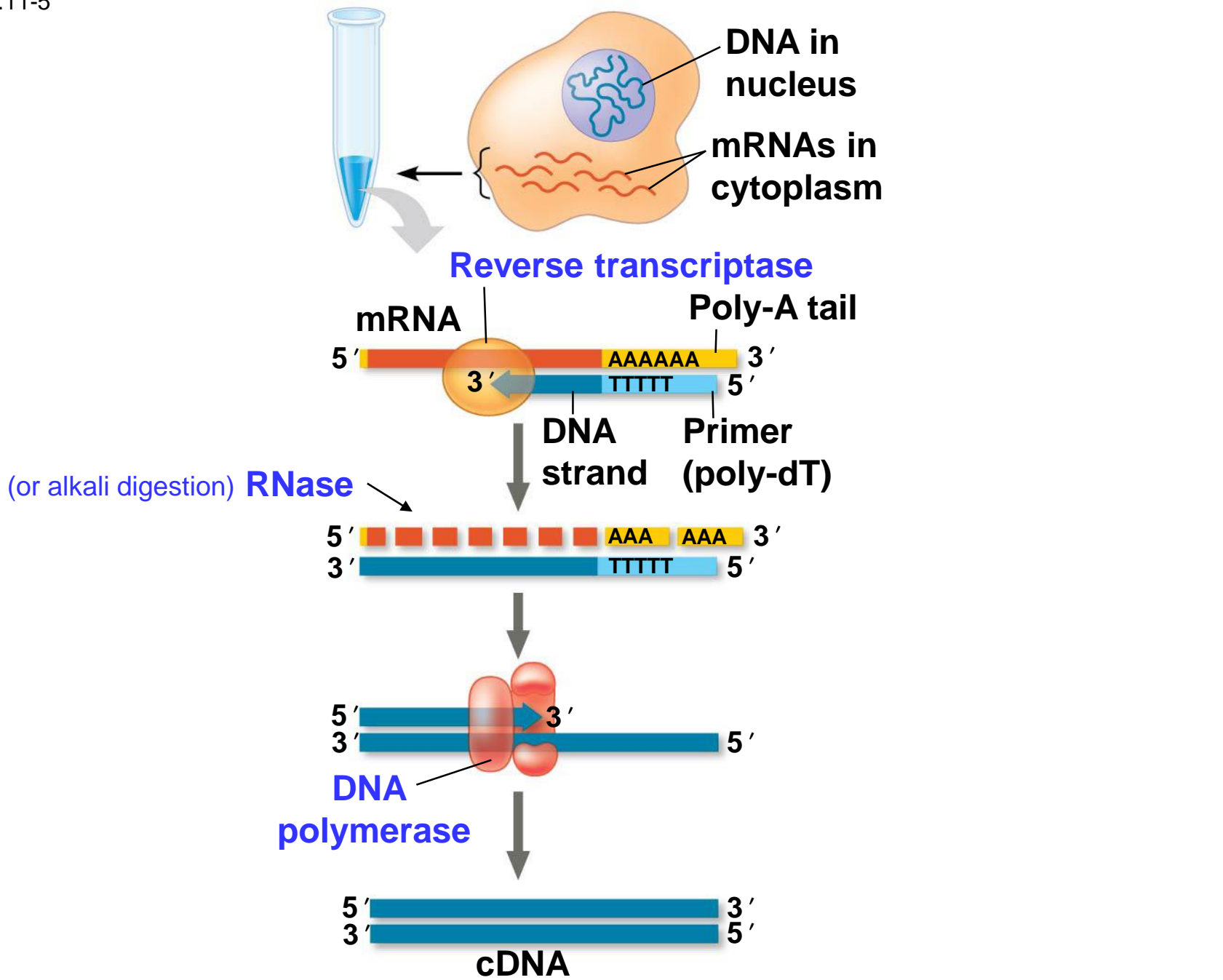
Making complementary DNA (cDNA)

Figure 19.11-4



Making complementary DNA (cDNA)

Figure 19.11-5



Making complementary DNA (cDNA)

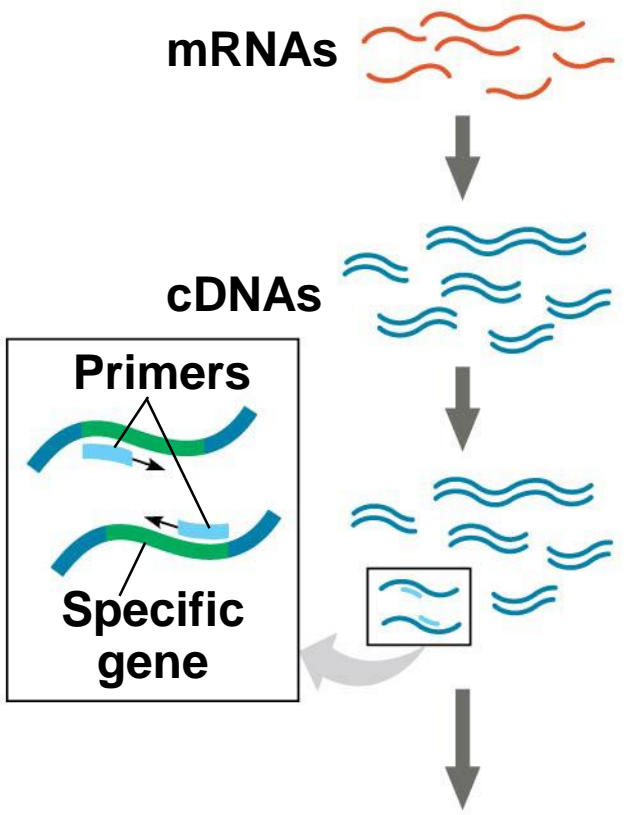
Figure 19.12

Technique

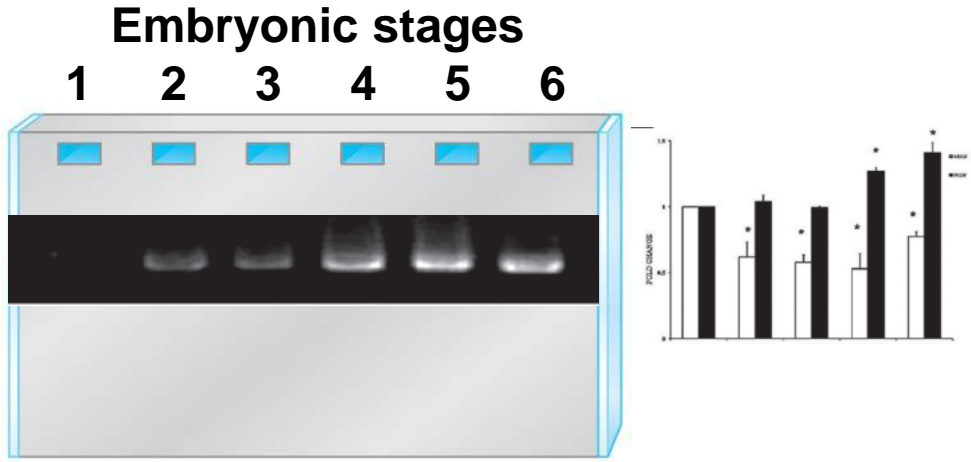
1 cDNA synthesis

2 PCR amplification

3 Gel electrophoresis



Results



RT-PCR analysis of the expression of genes

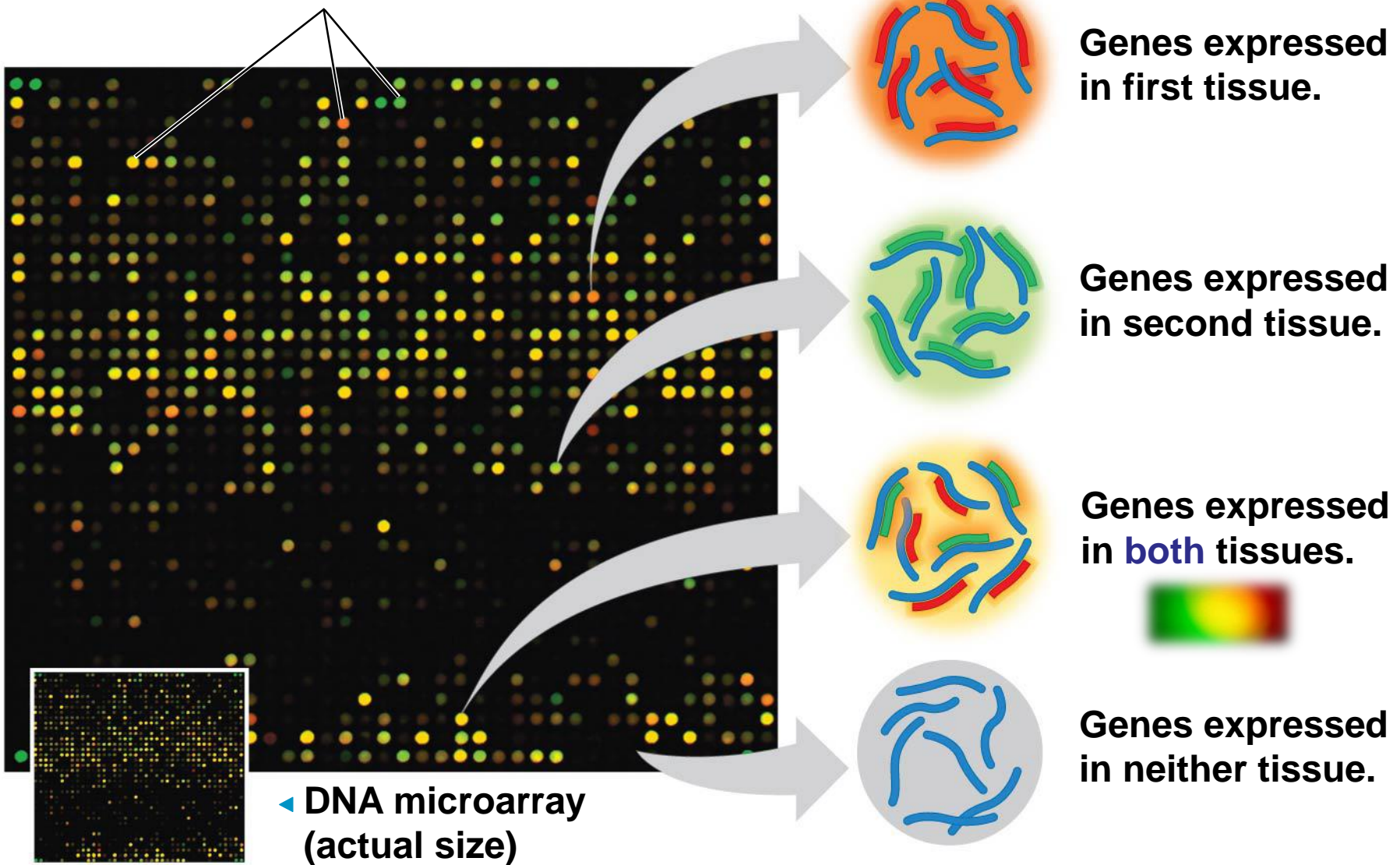
Studying the Expression of Interacting Groups of Genes

- **Automation** 自動化 has allowed scientists to measure the expression of thousands of genes at one time using DNA microarray assays
- **DNA microarray assays** 微陣列檢定 compare patterns of gene expression in different tissues, at different times, or under different conditions



Figure 19.13

Each dot is a well containing identical copies of DNA fragments that carry a specific gene.



Protocol next page

A typical DNA microarray co-hybridization (2 dye) experiment

Control Sample

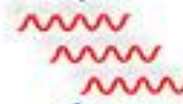
Experimental Sample



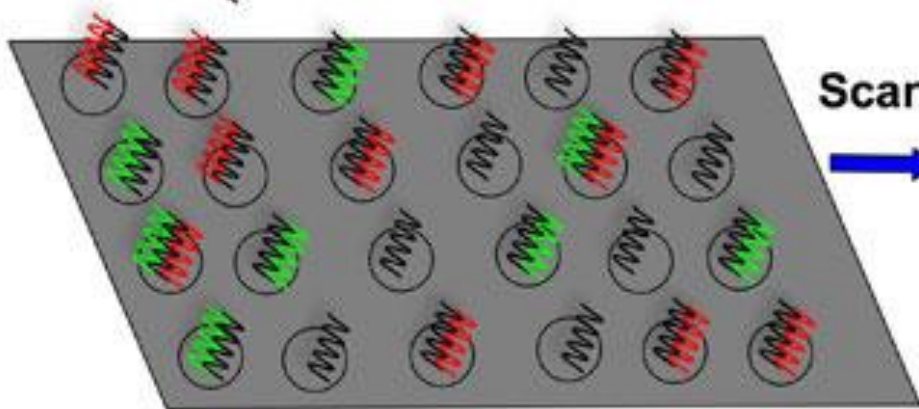
mRNA extraction



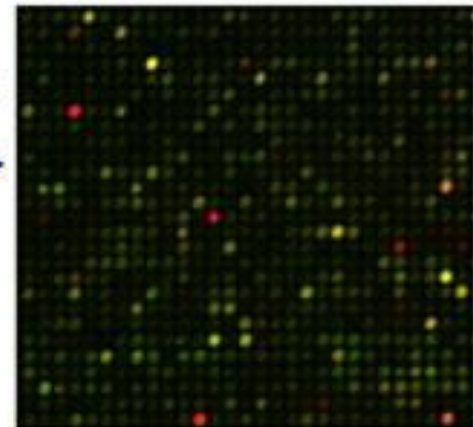
Reverse Transcription,
fluorescent labeling



Combine equal amounts
and hybridize



Scan

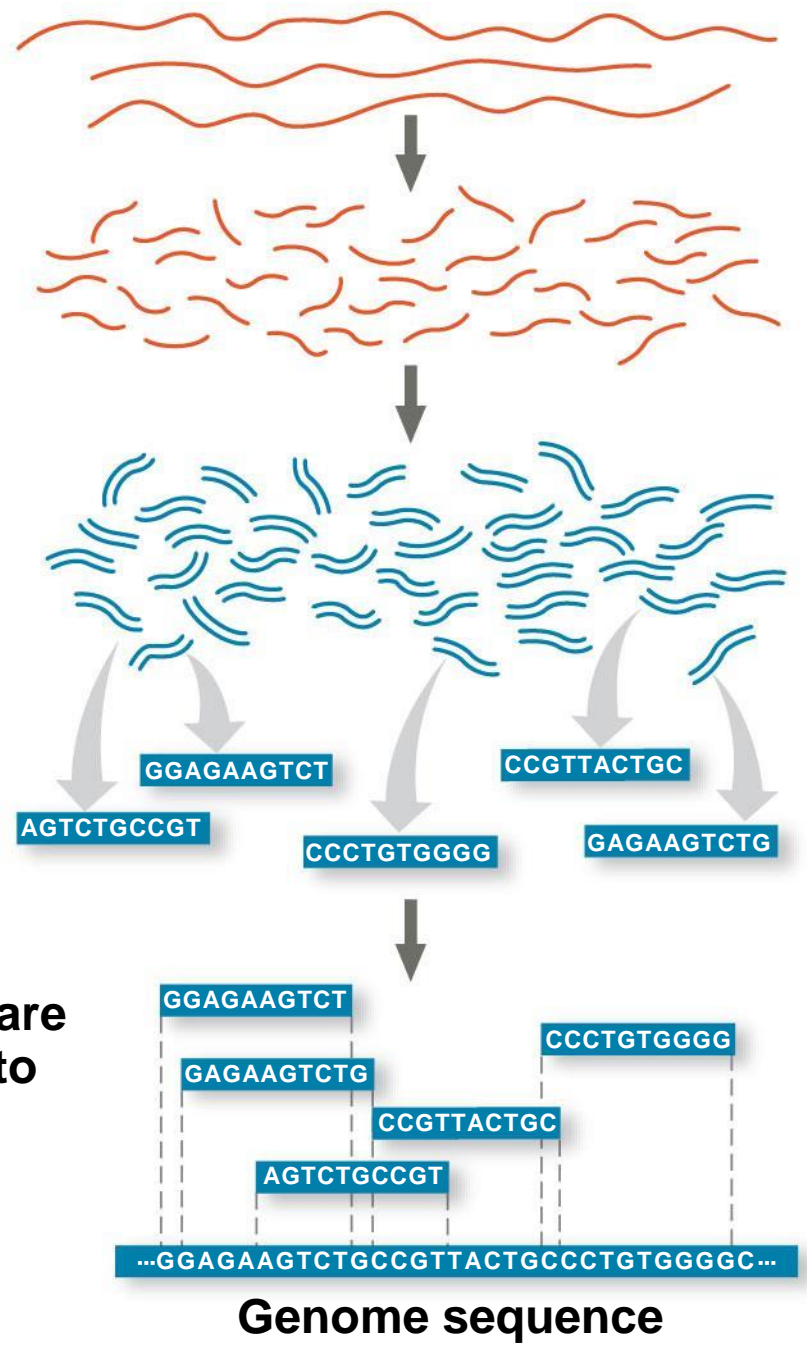


Use of **RNA sequencing** (RNA-Seq) to analyze expression of many genes

- With rapid and inexpensive sequencing methods, researchers can now just **sequence cDNA samples from different tissues** or embryonic stages to determine the **gene expression differences** between them
- This approach is called **RNA sequencing** or **RNA-seq**
- mRNAs are isolated, cut into short, similar-sized fragments, converted into cDNAs, and sequenced

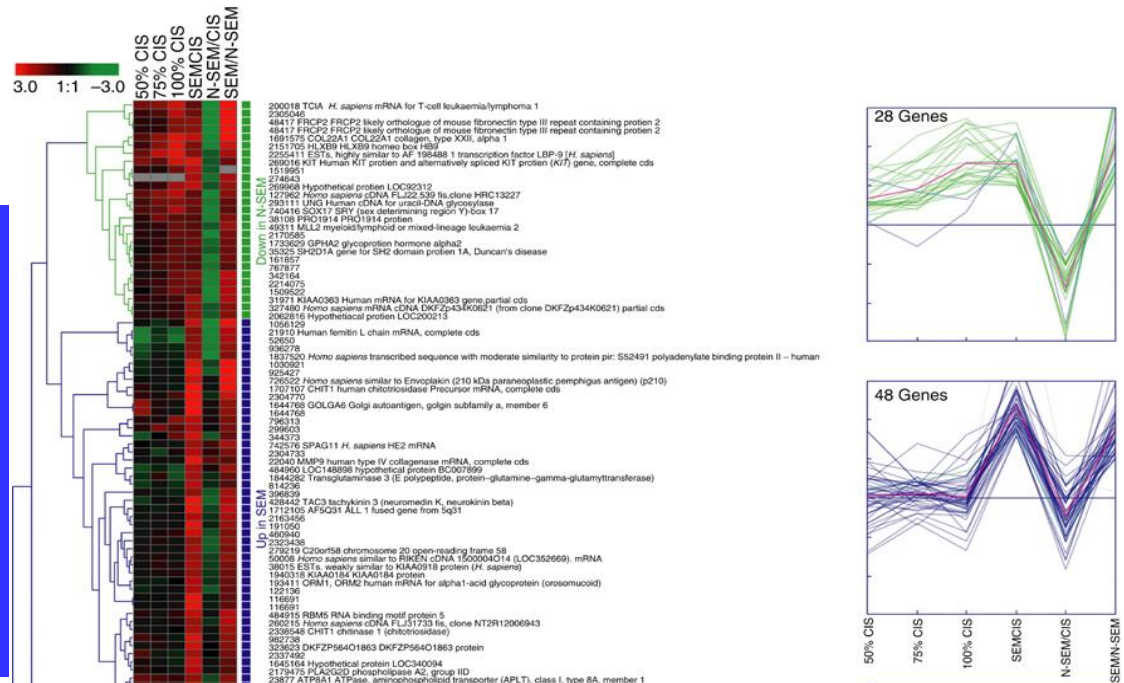
Figure 19.13

- 1 mRNAs are isolated from the tissue being studied.
- 2 mRNAs are cut into similar-sized, small fragments.
- 3 mRNAs are **reverse-transcribed** into cDNAs of the same size.
- 4 cDNAs are sequenced.
- 5 The short sequences are mapped by computer onto the genome sequence.



- Technique like RNA-seq enables researchers to determine the gene **expression differences** between different tissues or embryonic stages
- By uncovering **gene interactions** and **clues to gene function**, gene expression assays may contribute to understanding of disease and suggest new diagnostic targets

Gene interaction -
 Several different genes collaborate in the production of one phenotypic character (or related group of characters).



Example: Determining Gene Function

- One way to determine function is to **disable the gene** and observe the consequences – **loss-of-function** (vs. **gain-of-function**)
- How? Using ***in vitro* mutagenesis**,
 - mutations are introduced into a cloned gene, **altering or destroying its function**
 - when the **mutated gene** is returned to the cell, the normal gene's function might be determined by examining the **mutant's phenotype**

Wild type

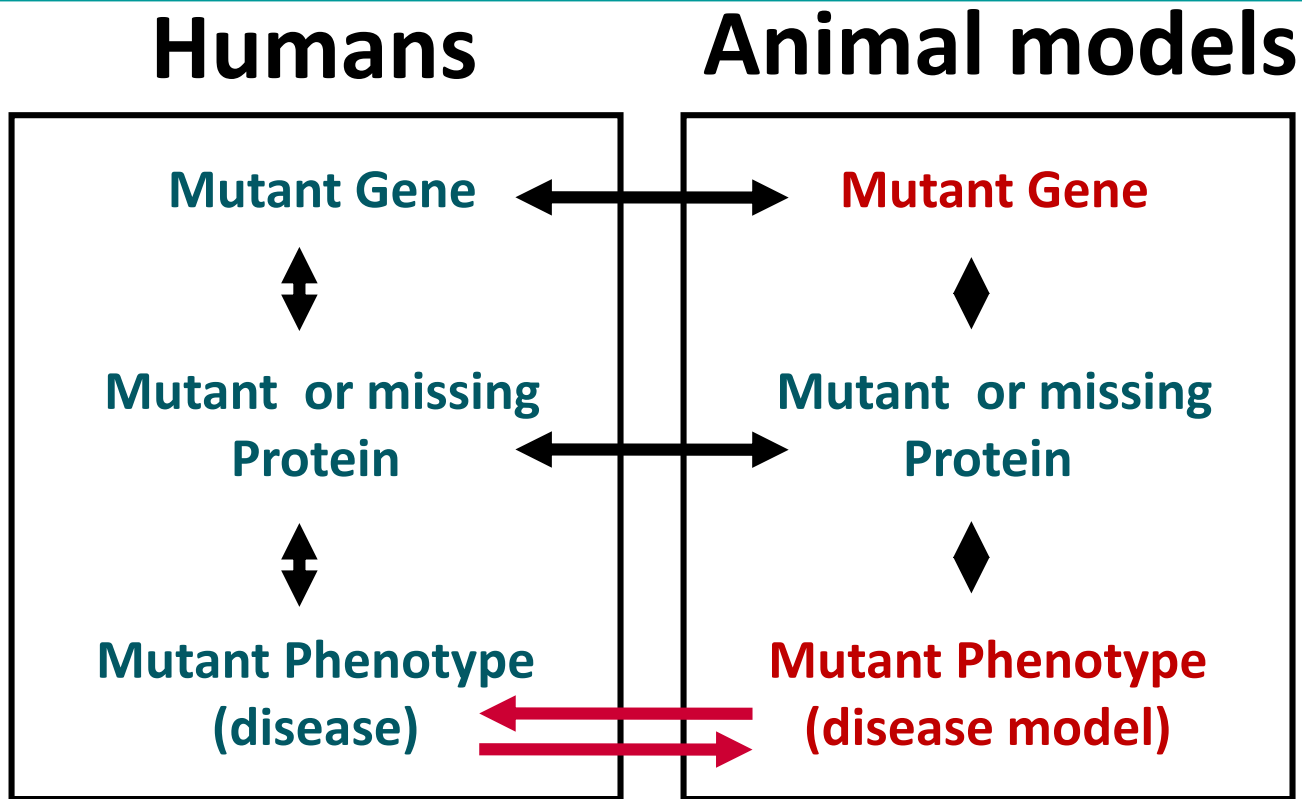
NACRE^{-/-}

ROY^{-/-}

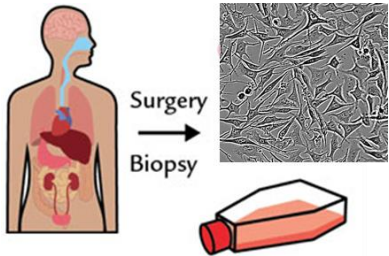
ROY^{-/-} ;NACRE^{-/-}



Animal models of human diseases



Cell lines of melanoma



Mouse model of melanoma



Zebrafish model of melanoma



Altering or Editing Genetic materials

- Using ***in vitro* mutagenesis** 誘發突變, specific mutations are introduced into a cloned gene, altering or destroying its function
- When the mutated gene is returned to the cell, the normal gene's function might be determined by examining the mutant's phenotype

in vitro: 試管內，體外

in vivo: 活體內

In situ: 原位，正位

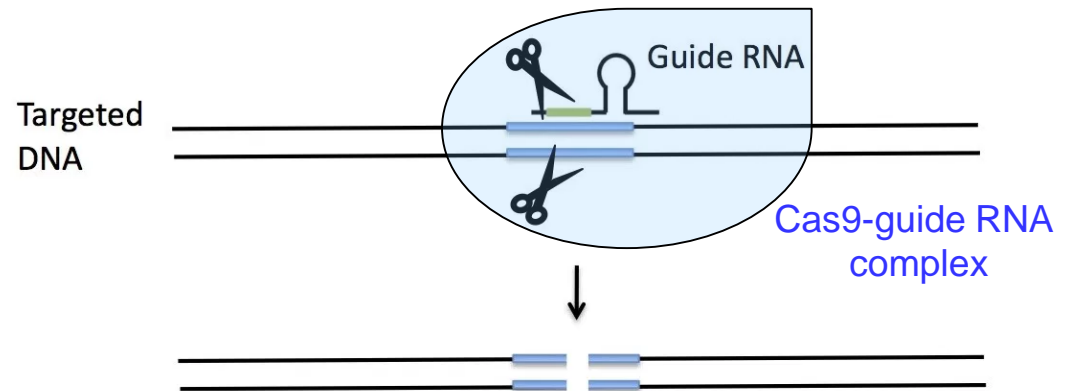
ex vivo: 生物活體之外

In utero: 在子宮裡

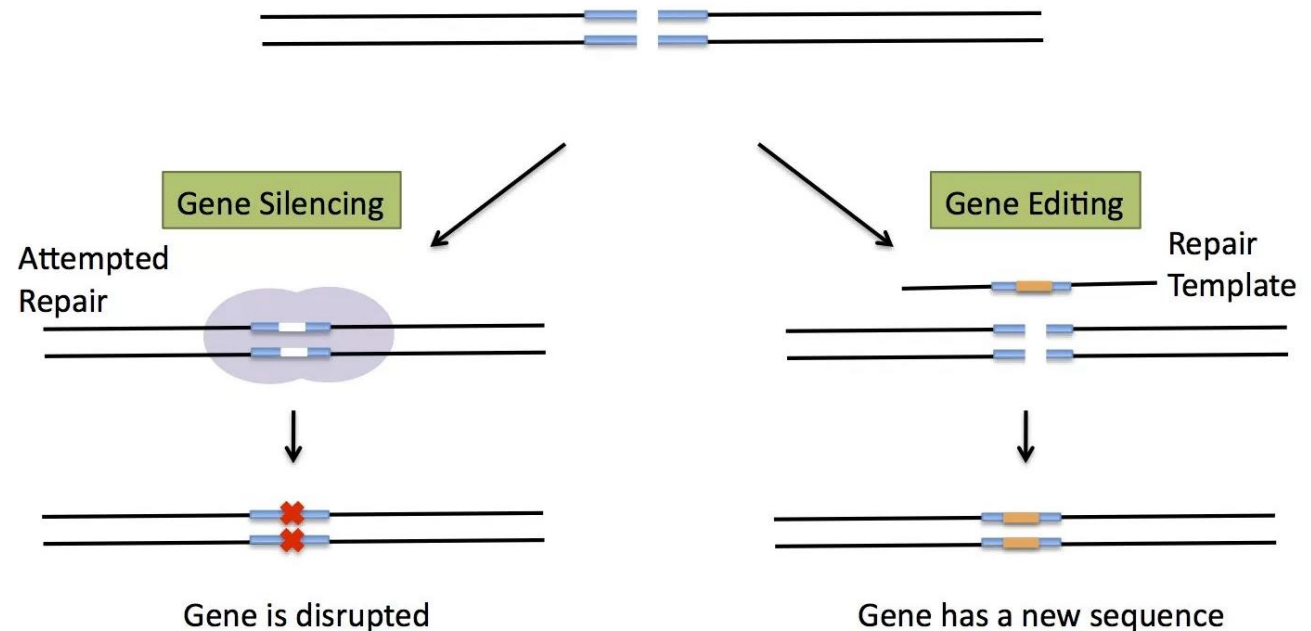
In silico: 經由電腦模擬

Genome editing 基因編輯

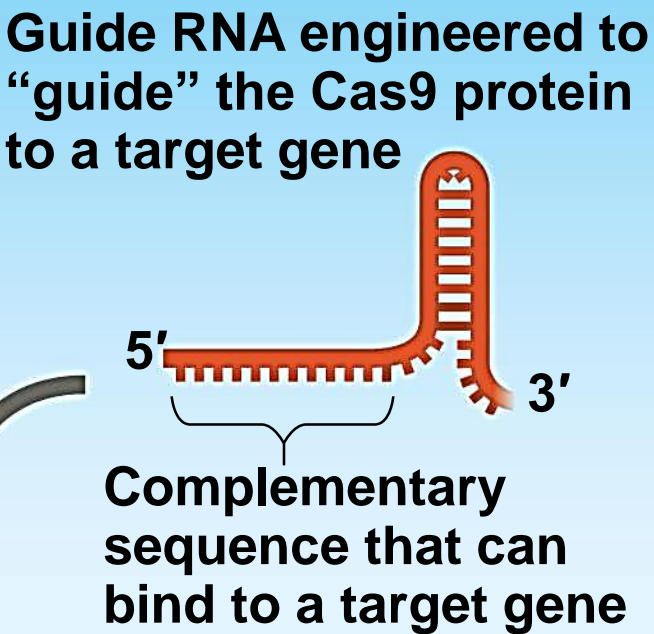
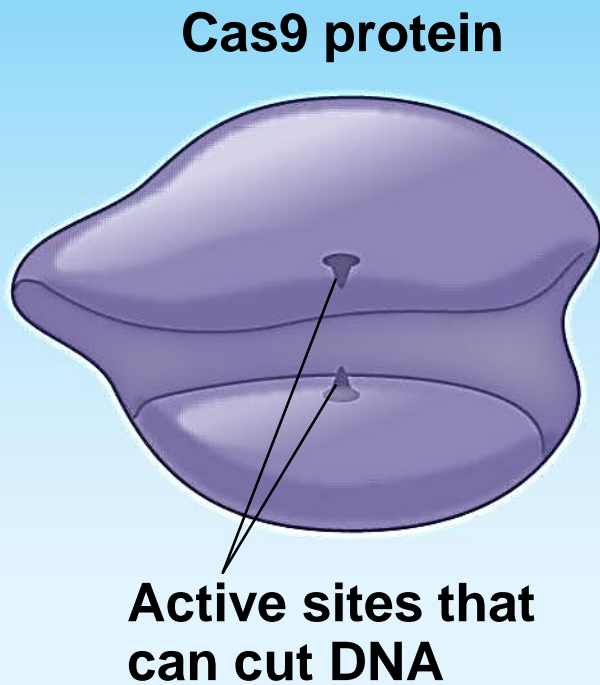
- **Bacterial defense against bacteriophage**
 - The **CRISPR-Cas9 system** is a powerful new technique for gene editing in living cells and organisms
 - **Cas9** acts together with a “**guide RNA**” made from the **CRISPR region** of the bacterial system; the **Cas9-guide RNA complex** will bind and cut both strands of any DNA sequence complementary to the guide RNA
- If the guide RNA is **engineered** to be complementary to the target gene, the target DNA will be cut



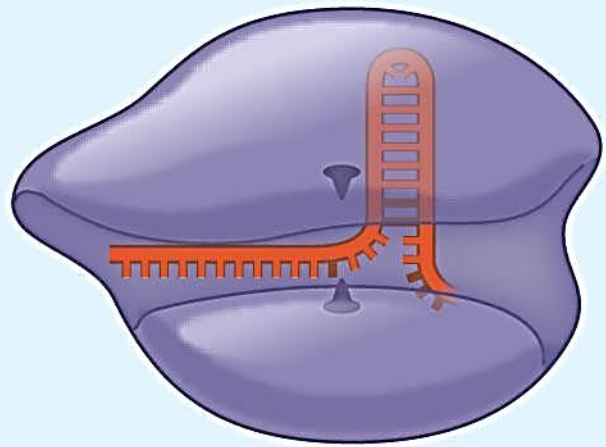
- When the cut DNA is **repaired**, nucleotides may be introduced or removed, causing the gene to be **inactivated**
- Researchers have also modified the technique so **a gene with a mutation in it can be repaired**
 - This is done by introducing **a segment of the wild-type gene**, which may be used as a template to repair the target DNA (i.e. **gene therapy**)



Gene editing using the CRISPR-Cas9 system



Cas9-guide RNA complex



1

Figure 19.14b

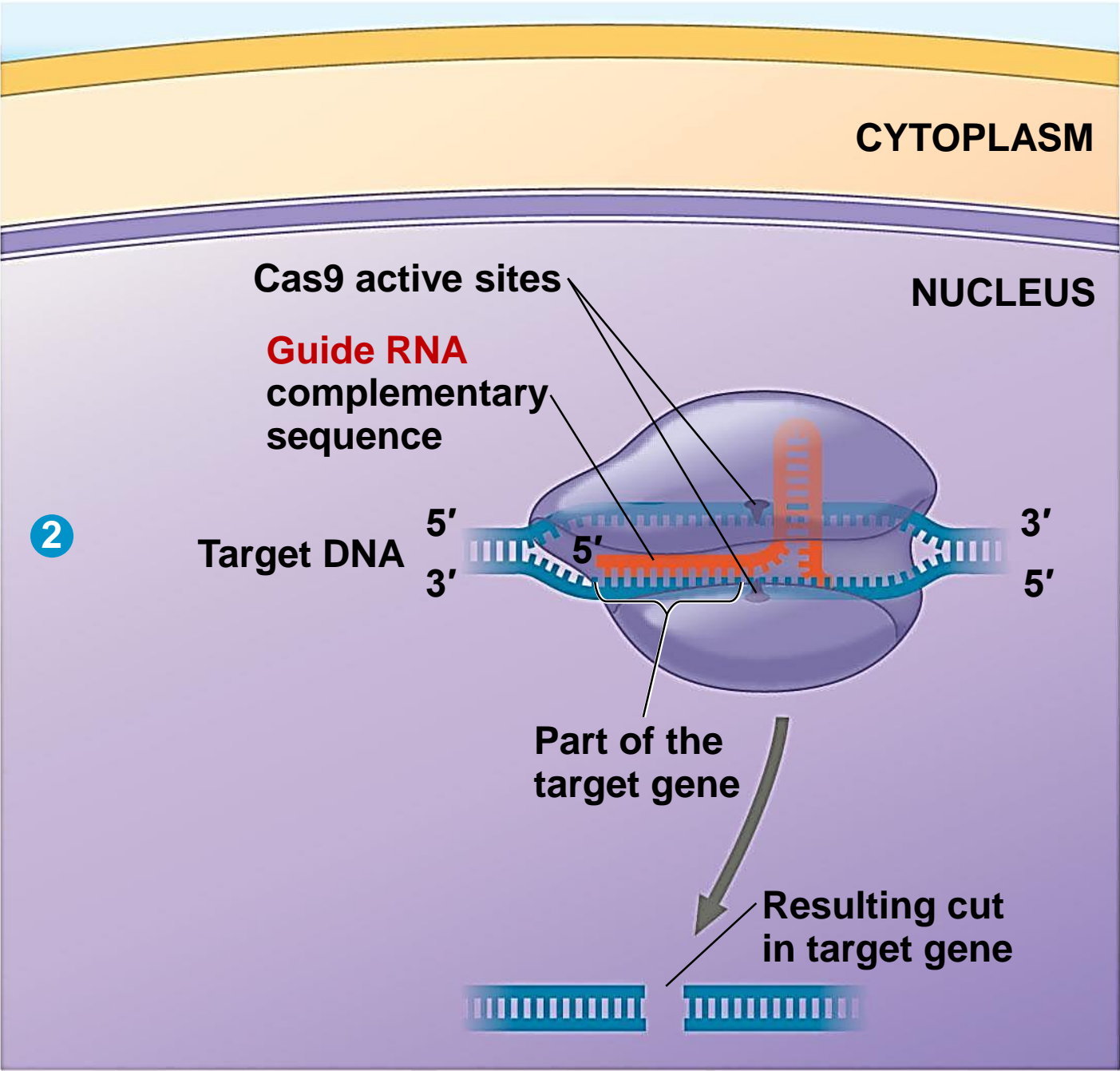
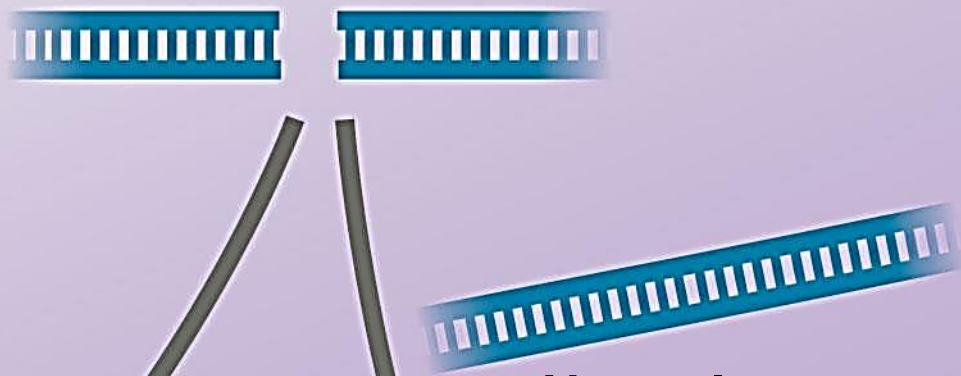


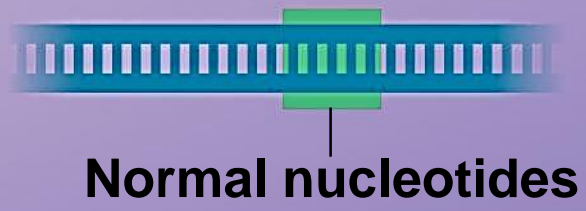
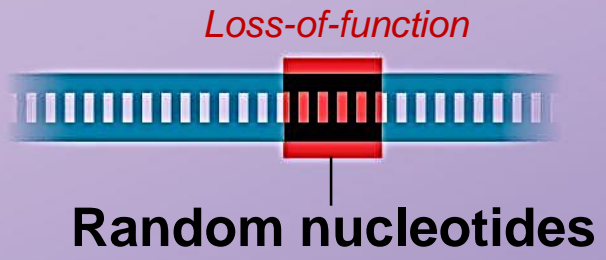
Figure 19.14c

3



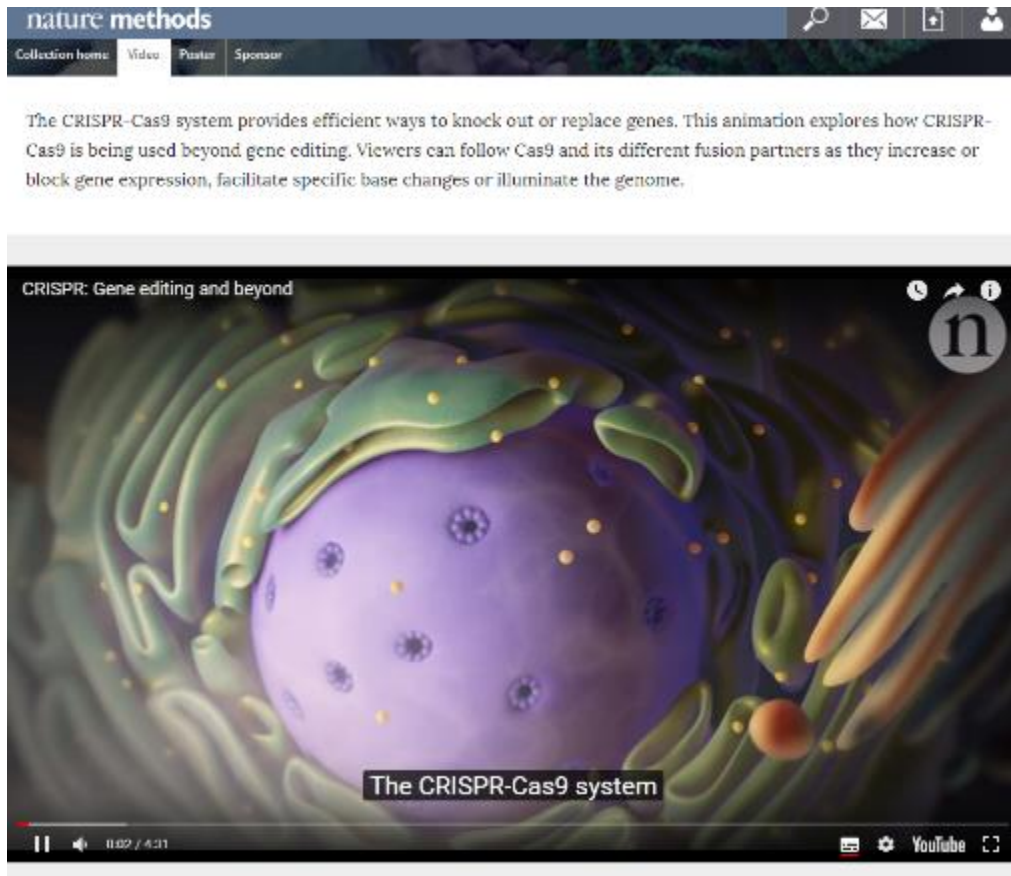
(a) Scientists can disable (“knock out”) the target gene to study its normal function.

(b) If the target gene has a mutation, it can be repaired.



補充: Expanding the CRISPR Toolbox

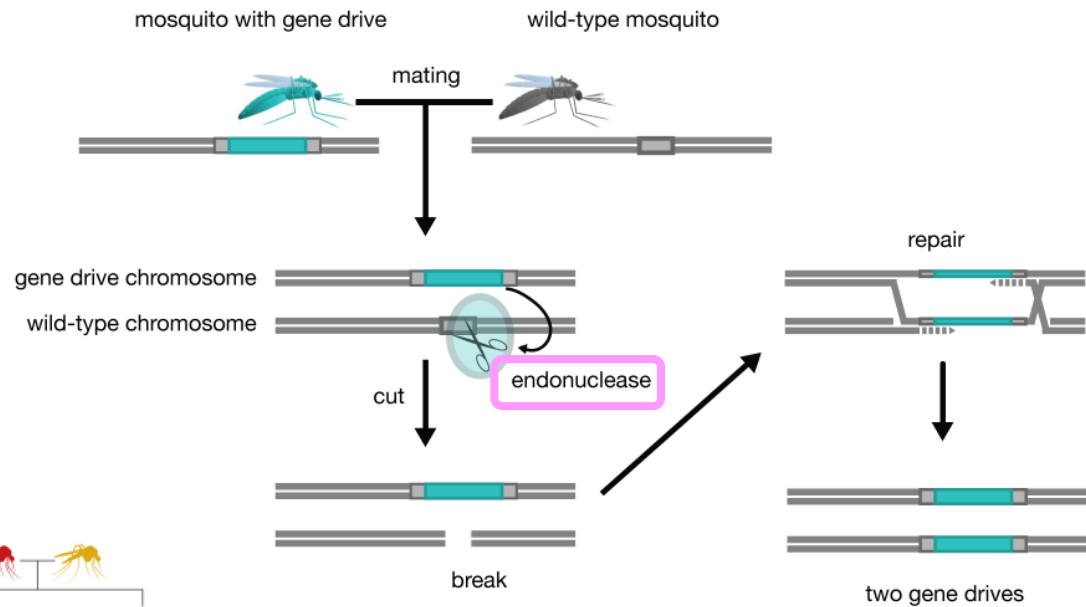
Animation on **CRISPR gene editing and beyond**
<https://www.nature.com/collections/txhdfs1xzh/videos>



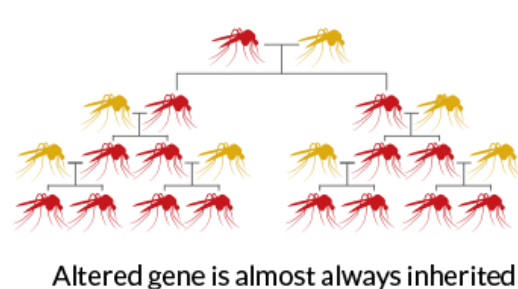
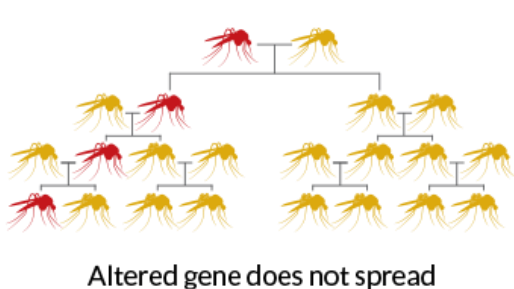
The CRISPR-Cas9 system is best known for its ability to knock out or replace specific genes, via targeted cleavage of the genome. But scientists are developing **many more applications**, typically by using an inactive Cas9 to target other enzymes to specific genomic sites. From transcriptional regulation to base editing, these developments are extending the range of biological questions that can be probed with CRISPR/Cas9.

Example Application of Genome Editing

- Alternating genes in insect to stop insect-borne disease (i.e. Mosquito)
- Engineering **new allele that is favored for inheritance** (i.e. **Gene drive 基因驅動** to introduce new allele through the population)



「刺激有遺傳偏差的基因去影響整個族群」的做法就稱作「基因驅動/gene drive」



- 2003 年 Imperial College London 的演化生物學家 Austin Burt，想到利用 **HEG** (**homing endonuclease gene**) 這個自私基因來改變整個族群的基因。HEG 的 H 指的是：「**endonuclease gene**」會自動返航 (**homing**)，跑到同源染色體 (**homologous chromosome**) 的相對位置。
- Burt 的想法是：首先用基因工程的方法改變 **HEG** 切割的位置，讓它去切一個對宿主很重要的基因，這個步驟叫**基因剔除** (**gene knockout**)。而這個被剔除的基因必須具備一個條件：**單一基因被剔除時宿主沒有生存的問題**，但**當兩個都被剔除時，宿主便無法存活**。
- 接著想辦法將 **HEG** 置於減數分裂啟動子 (**meiosis promoter**) 的控制下，使得宿主產生大量含 **HEG** 的配偶子 (**gametes**)，這樣帶有 **2 個 HEG 缺陷基因** 的子代便會**全數死亡**。估計用這種方法只要放出 **1%** 的蚊子，**20 個** 世代後便能完全消滅整個族群。
- CRISPR-Cas9: Gene Drives <https://wyss.harvard.edu/media-post/crispr-cas9-gene-drives/>

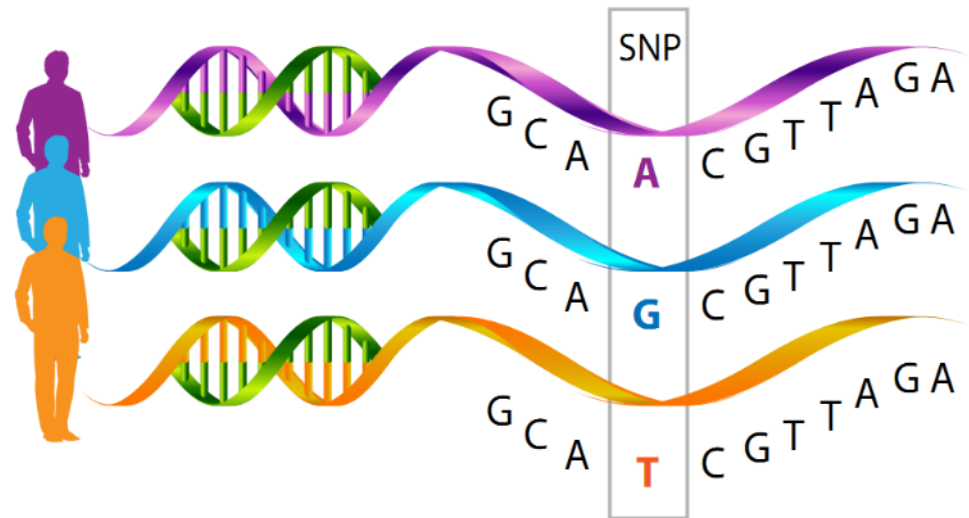
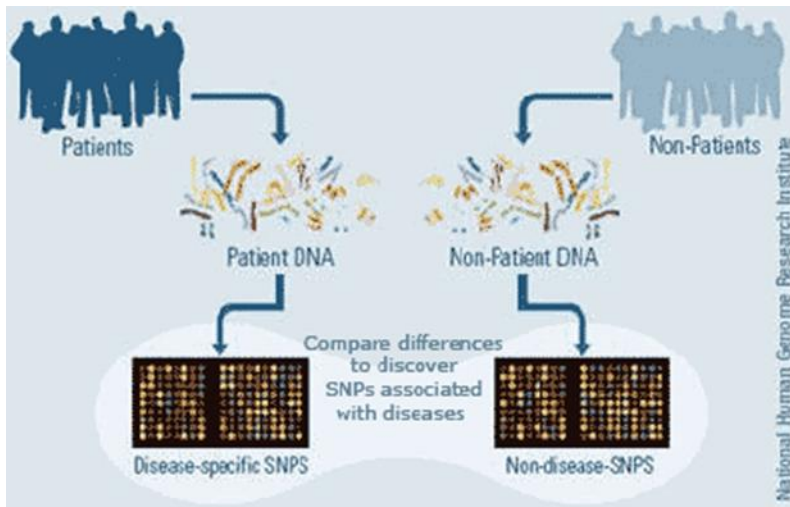
RNA interference (RNAi)

- Gene expression can also be silenced using **RNA interference (RNAi)**
 - Synthetic double-stranded RNA molecules matching the sequence of a particular gene are used to **break down or block the gene's mRNA**
 - **Reduction** in target **protein expression**
 - **Loss of Function** analysis on specific gene's product
 - Phenotypic effect; Effect on signaling/metabolic pathways, etc.
 - **Potential application for Disease treatment**
-

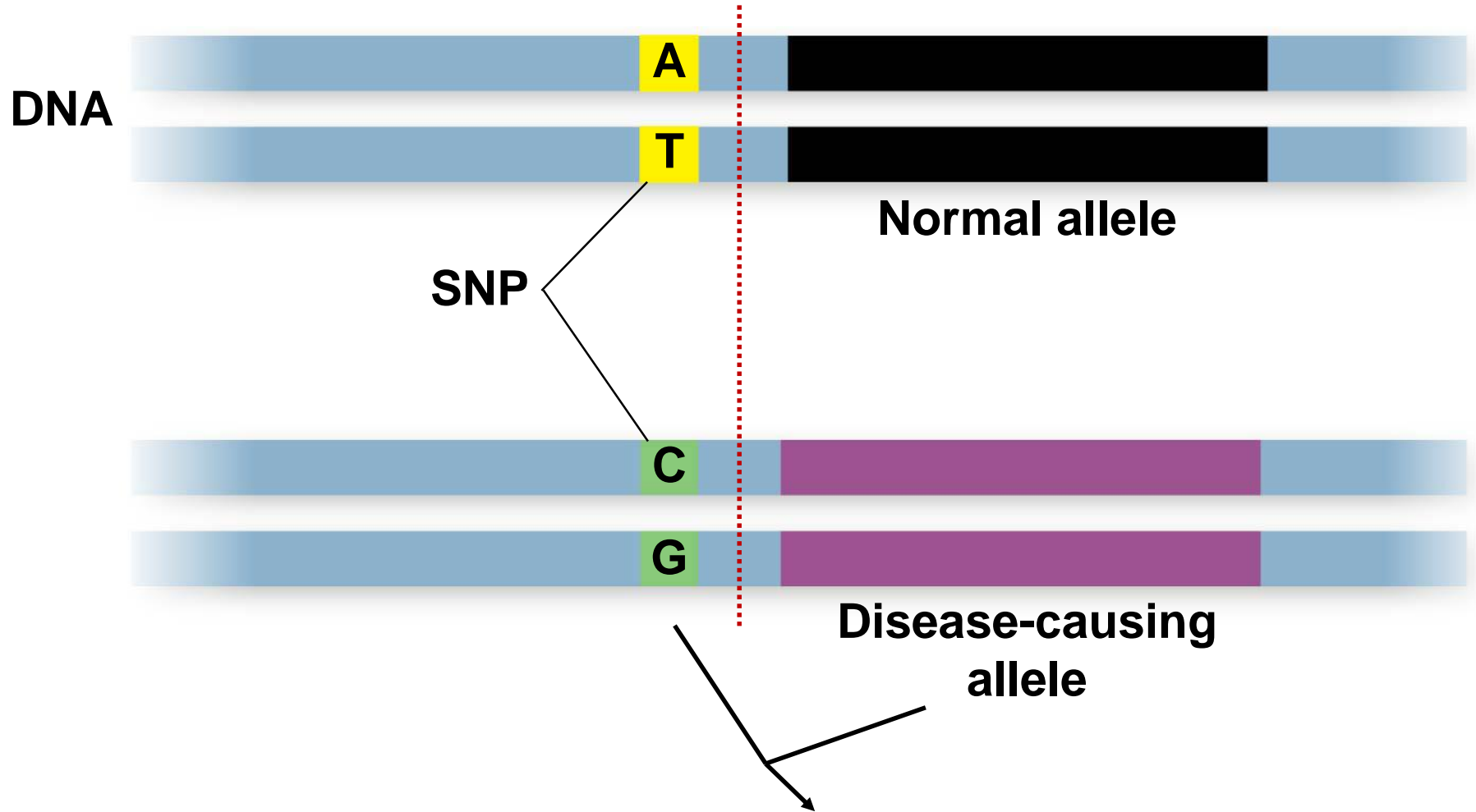
Genome-wide association studies

- In humans, researchers analyze the genomes of many people with a **certain genetic condition** to try to find nucleotide changes specific to the condition
 - **Genetic markers** called **SNPs** (**single nucleotide polymorphisms**) occur on average every 100–300 base pairs
 - SNPs can be detected by **sequencing, microarray** or **PCR**, and any SNP shared by people affected with a disorder but not among unaffected people may **locate the disease-causing gene**
-

- **SNP variants** that are found frequently associated with a particular inherited disorder alert researchers to **the most likely location for the disease-causing gene**
- SNPs are *rarely directly involved in the disease*; they are most often in **noncoding regions** of the genome



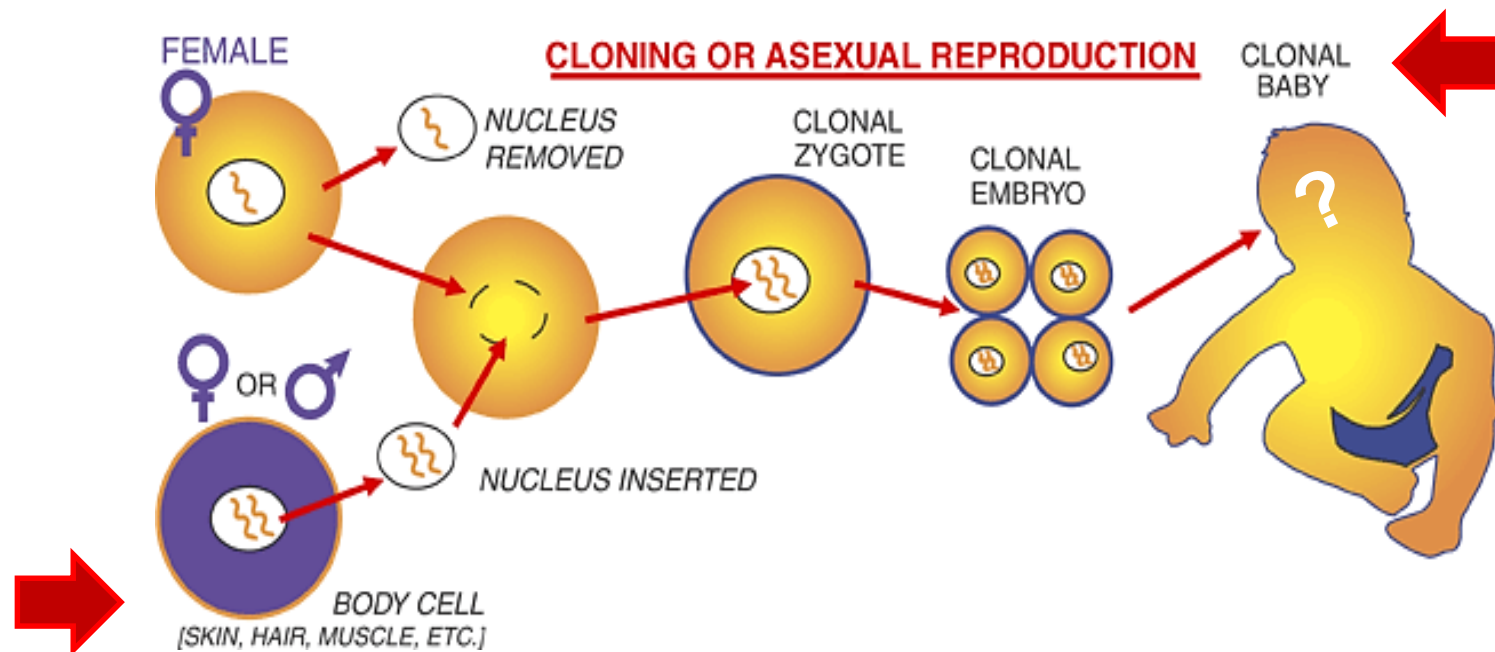
Single nucleotide polymorphisms (SNPs) as genetic markers for disease-causing alleles



Review Concept 15.3 for “linked genes”

Concept 19.3: Cloned organisms and stem cells are useful for basic research and other applications

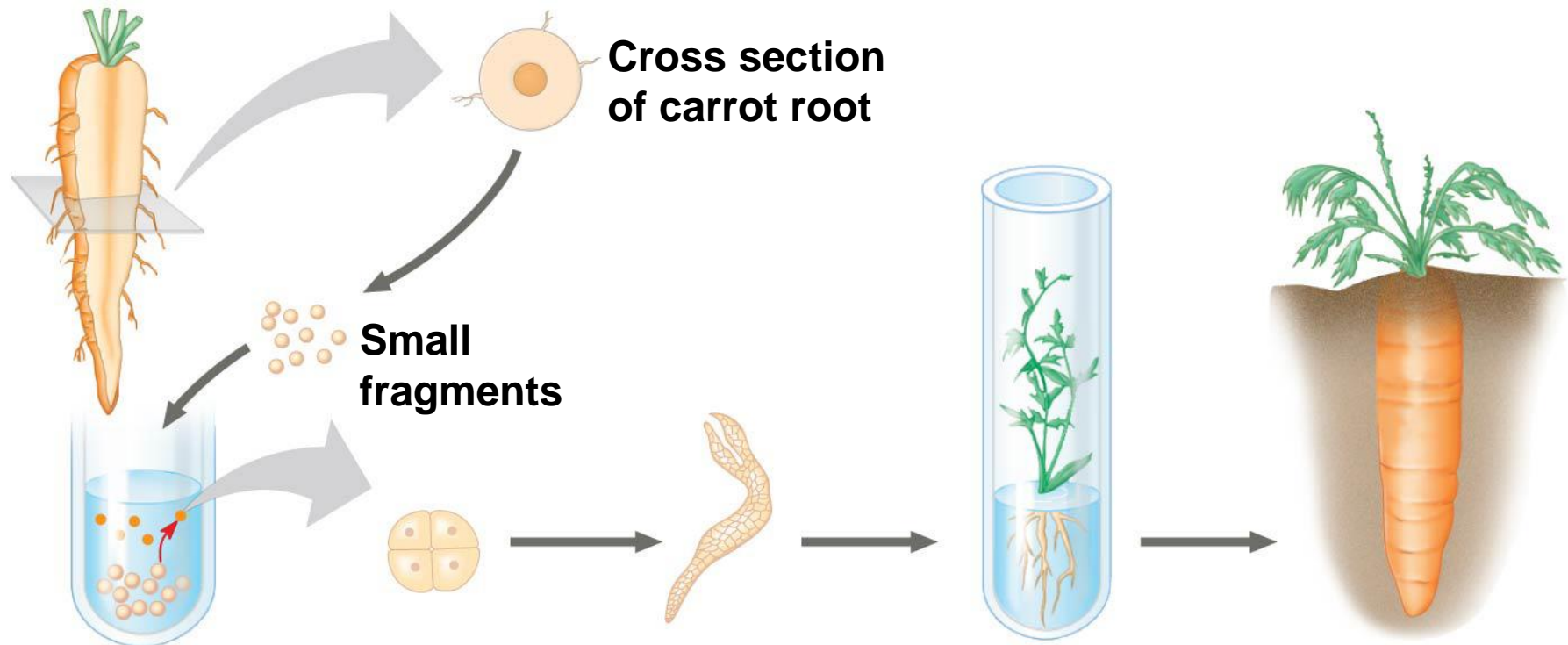
- **Organismal cloning** produces one or more organisms genetically identical to the “parent” that donated the single cell



Cloning Plants: Single-Cell Cultures

- **What is stem cell?**
 - A **stem cell** is a relatively **unspecialized cell** that can **reproduce** itself indefinitely, or under certain conditions can differentiate into one or more types of specialized cells
 - In plants, cells can dedifferentiate and then give rise to all the specialized cell types of the organism - *Plant cloning is used extensively in agriculture*
 - A **totipotent cell** (全能細胞) can generate a **complete new organism**
-

The cloning of a whole carrot plant from a single carrot cell



Fragments were cultured in nutrient medium; stirring caused single cells to shear off into the liquid.

Single cells free in suspension began to **divide**.

Embryonic plant developed from a cultured single cell.

Plantlet was cultured on agar medium. Later it was planted in soil.

Adult plant

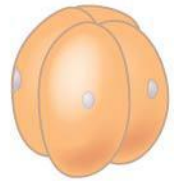
Cloning Animals: Nuclear Transplantation

- In **nuclear transplantation**, the **nucleus of an unfertilized egg cell** or zygote is replaced with the **nucleus of a differentiated cell**
 - Experiments with frog embryos have shown that a transplanted nucleus can often support normal development of the egg
 - However, **the older** the donor nucleus, **the lower** the percentage of **normally** developing tadpoles
-

Experiment

Can the nucleus from a differentiated animal cell direct development of an organism?

Frog embryo



Frog egg cell



Frog tadpole



Less differentiated cell

Fully differentiated (intestinal) cell

Donor nucleus transplanted

Enucleated egg cell

Donor nucleus transplanted

Egg with donor nucleus activated to begin development

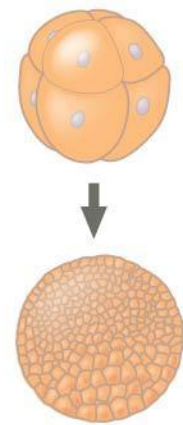
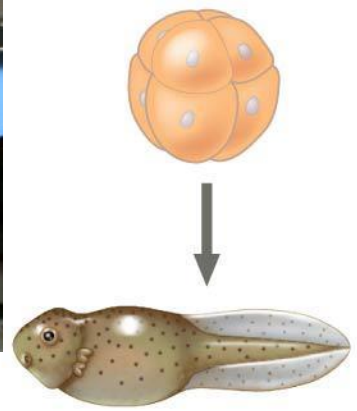
Results

Most develop into tadpoles.

Most stop developing before tadpole stage.



2012年諾貝爾醫學獎得主
John Gurdon
劍橋大學

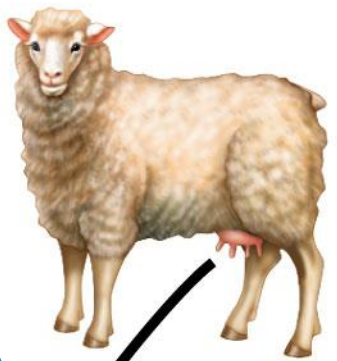


Reproductive Cloning of Mammals

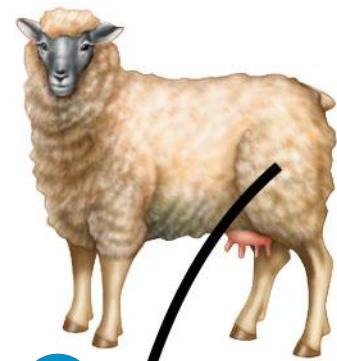
- In 1997, Scottish researchers announced the birth of **Dolly**, a lamb cloned from an adult sheep by nuclear transplantation from a differentiated mammary cell
 - Dolly's premature death in 2003, as well as her arthritis, led to speculation that her cells were not as healthy as those of a normal sheep, possibly reflecting **incomplete reprogramming** of the original transplanted nucleus
-

TECHNIQUE

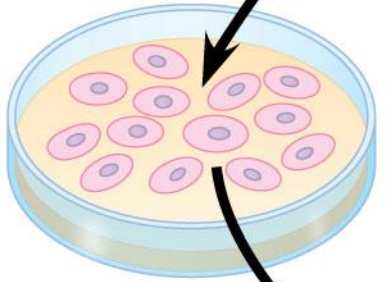
Mammary cell donor



Egg cell donor



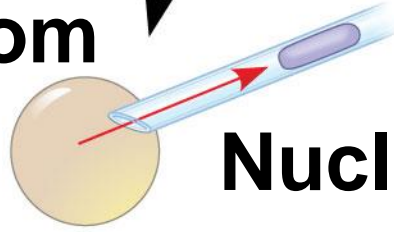
1



Cultured mammary cells

2

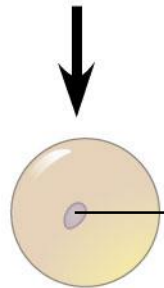
Egg cell from ovary



Nucleus removed

3

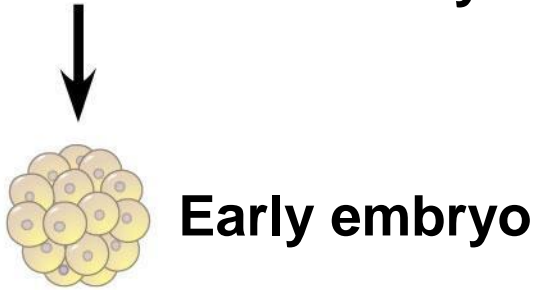
Cells fused



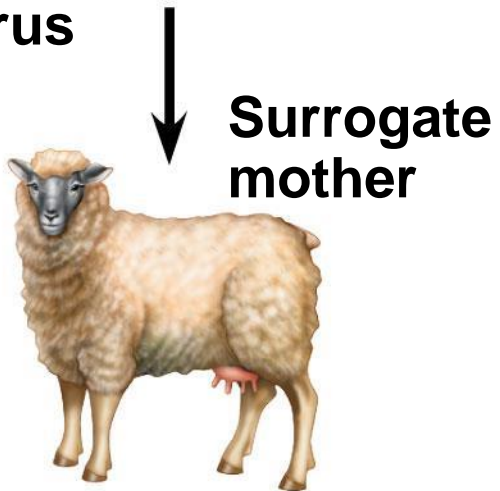
Nucleus from mammary cell



4 Grown in culture



5 Implanted in uterus of a third sheep



6 Embryonic development



RESULTS



CC, the first cloned cat

- Since 1997, cloning has been demonstrated in many mammals, including mice, cats, cows, horses, mules, pigs, and dogs
- **CC** (for **Carbon Copy**) was the first cat cloned; however, CC differed somewhat from her female “parent” due to random X chromosome inactivation.



Faulty Gene Regulation - Problems Associated with Animal Cloning

- In most nuclear transplantation studies, only a small percentage of cloned embryos have developed normally to birth
 - Many epigenetic changes, such as **acetylation of histones or methylation of DNA**, must be **reversed** in the nucleus from a donor animal in order for genes to be expressed or repressed appropriately for early stages of development
 - **Human Cloning?! – Chinese scientists claimed they have done it first @中南大學湘雅醫學院**
-

Stem Cells of Animals

- Stem cells isolated from early embryos at the **blastocyst** stage are called **embryonic stem cells**; these are able to differentiate into all cell types
- The adult body (bone marrow) also has stem cells, which replace non-reproducing specialized cells

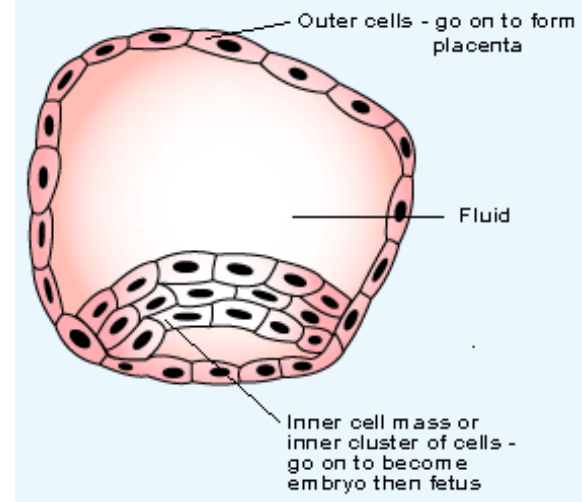
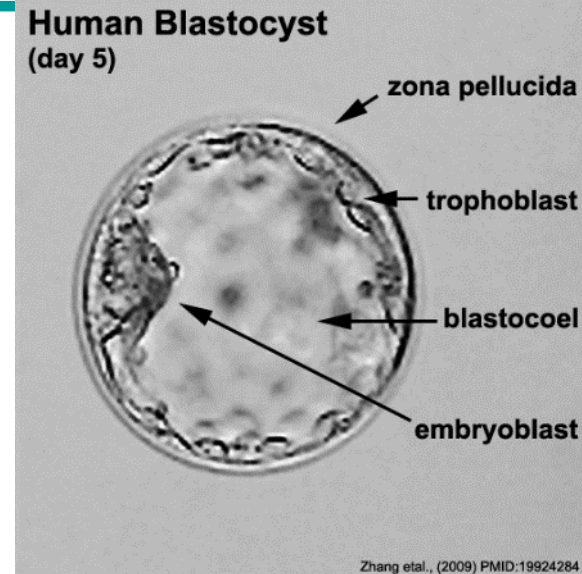
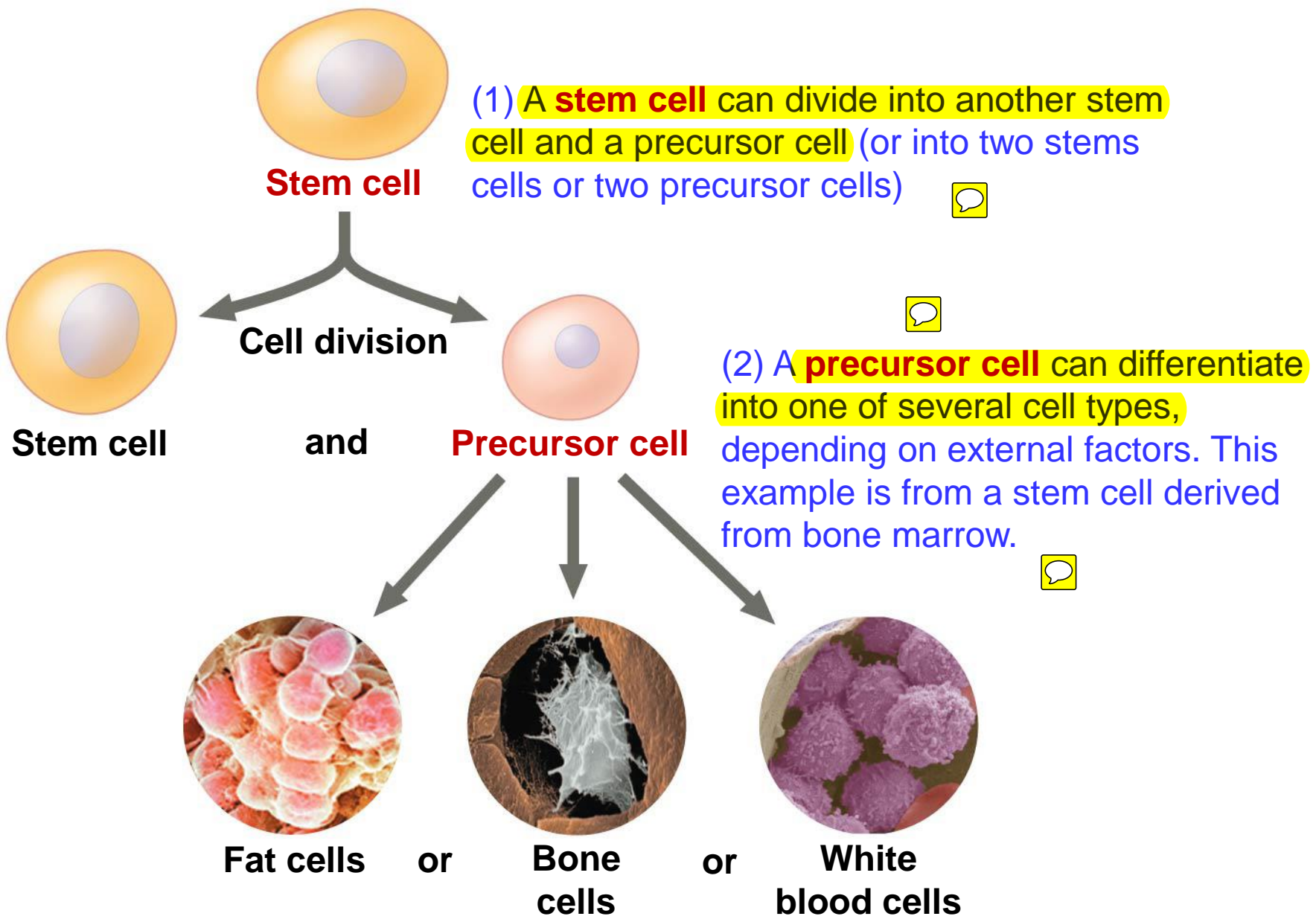


Figure 19.19

How stem cells maintain their own population and generate differentiated cells

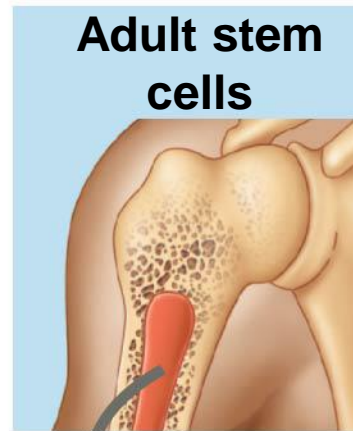
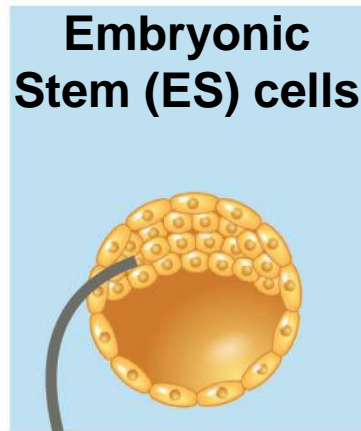


Embryonic vs. Adult Stem Cells

- Many early embryos contain stem cells capable of giving rise to **differentiated embryonic cells of any type**
- Review **Stem Cell** characteristics:
 - In culture, these embryonic stem cells **reproduce indefinitely**
 - Depending on culture conditions, they can be made to **differentiate into a variety of specialized cells**
- Adult stem cells can generate **multiple (but not all) cell types** and are used in the body to **replace** non-reproducing cells as needed

Figure 19.20

ES cells are **Pluripotent**, they can differentiate into many different cell types



Cells that can generate all embryonic cell types

Cells that generate a limited number of cell types

Cultured stem cells

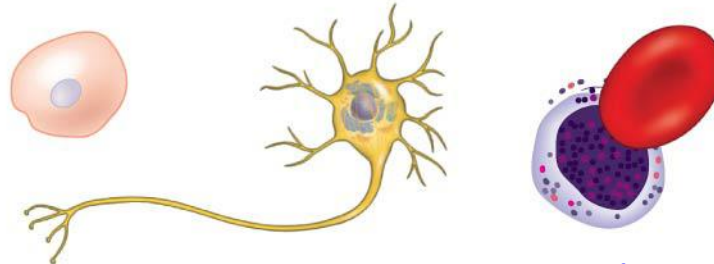
Different culture conditions

Liver cells

Nerve cells

Blood cells

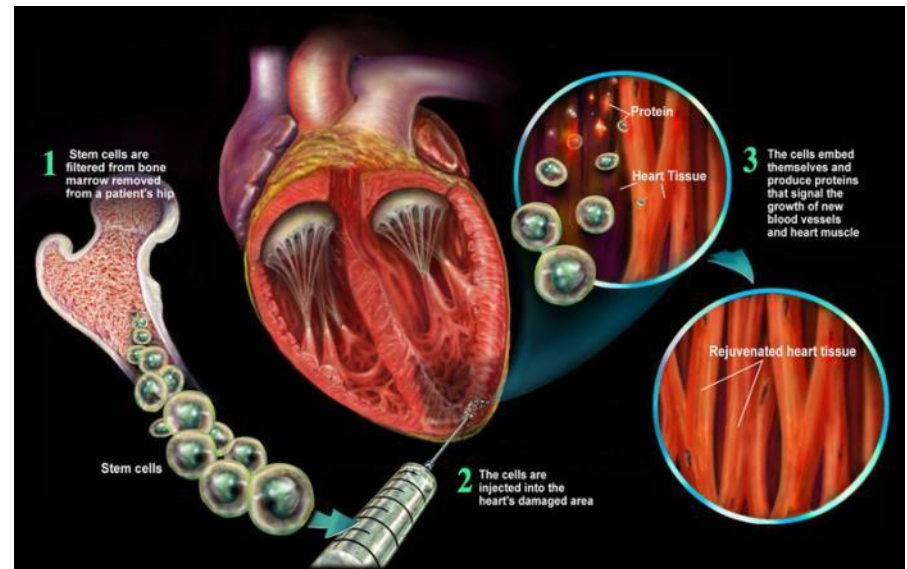
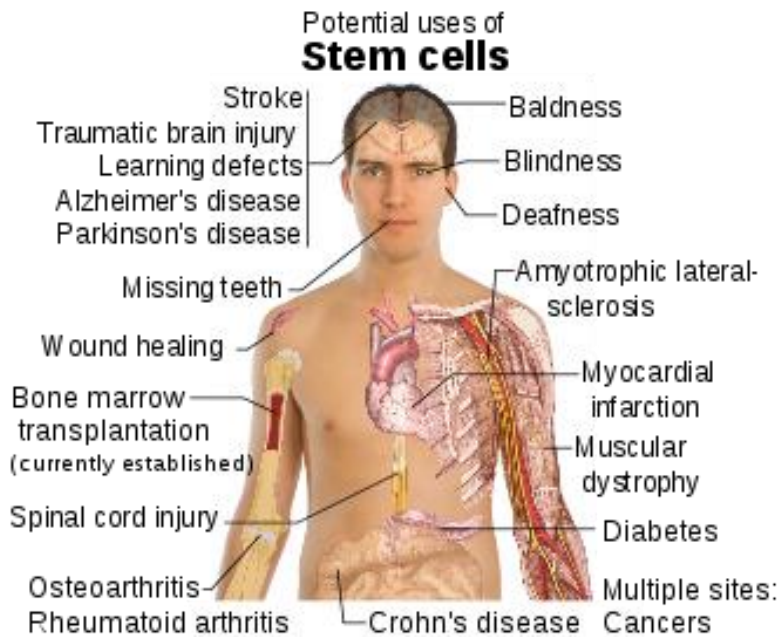
Different types of differentiated cells



Working with stem cells


Producing stem cells by therapeutic cloning

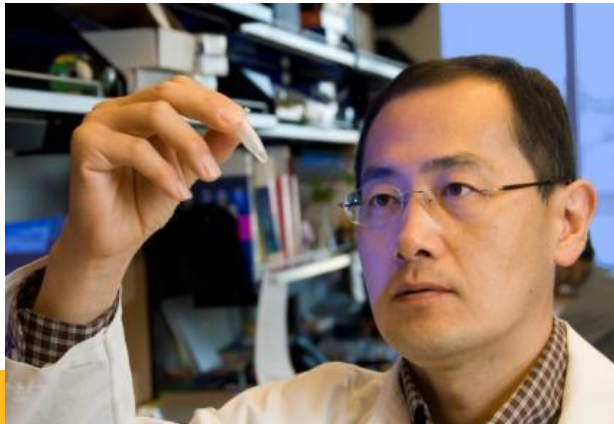
- The aim of stem cell research is to supply cells for the **repair of damaged or diseased organs** → 再生醫學



Cell therapy for myocardial infarction

Induced Pluripotent Stem (iPS) cells

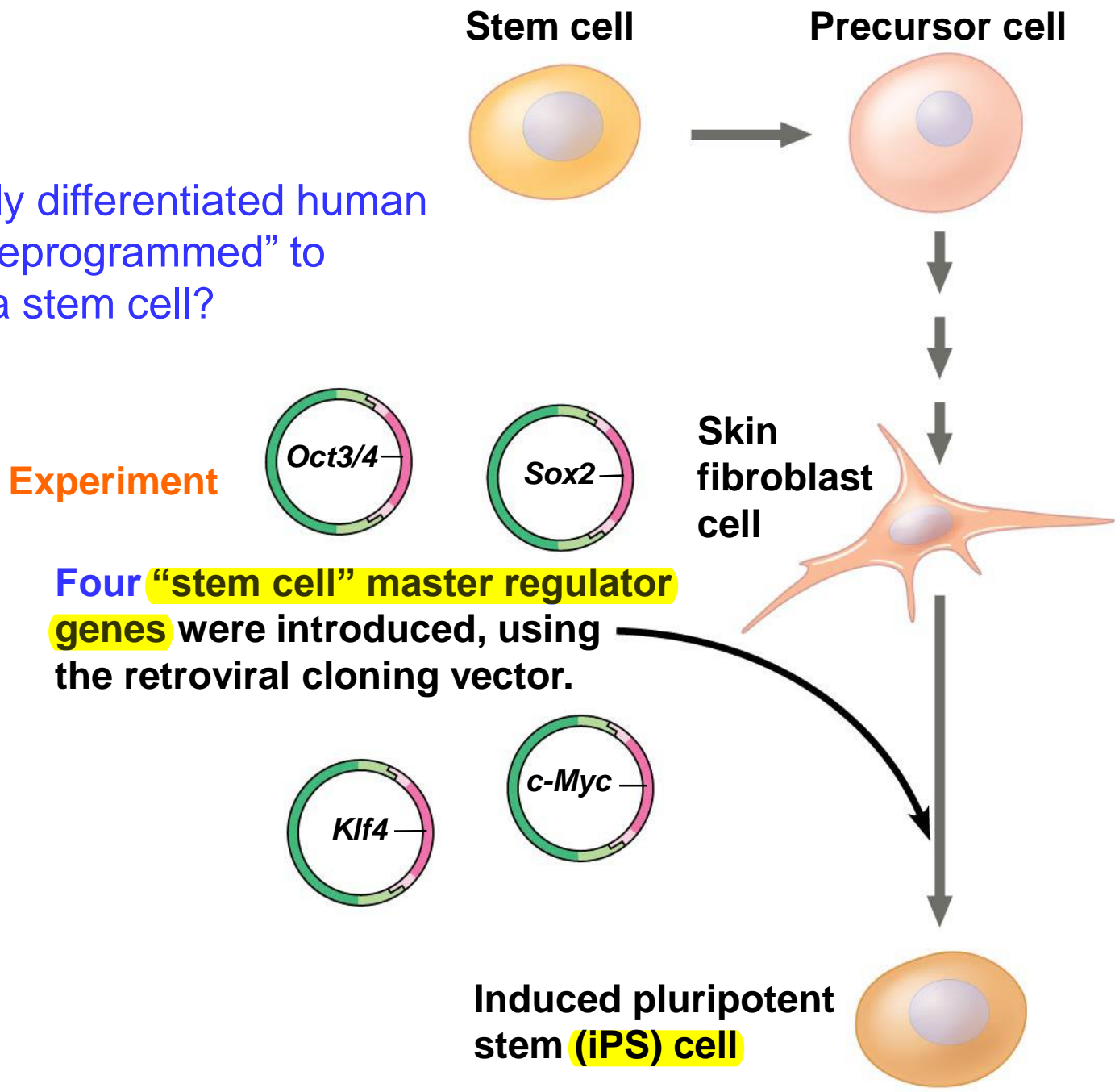
- Researchers can transform differentiated cells into ES cells by using **retroviruses** to introduce stem cell master regulatory genes 
- These transformed cells are called **iPS cells (induced pluripotent cells)** 誘導多能性幹細胞
- **iPS cells can perform most of the functions of ES cells**, and used as models for the study of certain diseases and potentially be used to replace nonfunctional cells for patients
 - **Eliminate many ethical problems!**



2012年諾貝爾醫學獎得主
Shinya Yamanaka 山中伸弥
京都大學

Figure 19.21

Can a fully differentiated human cell be “deprogrammed” to become a stem cell?



Experiment

Four “stem cell” master regulator genes were introduced, using the retroviral cloning vector.

Induced pluripotent stem (iPS) cell

One active research topic on iPSCs

Artificial sperm and egg cells of mice → 人造人? 客製人?

Cell

Reconstitution of the Mouse Germ Cell Specification Pathway in Culture by Pluripotent Stem Cells

Katsuhiko Hayashi,^{1,2,3*} Hiroshi Ohta,^{1,3} Kazuki Kurimoto,^{1,3} Shinya Aramaki,¹ and Mitunori Saitou^{1,2,3*}

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 Kyoto University, Yoshida-Konoe-cho, Sakyo-ku, Kyoto 606-8501, Japan
³JST, CREST, Yoshida-Konoe-cho, Sakyo-ku, Kyoto 606-8501, Japan
 *Correspondence: saitou@anat2.med.kyoto-u.ac.jp
 DOI 10.1016/j.cell.2011.06.052

人造精子

SUMMARY
 The generation of properly functioning gametes *in vitro* requires reconstitution of the multisteped PGC fate (Kurimoto et al., 2008; Ohinata et al., 2005; Vincent et al., 2005; Yamaj et al., 2008). Essentially all of the epiblast cells from E5.5–E6.0 are competent to express *Blimp1* and *Prdm14* in response to *Bmp4*, and the PGC-like cells induced from the epiblasts *in vitro* can form functional germ cells *in vivo*.

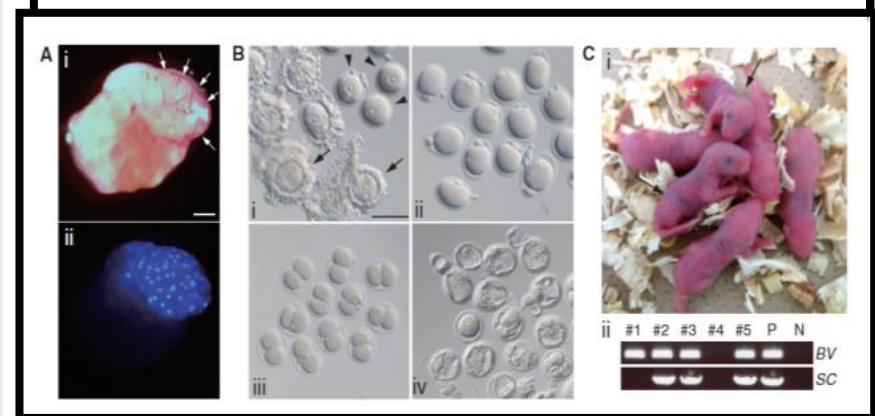
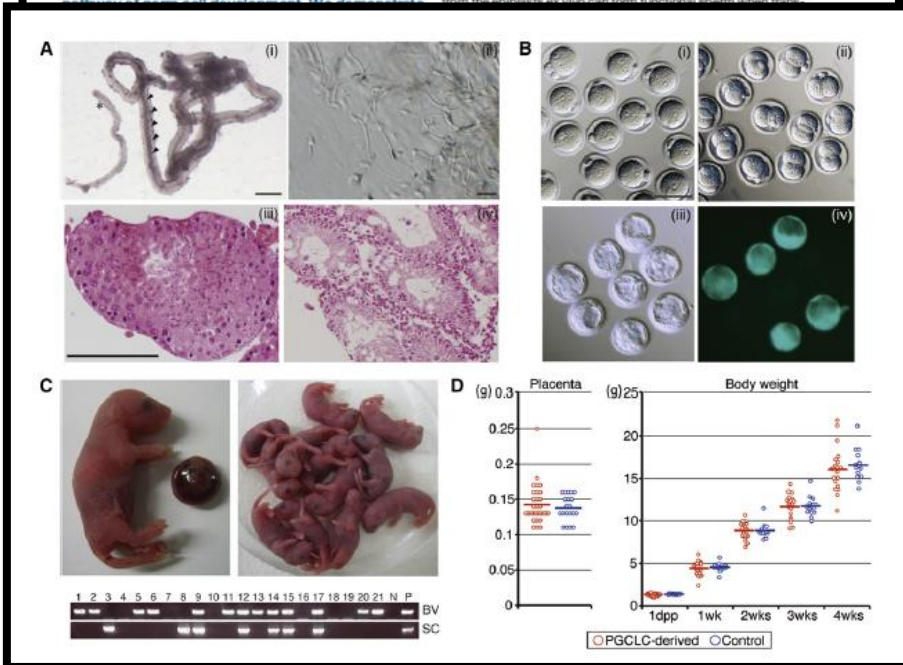
Offspring from Oocytes Derived from *In Vitro* Primordial Germ Cell-like Cells in Mice

Katsuhiko Hayashi,^{1,2,3*} Sugako Ogushi,^{1,4} Kazuki Kurimoto,^{1,5} So Shimamoto,¹ Hiroshi Ohta,^{1,5} Mitunori Saitou^{1,2,5,6*}

Reconstitution of female germ cell development *in vitro* is a key challenge in reproductive biology and medicine. We show here that female (XX) embryonic stem cells and induced pluripotent stem cells in mice are induced into primordial germ cell-like cells (PGCLCs), which, when aggregated with female gonadal somatic cells as reconstituted ovaries, undergo X-reactivation, imprint erasure, and cyst formation, and exhibit meiotic potential. Upon transplantation under mouse ovarian bursa, PGCLCs in the reconstituted ovaries mature into germinal vesicle-stage oocytes, which then contribute to fertile offspring after *in vitro* maturation and fertilization. Our culture system serves as a robust foundation for the investigation of key properties of female germ cells, including the acquisition of totipotency, and for the reconstitution of whole female germ cell development *in vitro*.

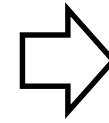
The germ cell lineage in mammals originates from pluripotent epiblasts as primordial germ cells (PGCs) and undergoes sexually dimorphic development, generating spermatozoa in males and oocytes in females. These cells fertilize to form zygotes with full developmental potential.

mag.org **SCIENCE** VOL 338 16 NOVEMBER 2012 **人造卵子** 971



In a world first, Japanese team creates eggs and sperm of endangered species from iPS cells

Science Advances 12 May 2017: Vol. 3, no. 5, e1602179, DOI: 10.1126/sciadv.1602179

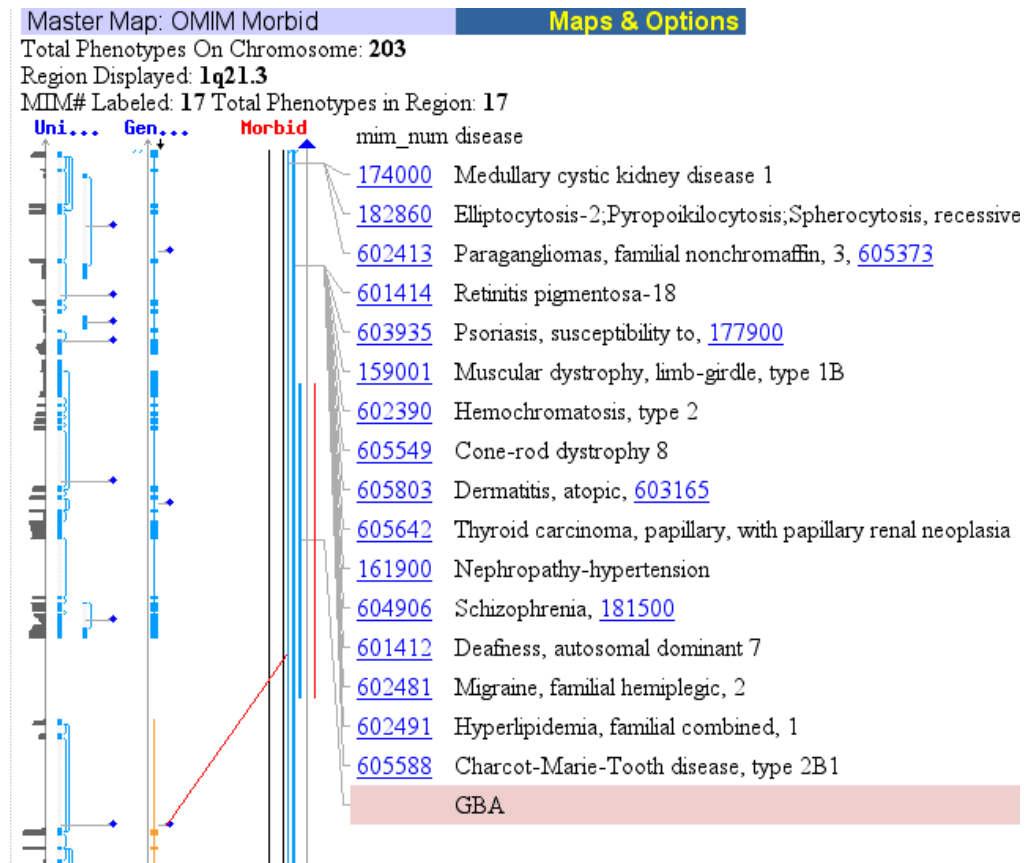


Concept 19.4: The practical applications of DNA-based technology affect our lives in many ways

- Many fields benefit from DNA technology and genetic engineering, some examples are:
 - Medical applications
 - Diagnosis of genetics diseases
 - Developing therapy or drug
 - Forensic evidence and genetic profiles
 - Environmental cleanup
 - Agriculture applications
-

Medical Applications

- One benefit of DNA technology is **identification of human genes** in which **mutation** plays a role in genetic diseases



Genes and Diseases

<http://www.ncbi.nlm.nih.gov/books/NBK22183/>

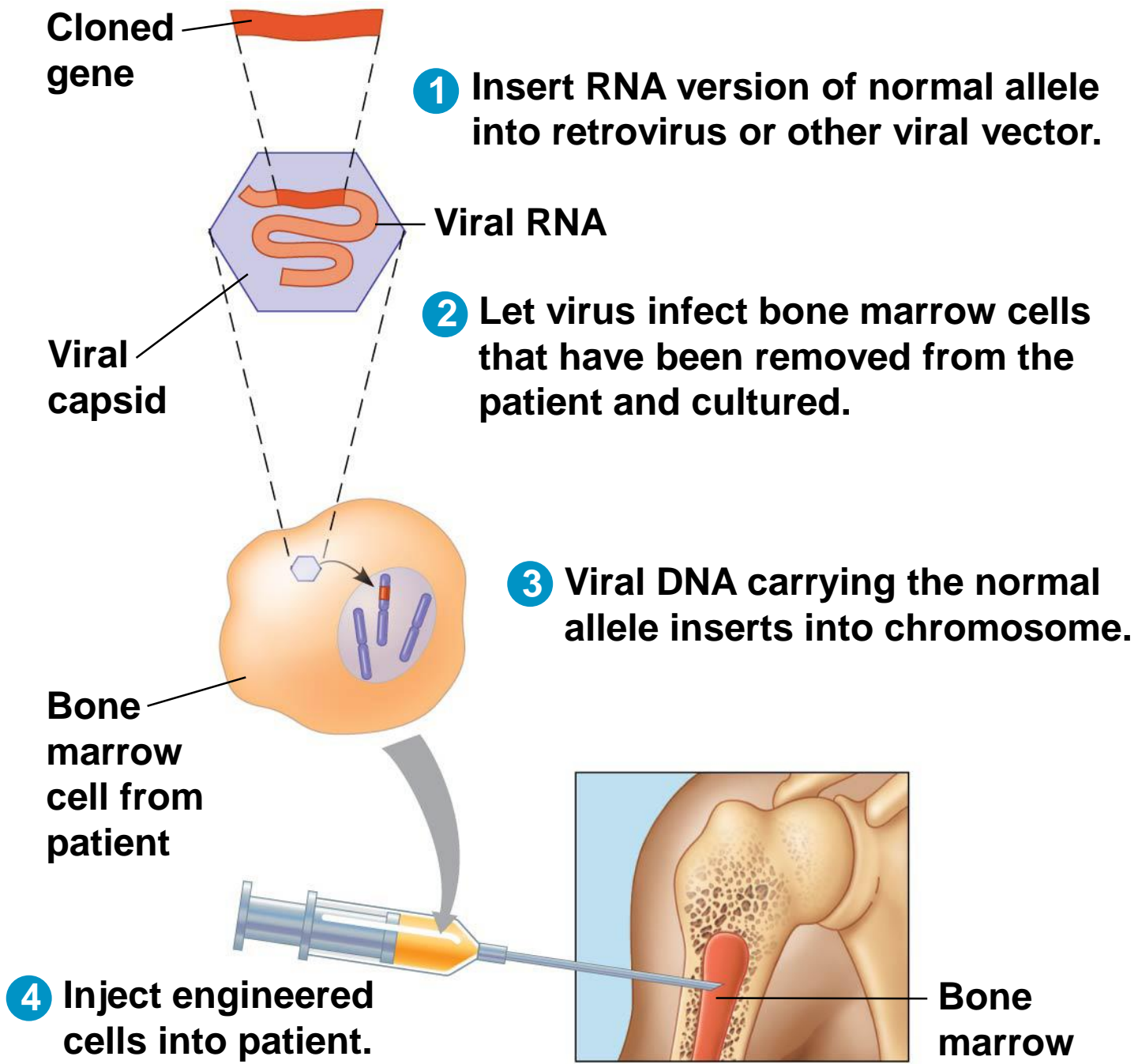
Diagnosis of Diseases

- Scientists can diagnose many human genetic disorders by using PCR and specific primers corresponding to cloned disease genes, then sequencing the amplified product to look for the disease-causing mutation
 - SNPs may be associated with a disease-causing mutation
 - SNPs may also be correlated with increased risks for conditions such as heart disease or certain types of cancer
-

Human Gene Therapy

- **Gene therapy** is the alteration of an afflicted individual's genes
 - Gene therapy holds great potential for treating disorders traceable to a single defective gene
 - **Vectors are used for delivery of genes into specific types of cells, for example bone marrow**
 - Gene therapy raises ethical questions, such as whether human germ-line cells should be treated to correct the defect in future generations
-

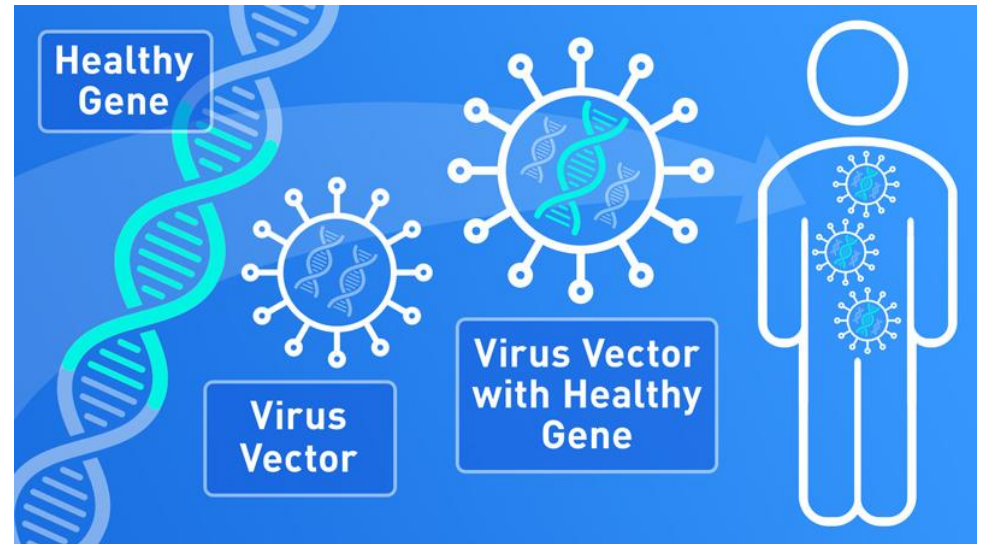
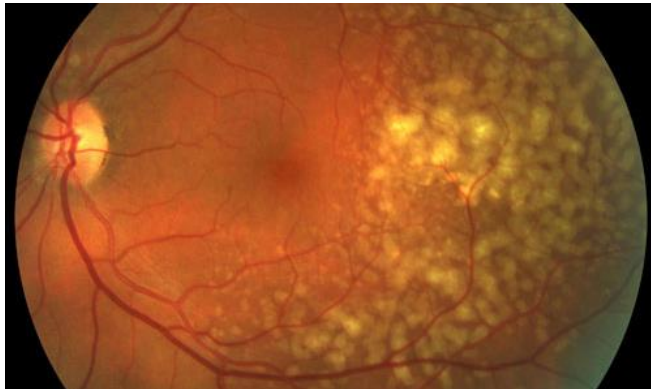
Figure 19.22



FDA approves novel gene therapy to treat patients with a rare form of inherited vision loss on 2017/12/19

補充

Hereditary retinal dystrophies are a broad group of genetic retinal disorders of varying severity and with differing inheritance patterns.



- **Luxturna** is the first gene therapy approved in the U.S. to target a disease caused by mutations in a specific gene
- It is an **adeno-associated virus vector-based gene therapy** indicated for the treatment of patients with confirmed bi-allelic **RPE65** mutation-associated retinal dystrophy. Patients must have viable retinal cells as determined by the treating physician(s).

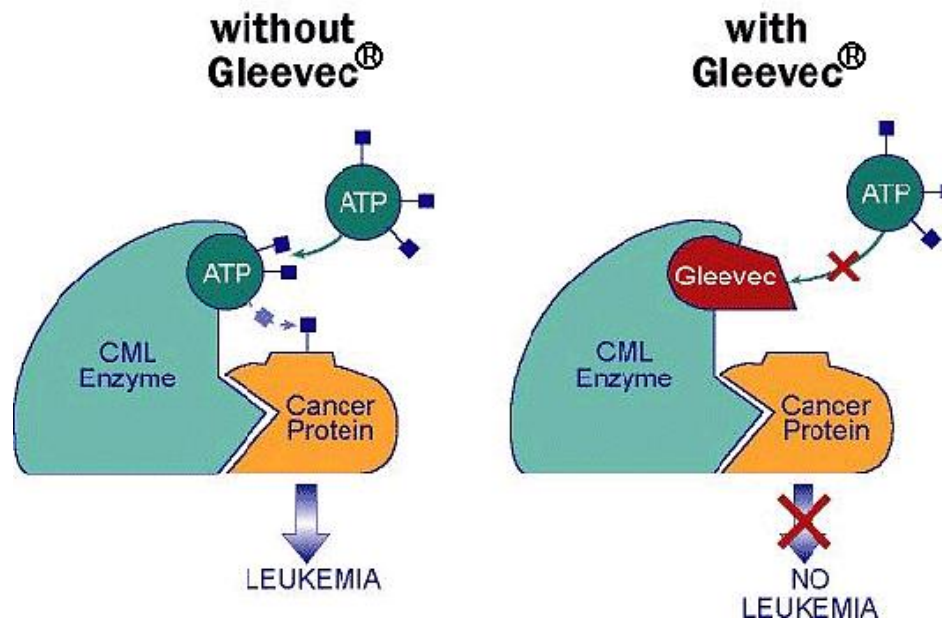
Pharmaceutical Products

- Advances in DNA technology and genetic research are important to the **development of new drugs** to treat diseases



Synthesis of Small Molecules for Use as Drugs

- The drug **imatinib** (Gleevec) is a small molecule that inhibits over-expression of a specific leukemia (CML)-causing **receptor tyrosine kinase**



建議閱讀

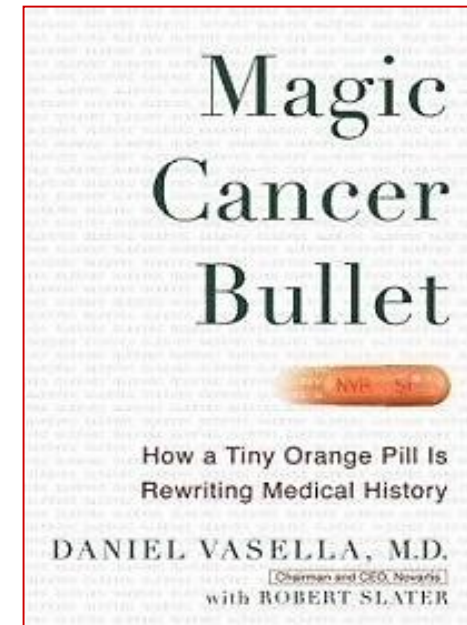
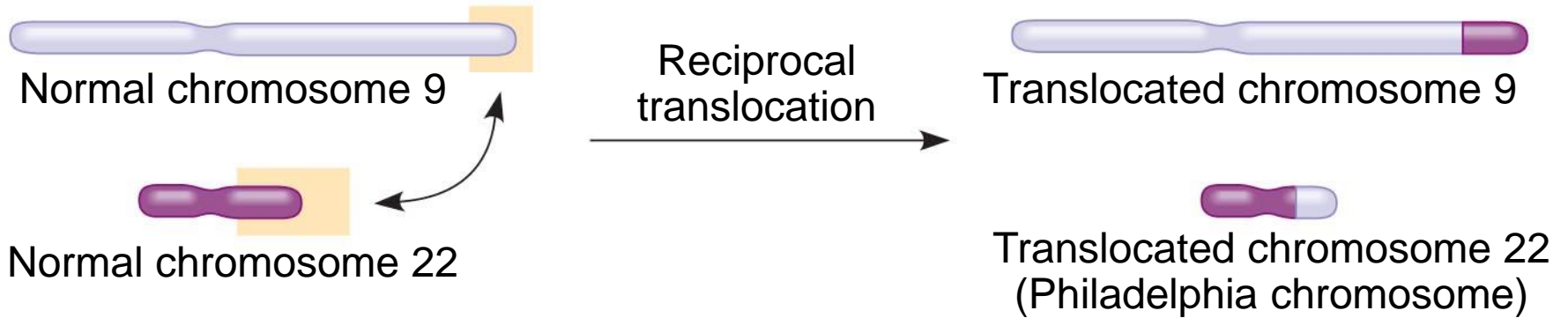
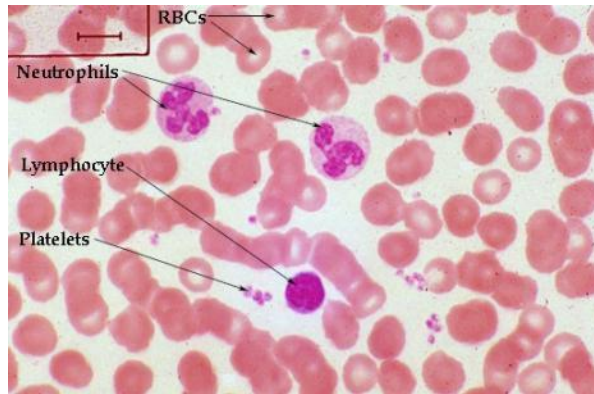


Fig.15-16

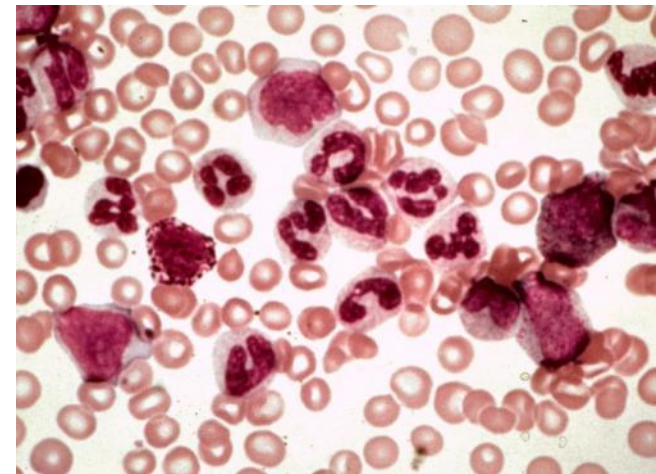
Translocation associated with chronic myelogenous leukemia (CML)



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Blood smear assay



Many **immature white blood cells**

Protein Drug Production in Cell Cultures

- Pharmaceutical products that are proteins can be synthesized on a large scale
 - Host cells in culture can be engineered to secrete a protein as it is made
 - This is useful for the production of **insulin, human growth hormones, antibodies and vaccines**
-

“Pharm” Animals and Plants

Protein Production by “Pharm” Animals and Plants

- **Transgenic** animals/plants are made by introducing genes from one species into the genome of another animal/plant
 - Transgenic animals/plants are **pharmaceutical “factories,”** producers of large amounts of otherwise rare substances for medical use
-

Goats as “pharm” animals – milking “antithrombin”



Protein structure of **Antithrombin**

Transgenic Goat bred to **secret antithrombin into its milk** - Such goat was produced by **Nuclear transfer**, performed in a laboratory dish, entails taking a cell carrying the new DNA, which codes for the desired therapeutic protein, and fusing it to a donor goats egg (that has had its genetic material removed). These eggs are then implanted into a surrogate female goat which she will carry to term.

THE JOURNAL OF BIOLOGICAL CHEMISTRY
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Vol. 276, No. 18, Issue of May 4, pp. 14961-14971, 2001
Printed in U.S.A.

Heparin Enhances the Specificity of Antithrombin for Thrombin and Factor Xa Independent of the Reactive Center Loop Sequence

EVIDENCE FOR AN EXOSITE DETERMINANT OF FACTOR Xa SPECIFICITY IN HEPARIN-ACTIVATED ANTITHROMBIN*

Received for publication, December 21, 2000
Published, JBC Papers in Press, February 7, 2001, DOI 10.1074/jbc.M011550200

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Regulation of the Blood Clotting Cascade (凝血級聯機制) By Antithrombin

Intrinsic Pathway

Extrinsic Pathway (Tissue Damage)

Factor IX

Factor XIa

Factor IXa

Factor X

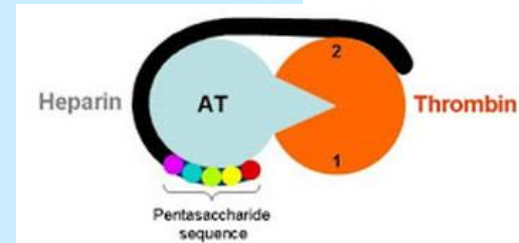
Factor Xa

Prothrombin

Thrombin

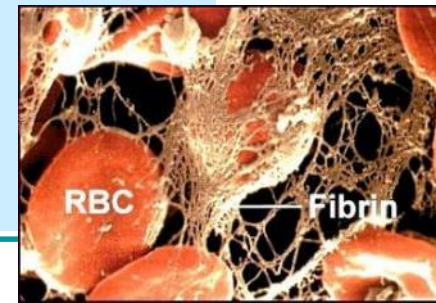
Fibrinogen

Fibrin Clot



**Antithrombin
(+Heparin)**

*Inhibition in Red



Forensic Evidence and Genetic Profiles

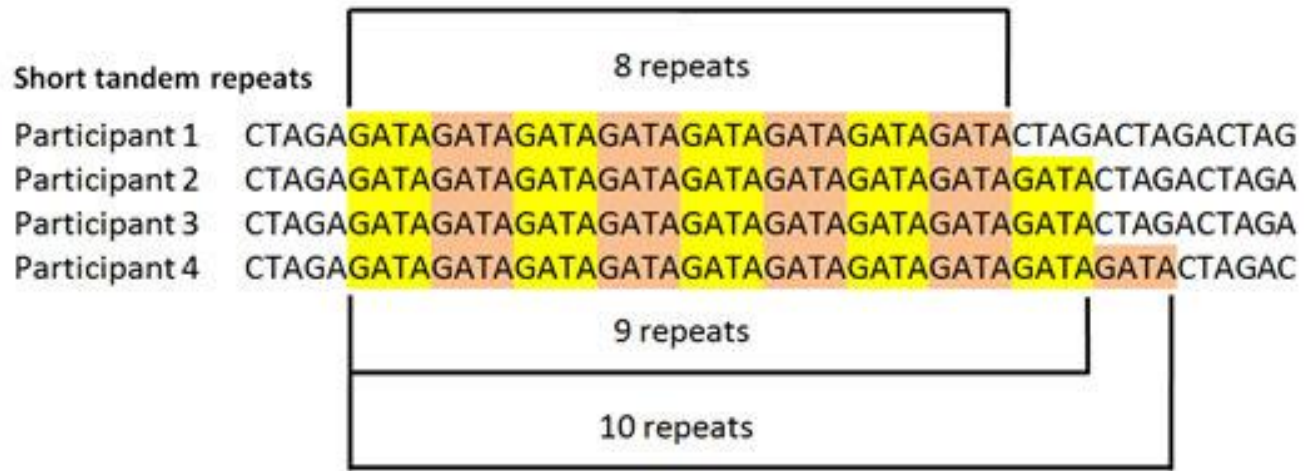
- An individual's unique DNA sequence, or **genetic profile**, can be obtained by analysis of tissue or body fluids
 - DNA testing can be used to provide evidence in criminal and paternity cases and to identify human remains
 - Genetic profiles are currently analyzed using genetic markers called **short tandem repeats (STRs)**
-

Short tandem repeats (STRs)

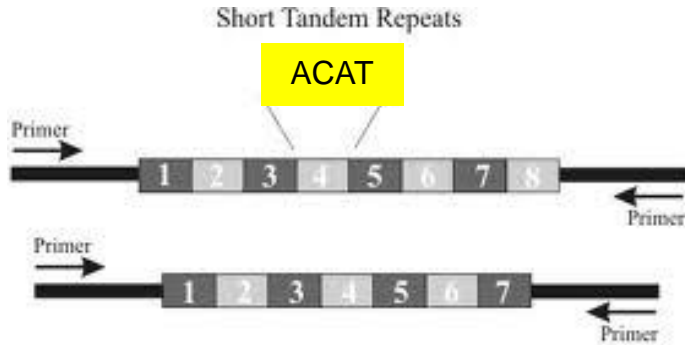
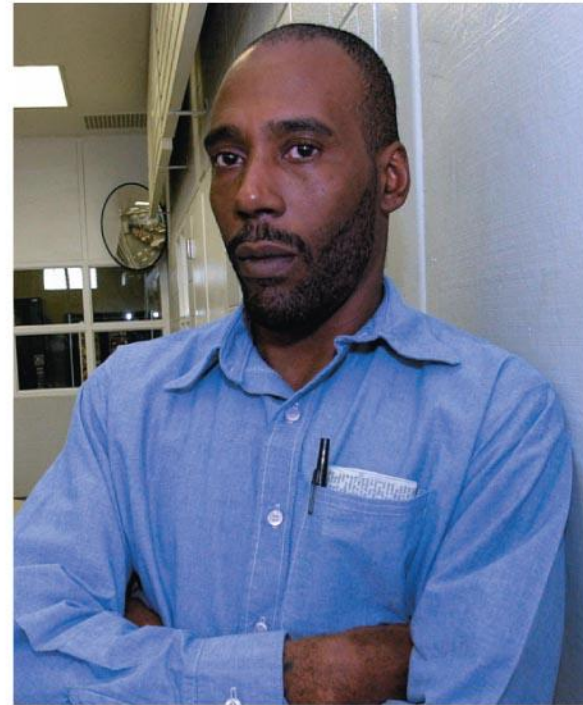
- **STRs** are variations in the number of repeats of specific DNA sequences (Explanation on next page)
 - PCR and gel electrophoresis are used to amplify and then identify STRs of different lengths
 - The probability that two people who are not identical twins have the same STR markers is exceptionally small
-

Short tandem repeat

- STR is a series of repeating units of 2 to 5 (or 2~16) nucleotides
- The repeat number for STRs can vary among sites (within a genome) or individuals



(a) This photo shows Earl Washington just before his release in 2001, after 17 years in prison.



The flanking regions where PCR primers bind are constant

Homozygote = both alleles are the same length

Heterozygote = alleles differ and can be resolved from one another

Source of sample	STR marker 1	STR marker 2	STR marker 3
Semen on victim	17, 19	13, 16	12, 12
Earl Washington	16, 18	14, 15	11, 12
Kenneth Tinsley	17, 19	13, 16	12, 12

} No. of repeats

(b) These and other STR data exonerated Washington and led Tinsley to plead guilty to the murder.

Environmental Cleanup

- Genetic engineering can be used to modify the **metabolism of microorganisms**
 - Some modified microorganisms can be used to **extract minerals** from the environment or **degrade potentially toxic waste materials (or oil spills)**
- **Biofuels** make use of crops such as corn, soybeans, and cassava (木薯) to replace fossil fuels



Animal Husbandry

- Genetic engineering of transgenic animals speeds up the selective breeding process
- **Beneficial genes** can be transferred **between varieties or species**



Agricultural Applications

- DNA technology is being used to improve **agricultural productivity** and **food quality**
-

Meanwhile, we need to preserve the nature's vanishing **biodiversity** with their DNA codes/Genomes → *Conservation Biology*



李家維教授 熱帶植物保種中心 <http://www.kbcc.org.tw/>

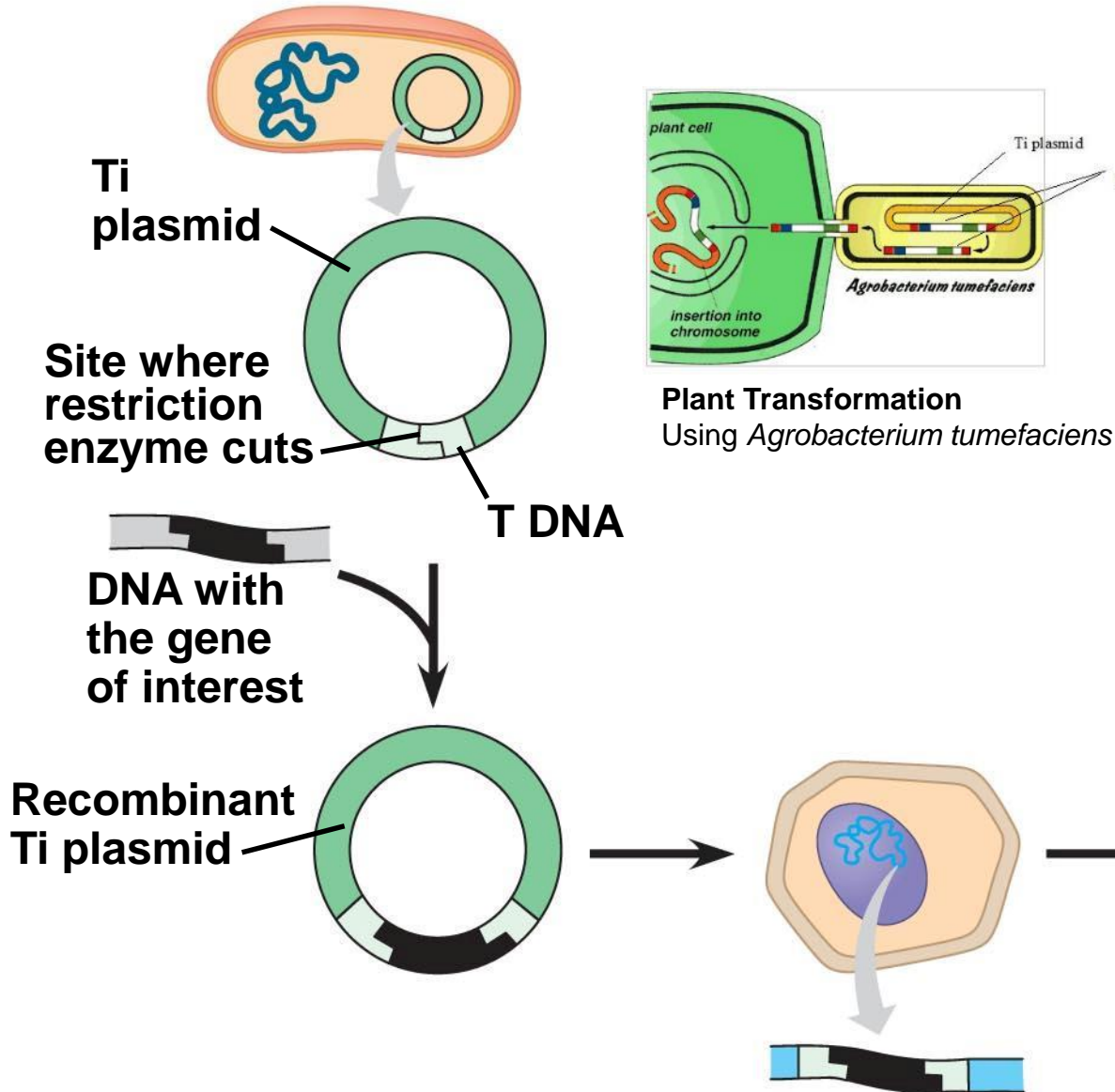
Genetic Engineering in Plants

- Agricultural scientists have endowed a number of crop plants **with genes for desirable traits**
 - The **Ti plasmid** is the most commonly used vector for **introducing new genes into plant cells**
 - Recently, CRISPR Cas9 system has been tried for plant genome editing as well
 - Genetic engineering in plants has been used to transfer many useful genes including those for **herbicide resistance**, increased **resistance to pests**, increased **resistance to salinity**, and improved **nutritional value** of crops.
-

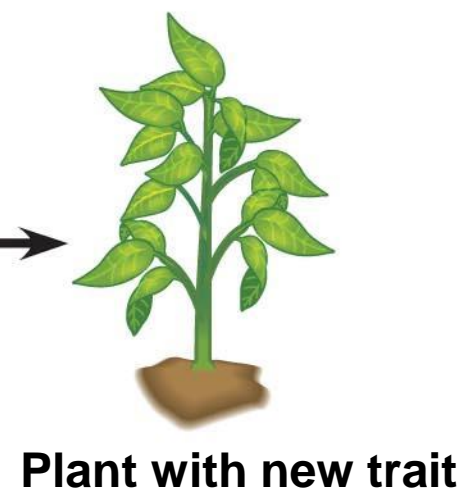
TECHNIQUE

Using the Ti plasmid to produce transgenic plants

Agrobacterium tumefaciens 農桿菌



RESULTS



Safety and Ethical Questions Raised by DNA Technology

- Potential **benefits** of genetic engineering must be weighed against potential **hazards** of creating harmful products or procedures
 - Guidelines are in place in the United States and other countries to ensure **safe practices for recombinant DNA technology**
-

GMO – genetically modified organism

- Most public concern about possible hazards centers on **genetically modified (GM) organisms** used as food
- Some concerns: (1) the creation of “**super weeds**” from the transfer of genes from GM crops to their wild relatives; (2) allergic reactions



GMO from the nature:

Solar-powered sea slug harnesses stolen plant genes



Horizontal gene transfer of the algal 藻 nuclear gene *psbO* to the photosynthetic sea slug *Elysia chlorotica* 海蛞蝓

PNAS (2008) Vol 18 No 46 P17867-17871



Ethical guidelines

- As biotechnology continues to change, so does its use in agriculture, industry, and medicine
- National agencies and international organizations strive to set **guidelines for safe and ethical practices** in the use of biotechnology

In May 2016, “**No substantiated evidence of a difference in risks to human health between GMO crops & conventional crops**” was declared by National Academics of Science, Engineering, and Medicine. <https://www.ncbi.nlm.nih.gov/books/NBK424543/>

You should now be able to:

1. Describe the natural function of restriction enzymes and explain how they are used in recombinant DNA technology
 2. Outline the procedures for cloning a eukaryotic gene in a bacterial plasmid
 3. Define and distinguish between genomic libraries using plasmids, phages, and cDNA
 4. Describe the polymerase chain reaction (PCR) and explain the advantages and limitations of this procedure
-

-
5. Explain how gel electrophoresis is used to analyze nucleic acids and to distinguish between two alleles of a gene
 6. Describe and distinguish between the Southern blotting procedure, Northern blotting procedure, and RT-PCR
 7. Distinguish between gene cloning, cell cloning, and organismal cloning
 8. Describe how nuclear transplantation was used to produce Dolly, the first cloned sheep
-

-
9. Describe the application of DNA technology to the diagnosis of genetic disease, the development of gene therapy, vaccine production, and the development of pharmaceutical products
 10. Define a SNP and explain how it may produce a RFLP
 11. Explain how DNA technology is used in the forensic sciences
-

12. Discuss the safety and ethical questions related to recombinant DNA studies and the biotechnology industry

Supporting information

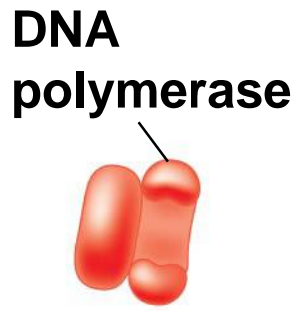
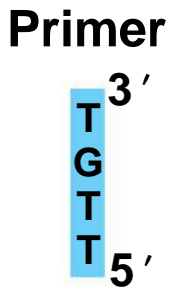
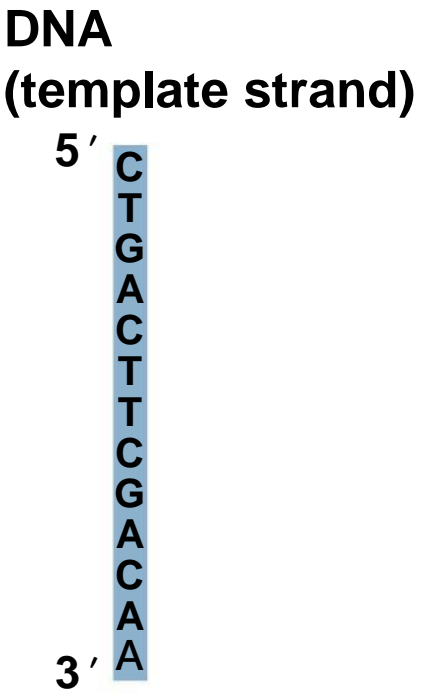
DNA Sequencing (Sanger method)

- Relatively short DNA fragments can be sequenced by the ***dideoxy or chain termination sequencing***, developed by Sanger
 - Modified nucleotides called **dideoxynucleotides (ddNTP)** attach to synthesized DNA strands of different lengths
 - Each type of ddNTP is **tagged with a distinct fluorescent label** that identifies the nucleotide at the end of each DNA fragment
 - The DNA sequence can be read from the resulting spectrogram
-

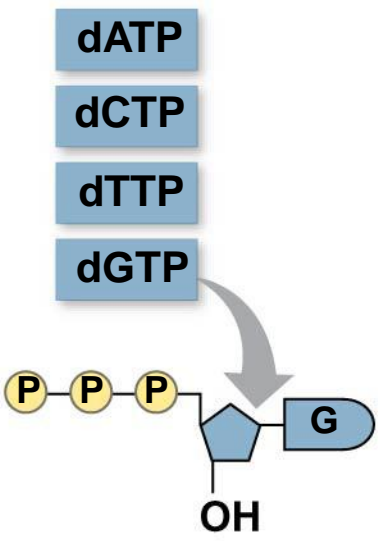
Figure 19.3a

DNA sequencing by Dideoxy or Chain Termination method

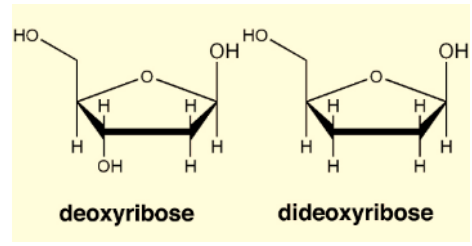
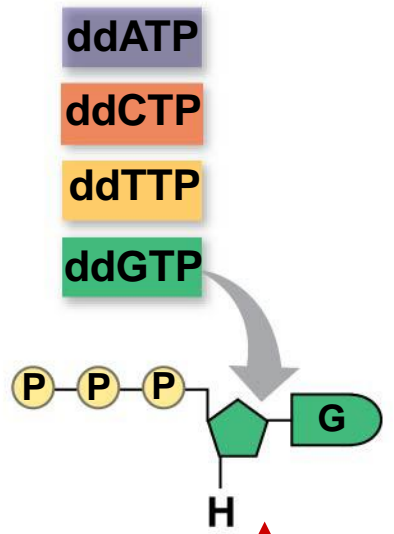
Technique



Deoxyribo-nucleotides



Dideoxynucleotides (fluorescently tagged)



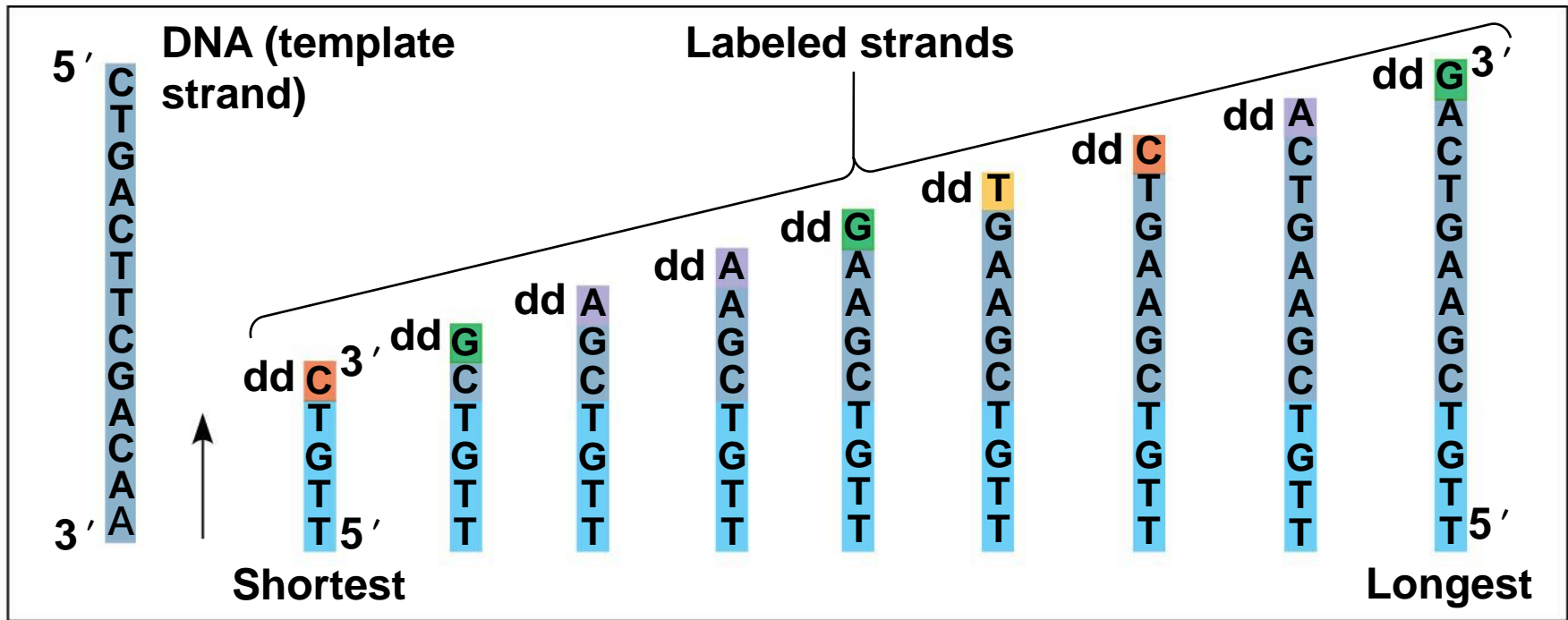
Key reagents and reactions

Chain reaction terminated

Figure 19.3b

- In the test tube, synthesis of each new strand starts at the same 3' end of the template.
- Due to random termination of the chain reaction, **a set of labeled strands of every possible length** is generation.

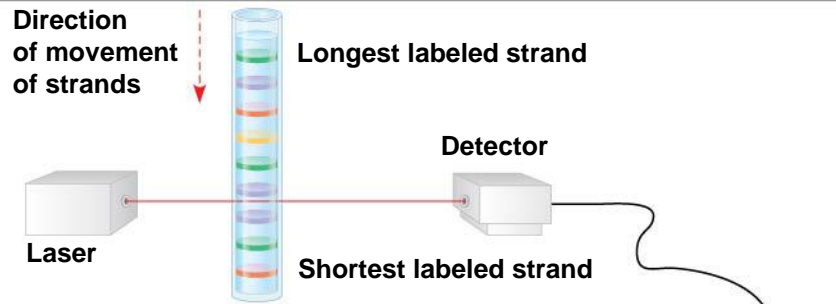
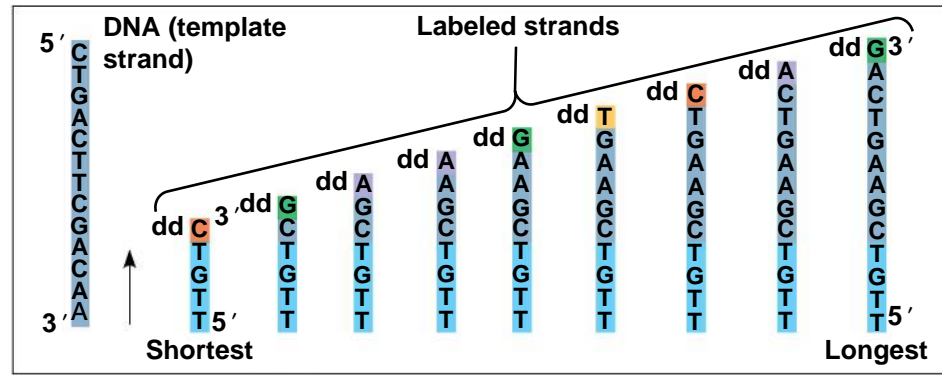
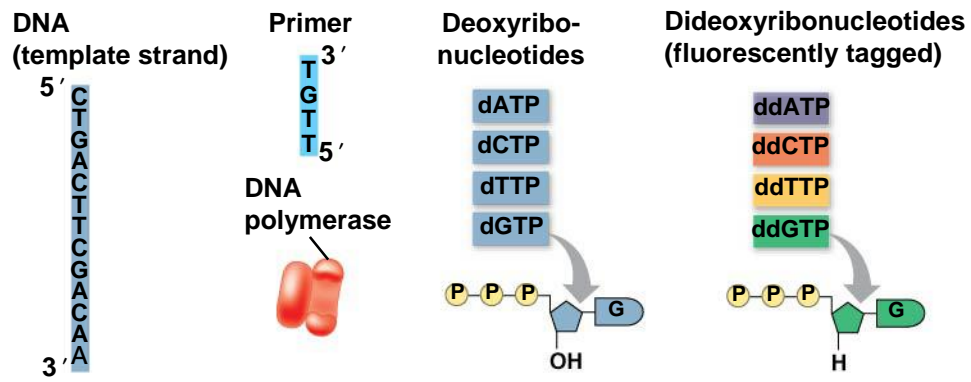
Technique



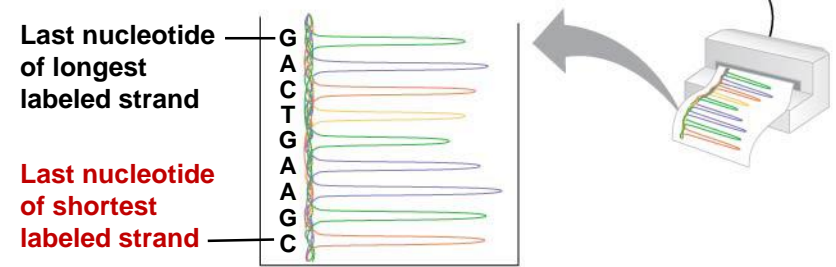
The color of the terminal tag (i.e., fluorescent molecule) represents the **last nucleotide** in each strand along the sequence.

Figure 19.3

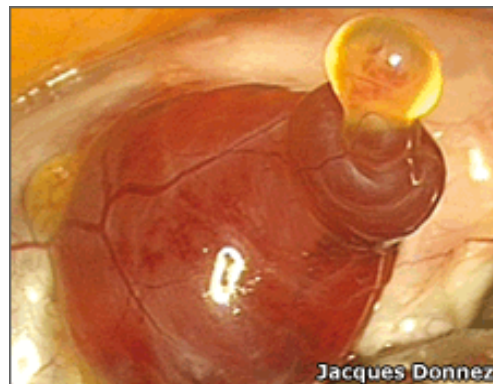
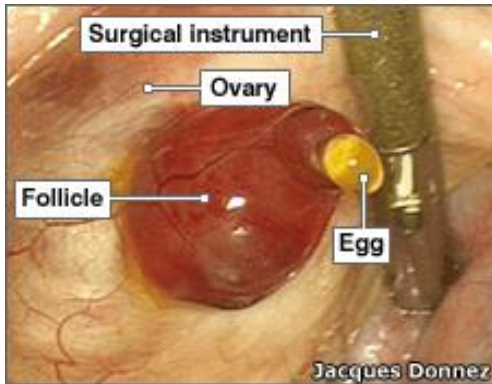
Technique



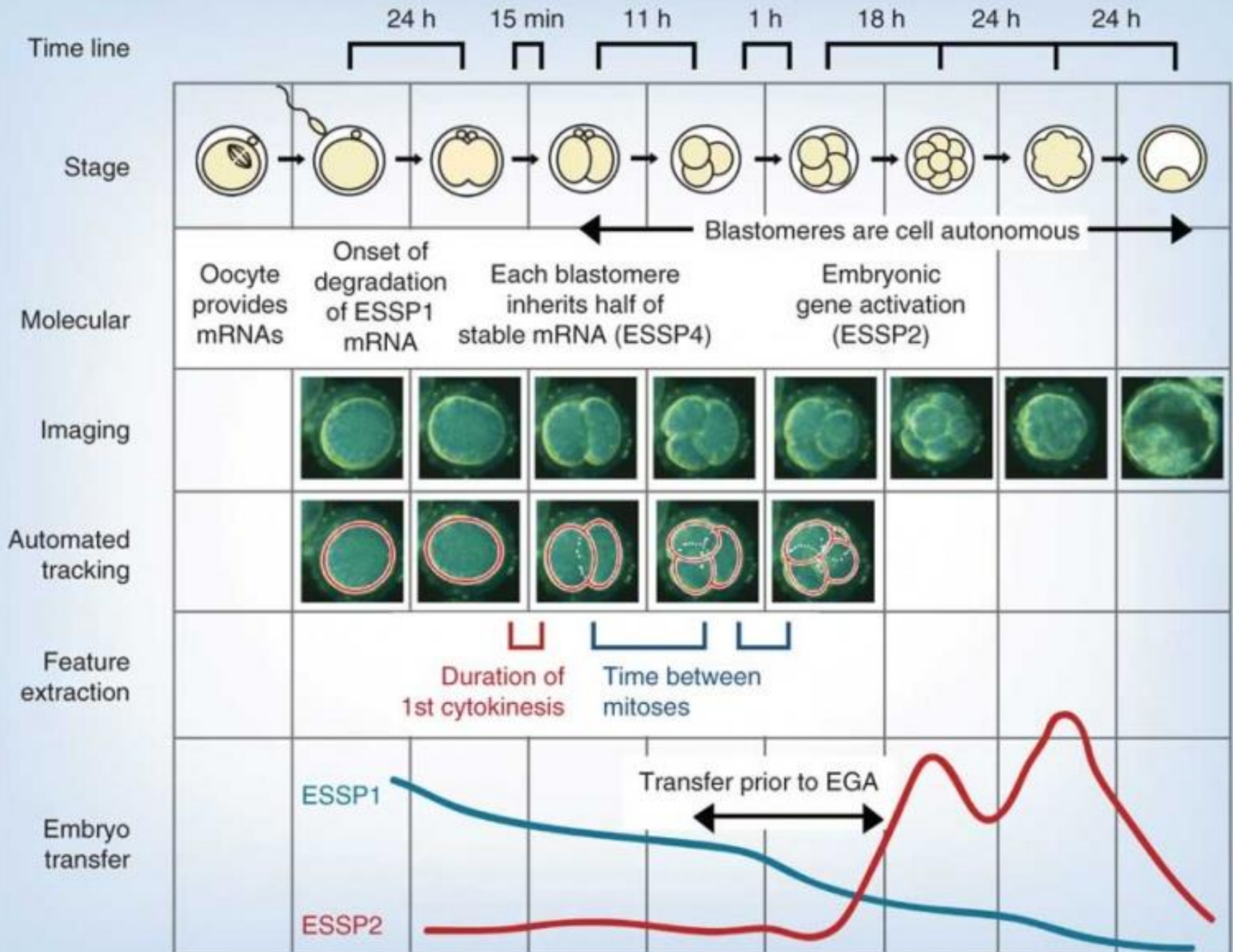
Results



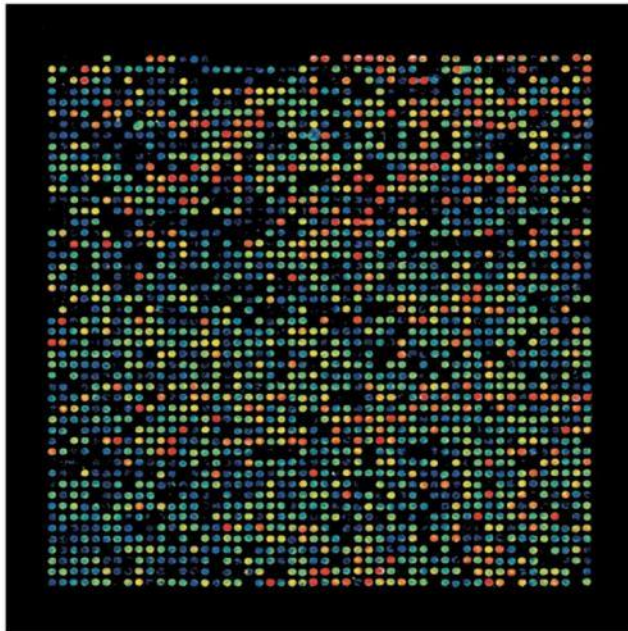
Human Ovulation



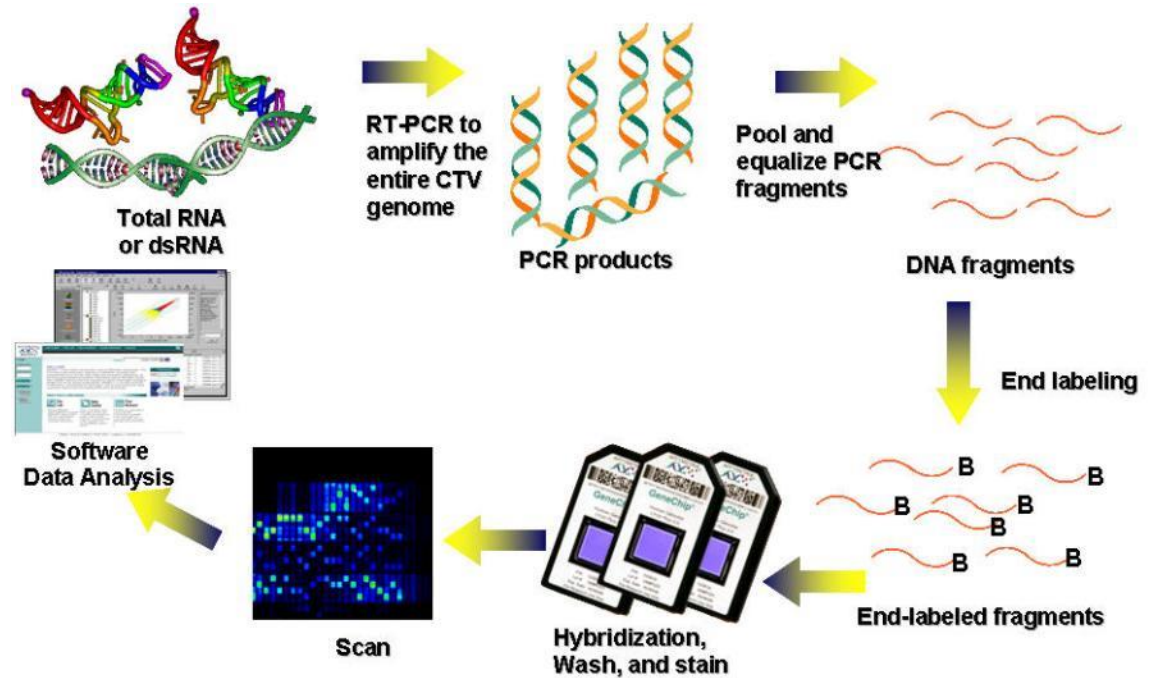
During a hysterectomy of a 45-year-old Belgian woman
Dr. Jacques Donnez, Catholic University of Louvain, Belgium



An example of DNA technology is the **microarray**, a measurement of gene expression of tens of thousands of different genes

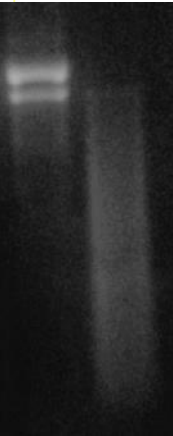


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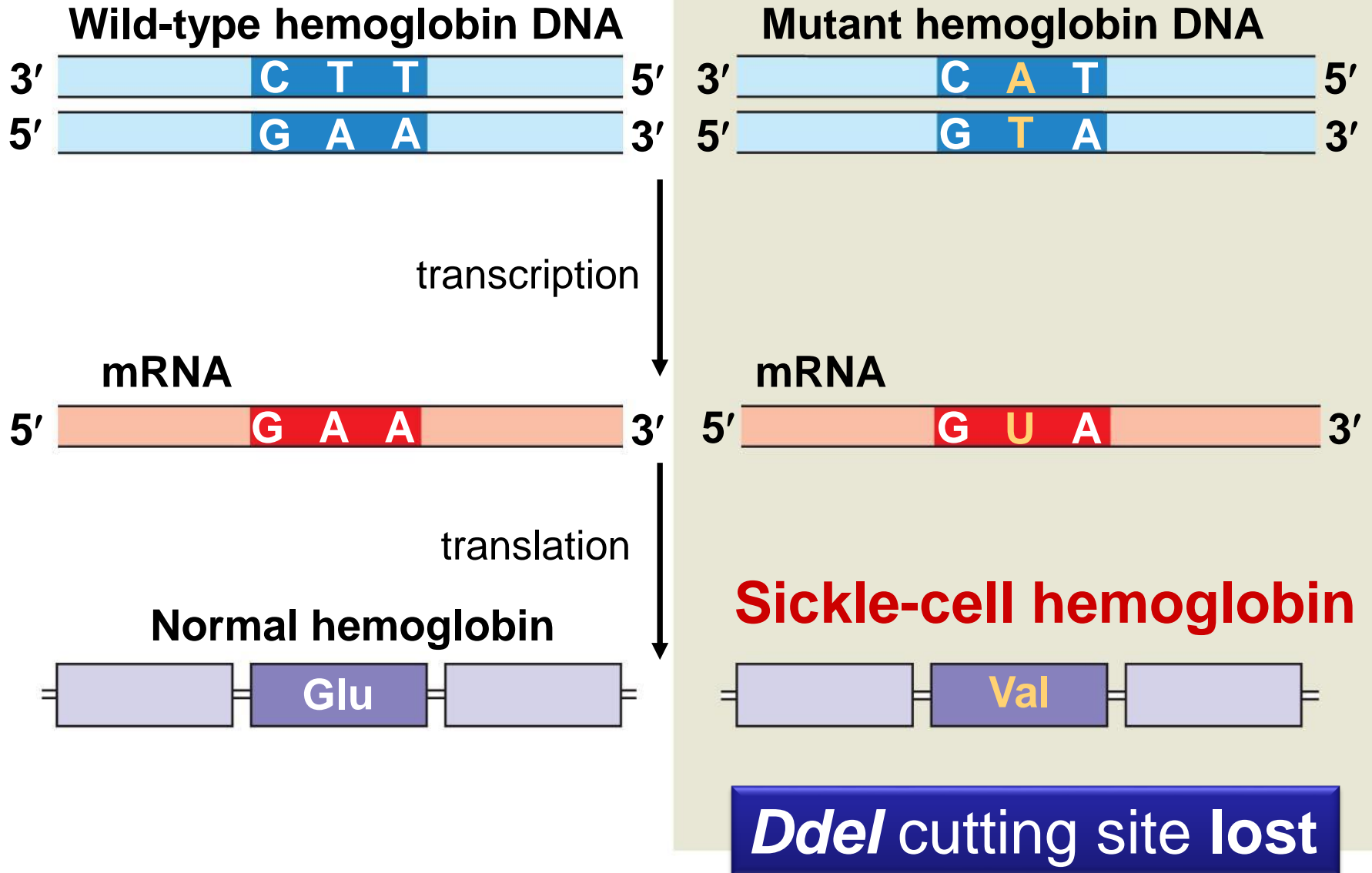


Restriction fragment analysis (RE digestion diagnostics)

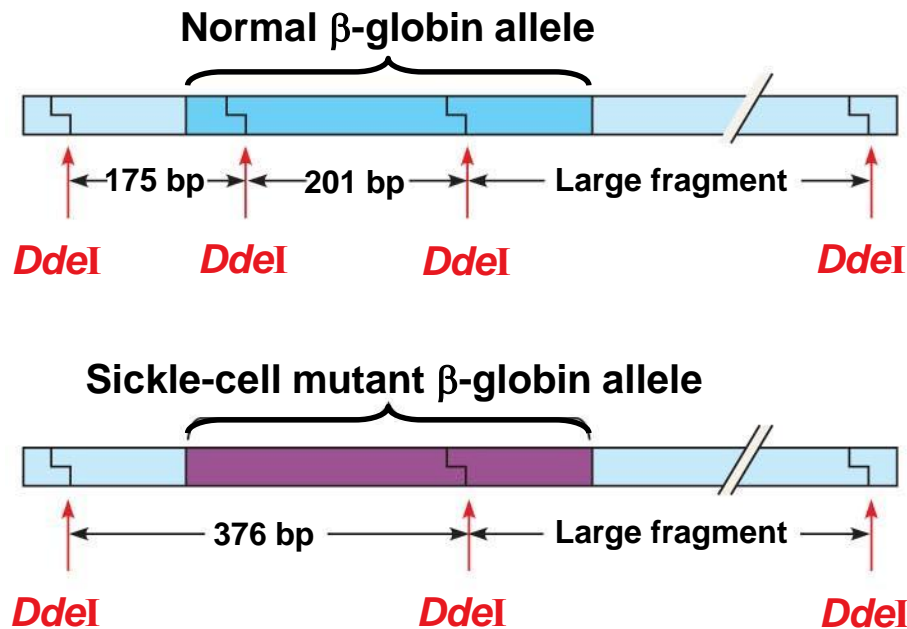
- In *restriction fragment analysis*, DNA fragments produced by **restriction enzyme** digestion of a DNA molecule are sorted by gel electrophoresis
 - A restriction enzyme will usually make many cuts in a DNA molecule
 - Yielding a set of **restriction fragments**
 - Gel electrophoresis to Separates DNA restriction fragments of different lengths:
 - Small DNA molecules from virus, plasmid – **discrete bands**
 - Large DNA molecules from eukaryotic chromosome --- **smear**
- Restriction fragment analysis is useful for comparing two different DNA molecules, such as two alleles for a gene – **restriction fragment length polymorphism (RFLP)**



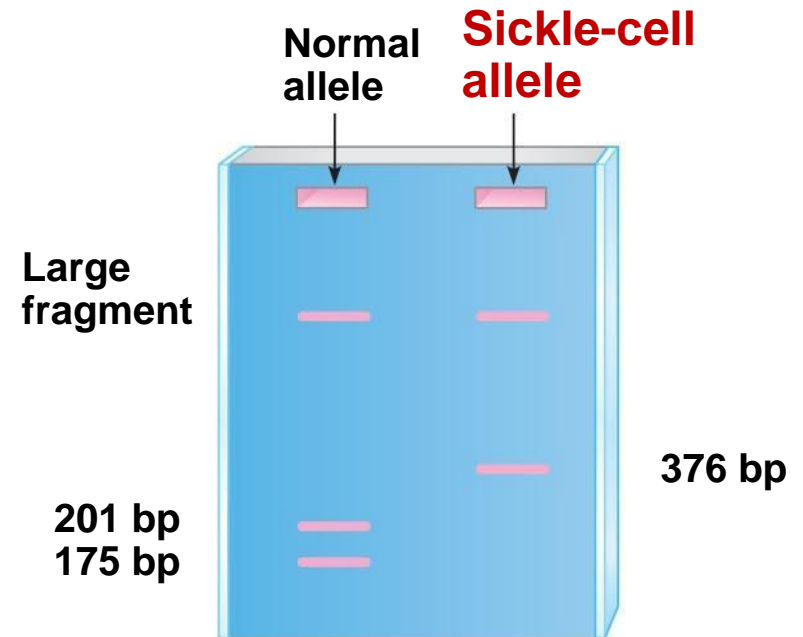
The molecular basis of sickle-cell disease: a point mutation



Using restriction fragment analysis to distinguish the normal and sickle-cell alleles of the β -globin gene



(a) *DdeI* restriction sites in normal and sickle-cell alleles of β -globin gene



(b) Electrophoresis of restriction fragments from normal and sickle-cell alleles

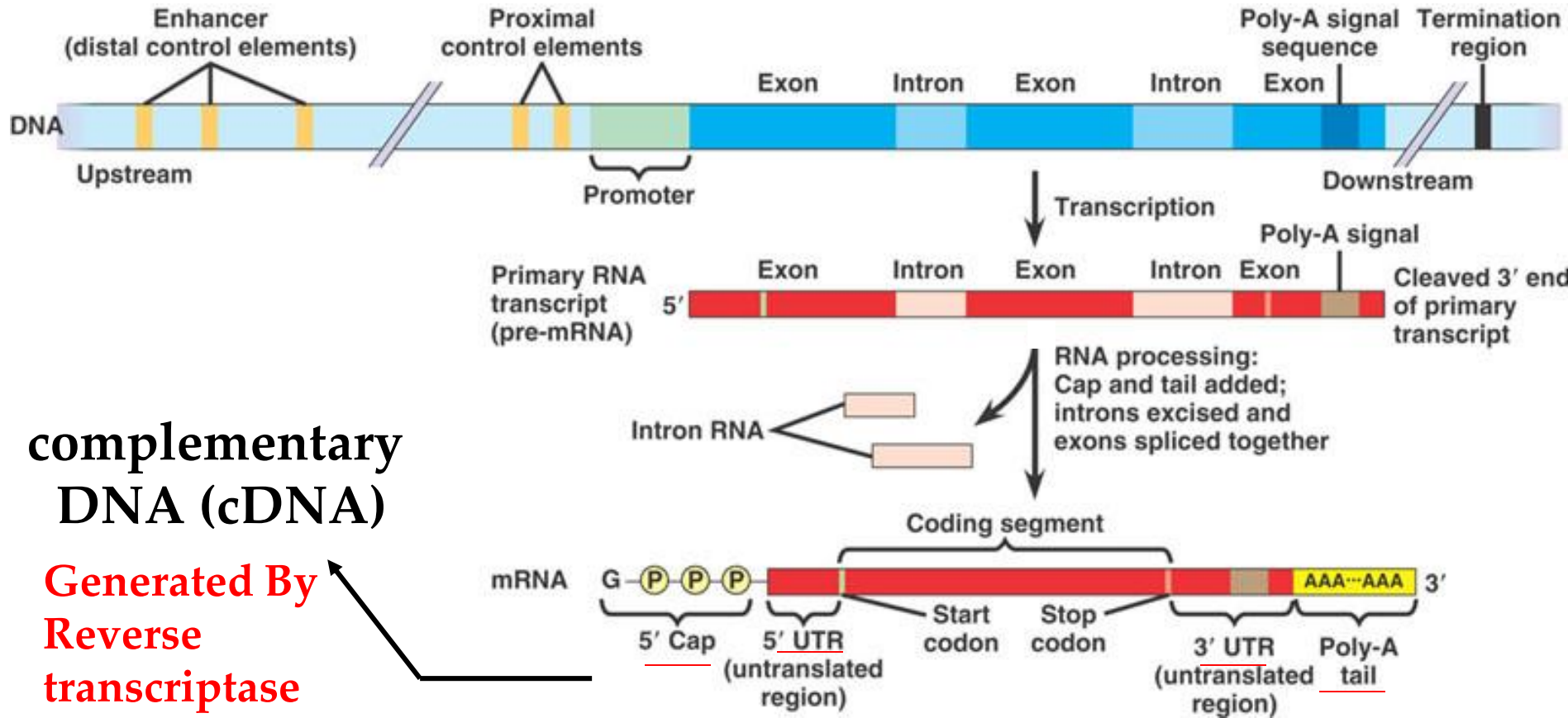
Molecular Diagnosis (分子診断)

Storing Genetic information in DNA Libraries

- A **genomic library** that is made using bacteria is the collection of recombinant vector clones produced by cloning DNA fragments from “**an entire genome**” - complete sequence with regulatory region (enhancers), exon, introns, noncoding region, etc.
 - A genomic library that is made using bacteriophages is stored as a **collection of phage clones**
-

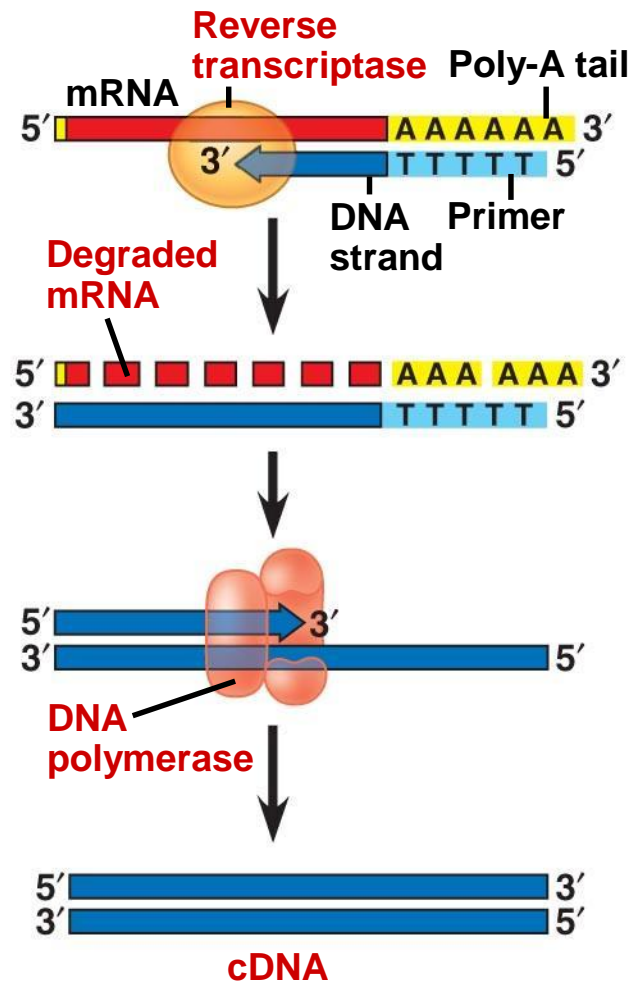
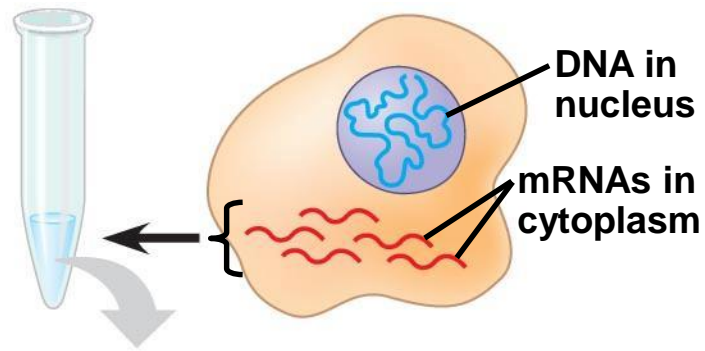
Organization of eukaryotic genes: with multiple control elements

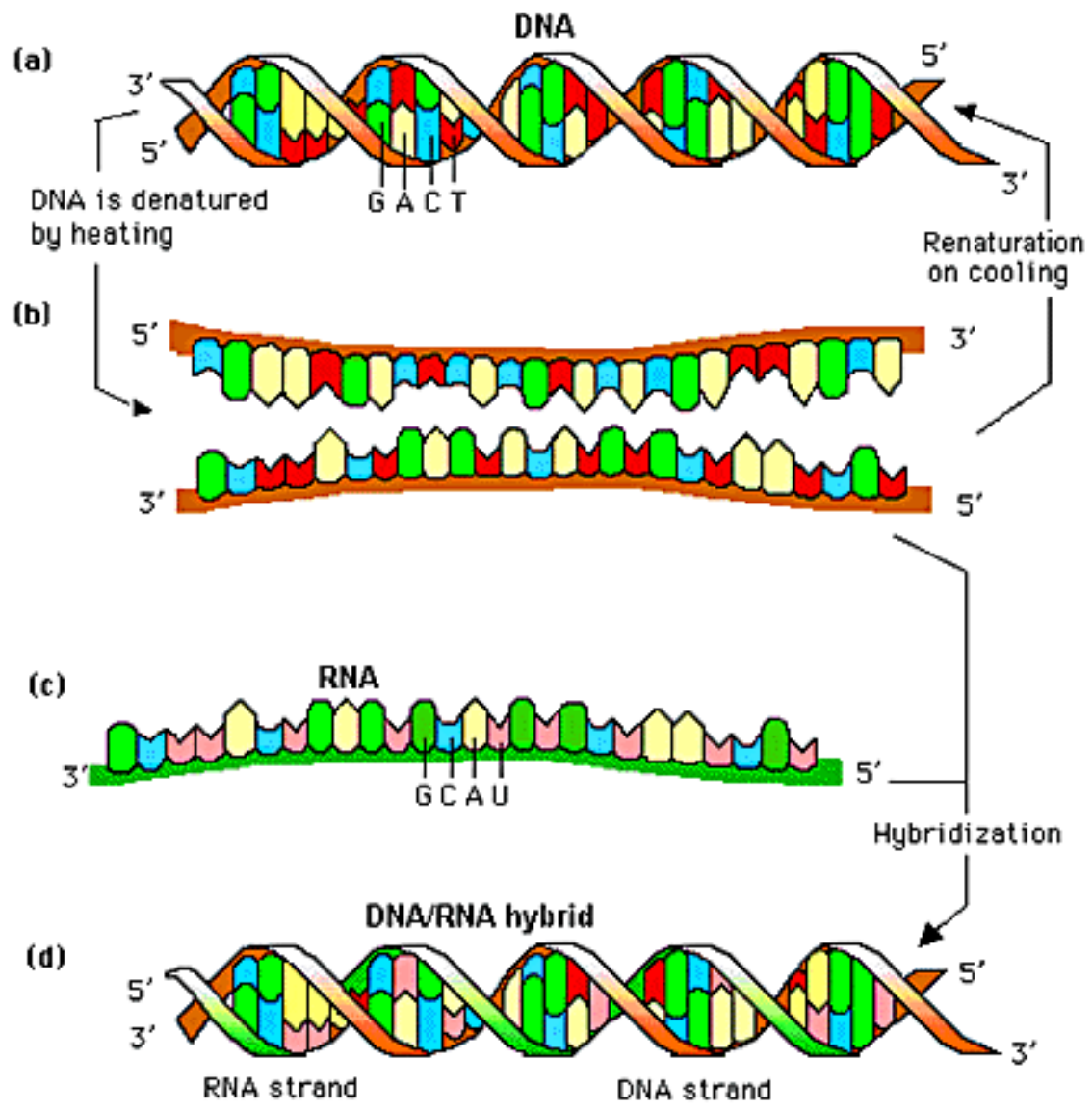
noncoding control elements coding region



cDNA library (from mRNAs, 只含已被表現的基因)

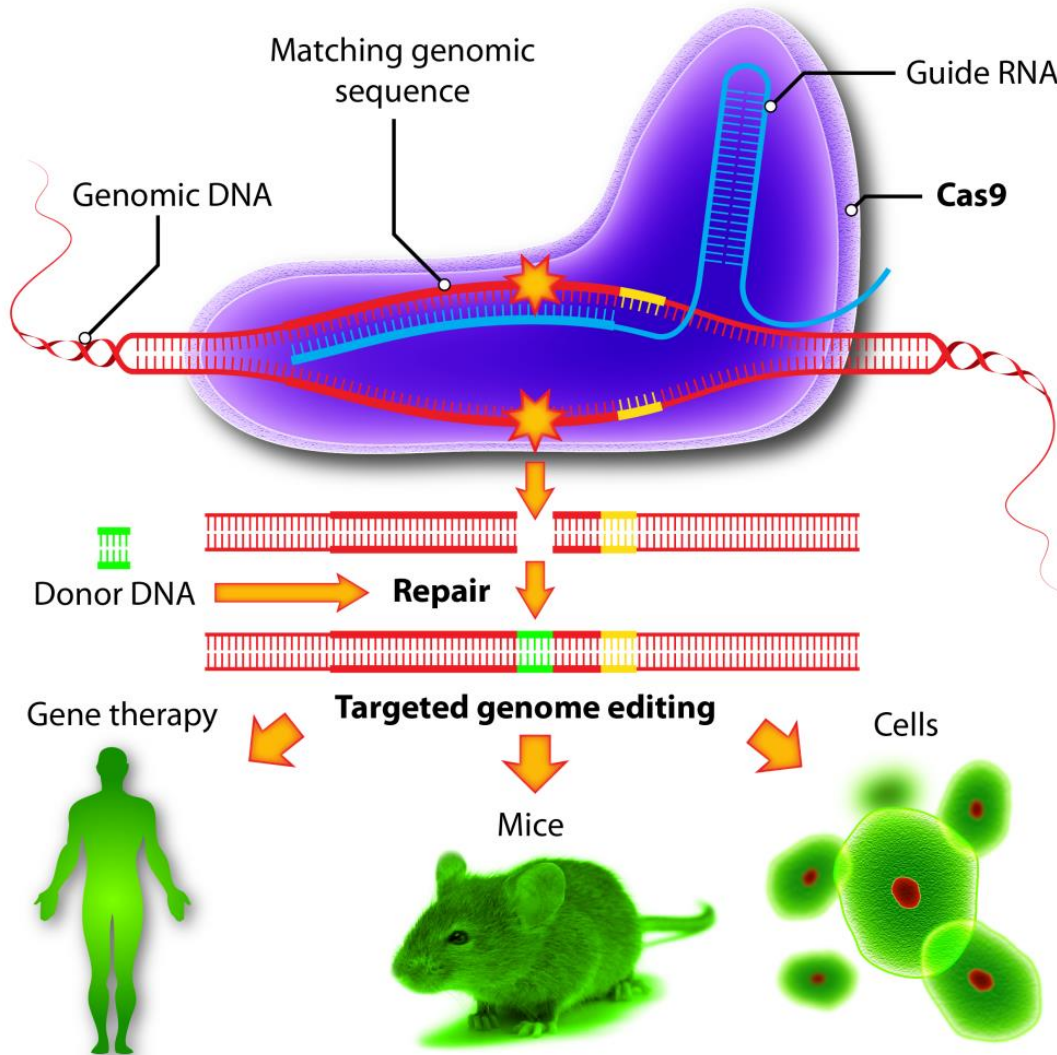
- A **complementary DNA (cDNA)** library is made by cloning DNA made *in vitro* by **reverse transcription of all the mRNA** produced by a particular cell or tissue
 - A **cDNA library** represents only part of the genome — **only the subset of genes** transcribed into mRNA in the original cells
-






Nucleic Acid Hybridization

Genome Editing Technology – CRISPR/Cas9 system



- **CRISPR/Cas9 system** has sparked a revolution in the field of genetic manipulation. Its applications includes targeted gene mutation, transgenic animals/plants production, endogenous gene labeling, and gene therapy.
- CRISPR/Cas system is a form of acquired immune system used by many bacteria and archaea species to provide resistance to viruses and other foreign genetic material.

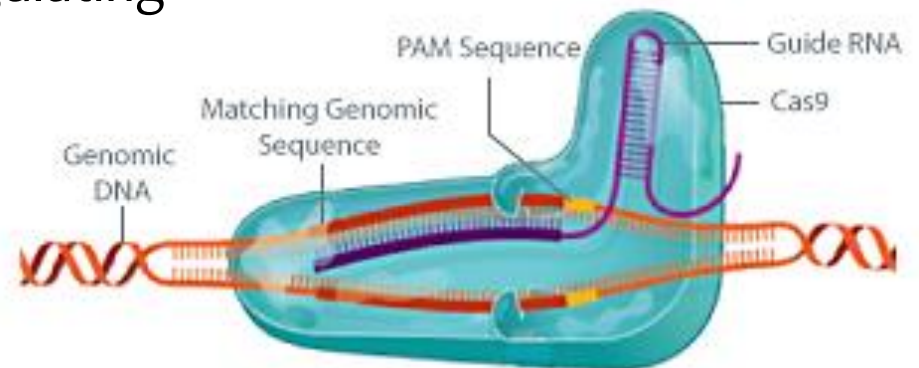


CLUSTERED REGULARLY INTERSPACED SHORT
PALINDROMIC REPEATS (CRISPRS) AND CRISPR-
ASSOCIATED (CAS) PROTEINS

CRISPR/CAS9 SYSTEM

■ CRISPR/Cas9 System

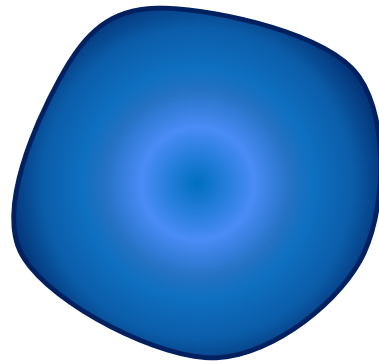
- ❖ derived from a natural process found in bacteria to protect themselves from pathogens
- ❖ targets genes for editing and regulating
- ❖ comparable to Photoshop



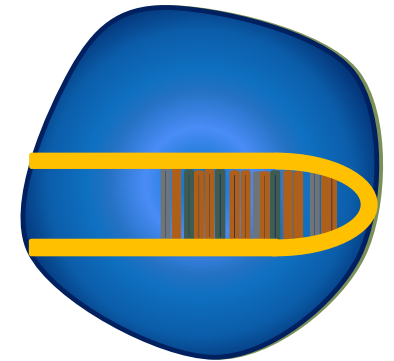
■ CRISPR/Cas9 System



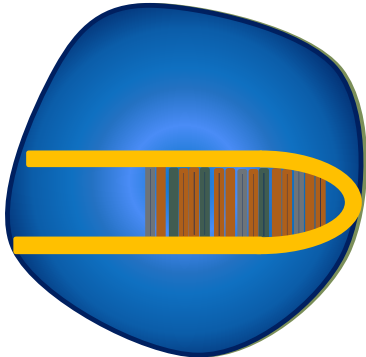
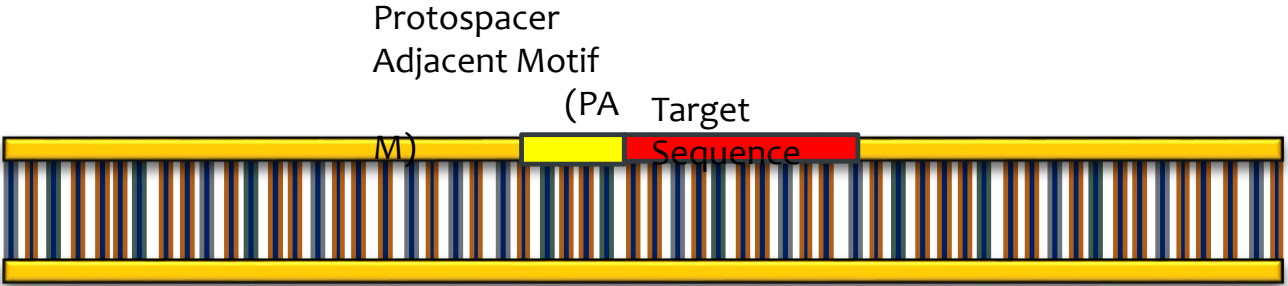
sgRNA
(single guide RNA)



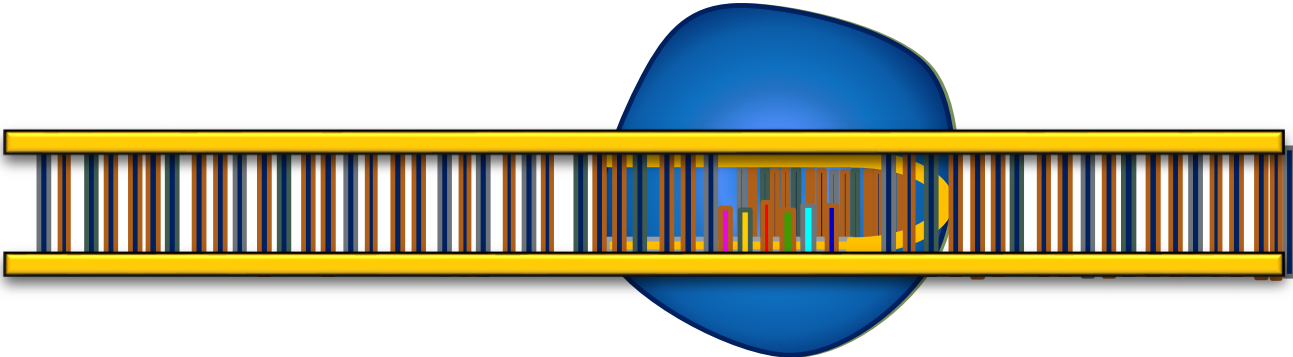
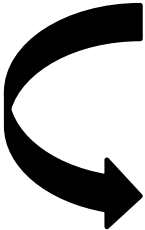
Cas9 nuclease

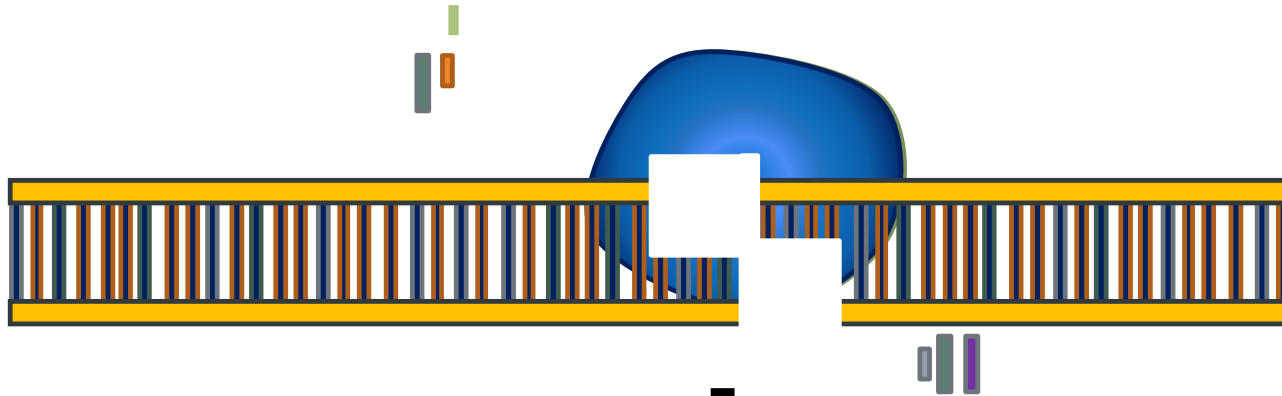


Cas9 complex



Gene of Interest



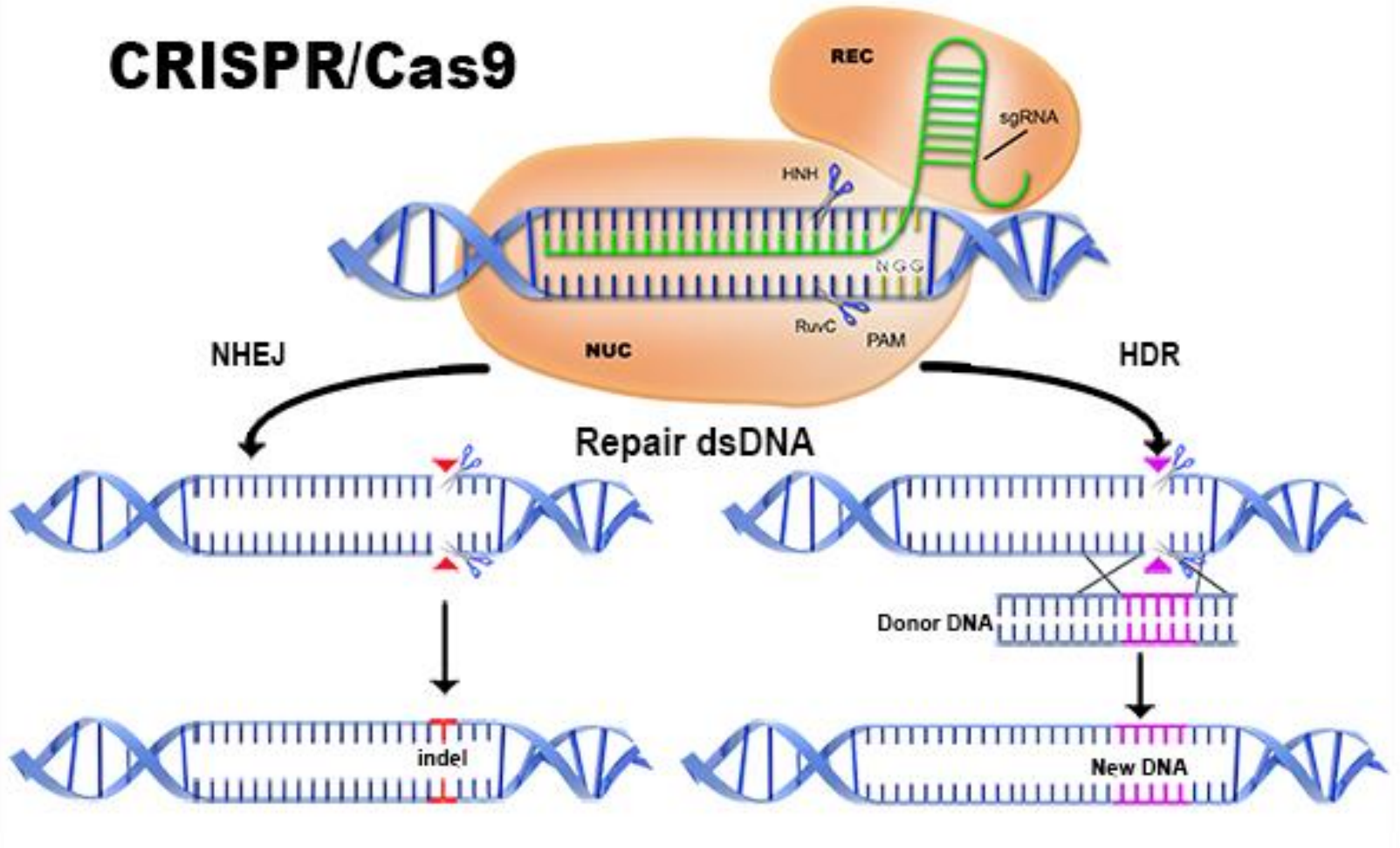


**Non-Homologous End
Joining (NHEJ) DNA
repair pathway**



Stop Codon

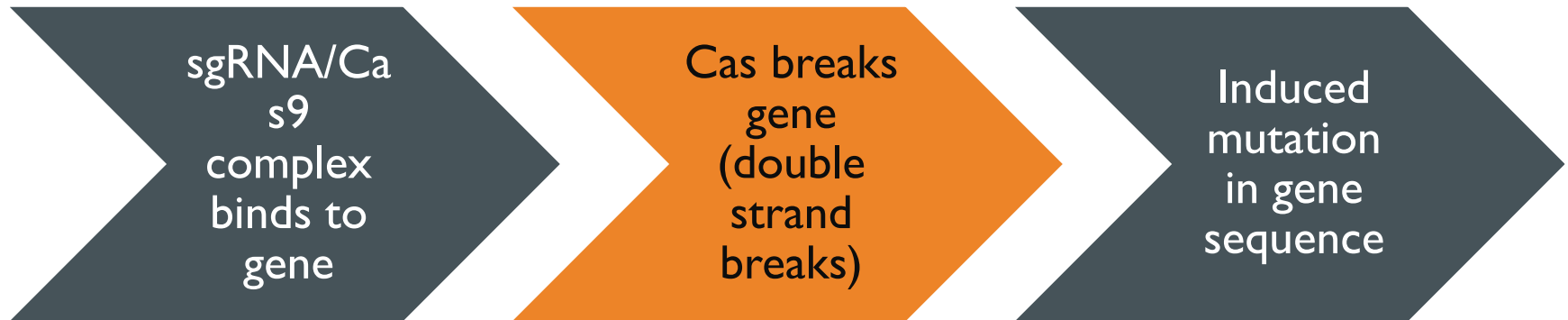
CRISPR/Cas9



Non-homologous end joining (NHEJ)

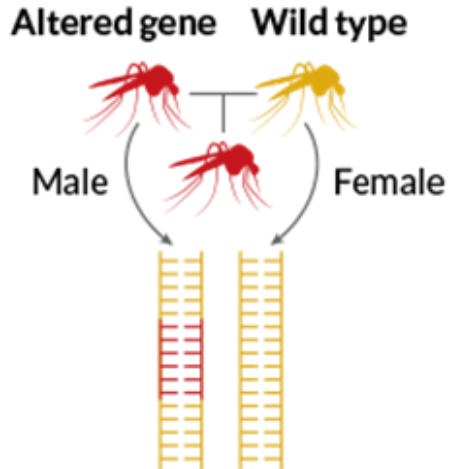
Homology directed repair (HDR)

SUMMATION: CRISPR/CAS SYSTEM

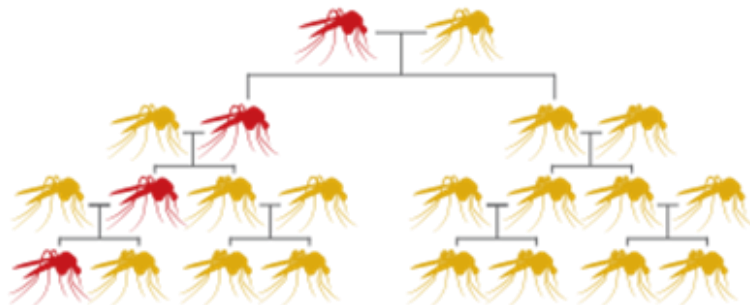


altered gene sequence → dysfunctional CAD gene

Normal inheritance

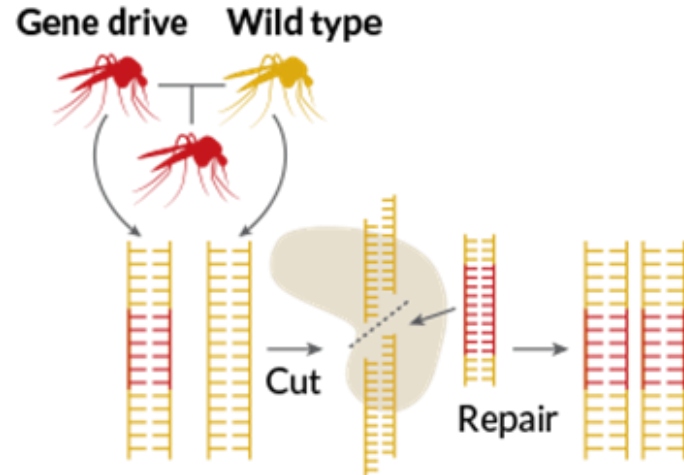


Altered gene without gene drive: One copy inherited from one parent. 50 percent chance of passing it on.

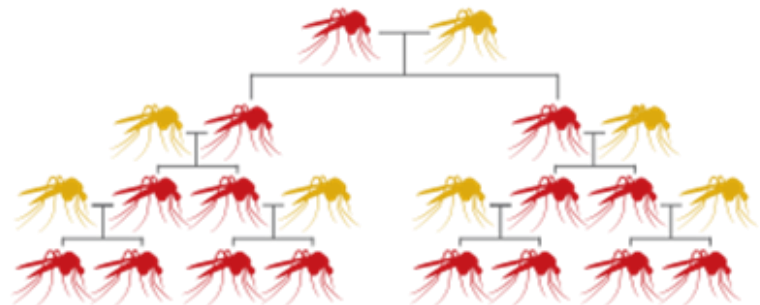


Altered gene does not spread

Gene drive inheritance



Altered gene as gene drive: One copy converts gene inherited from other parent. More than 50 percent chance of passing it on.



Altered gene is almost always inherited