

#### 4. Recombination

- Recombination is essential for generating genetic diversity and for maintaining genome integrity (完整性).
- DNA recombination is the exchange of DNA strands to produce new nucleotide sequence arrangements.

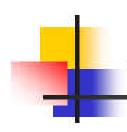
DNA重组是指由于DNA链的交换形成新的 核苷酸排列方式(新的DNA分子)的过程。



## Homologous recombination (同源重组)

Site-specific recombination (位点特异性重组)

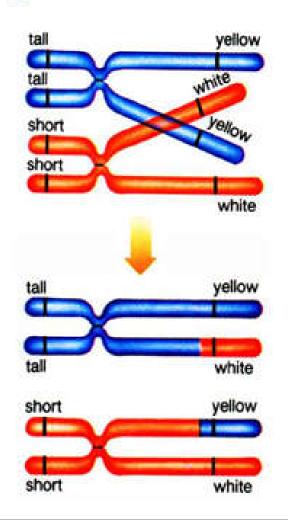
Transposition (转座)

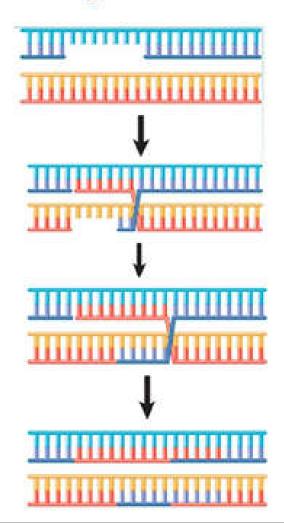


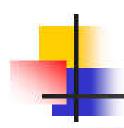
### 4.1 Homologous recombination (HR)

 Homologous recombination is the exchange of DNA strands of similar or identical (homologous) nucleotide sequence, also known as general recombination.

同源重组是相同或相似(同源)序列的 DNA片段之间的交换,也称为一般重组。 • HR is essential in meiosis (减数分裂) for generating diversity and for chromosome segregation, and in mitosis to repair DNA damage.







#### 4.1.1 Conditions for HR

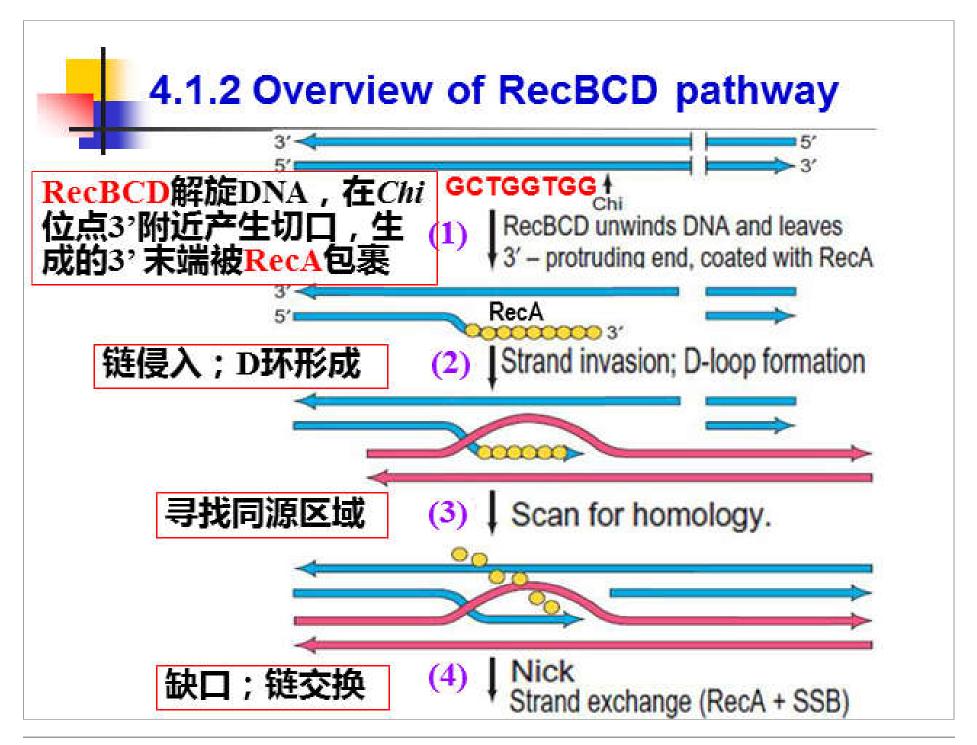
- (1) Homologous DNA sequences
- (2) Synapsis (联会)

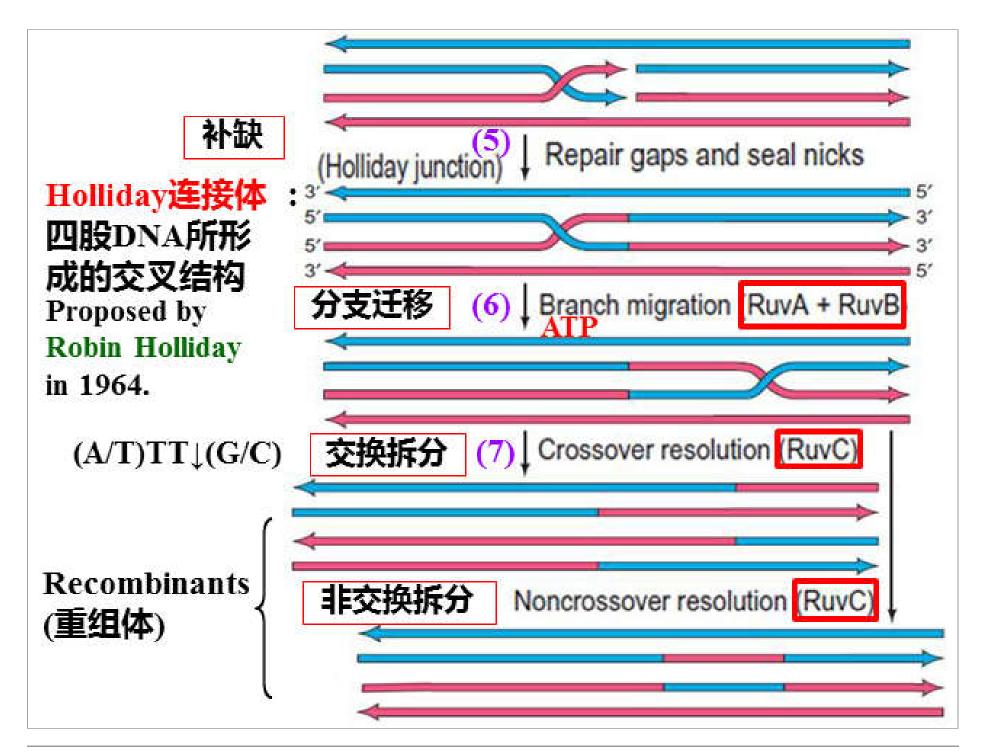
任意两个同源DNA片段在细胞内均可能重组,但是要做到这一点,两个同源双链DNA分子必须靠近,相对应的顺序方能交换,即发生联会。

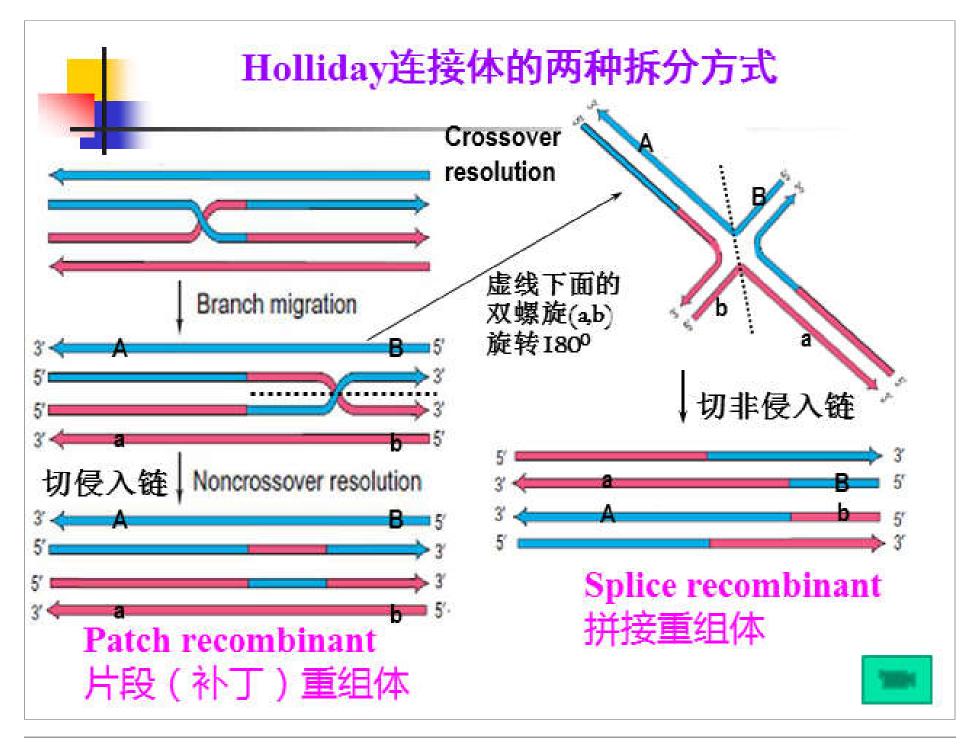
(3) Nick or gap

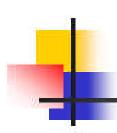
重组时,两个DNA片段必须至少有一个发生断裂(形成nick)或有一段单链缺失(形成gap)。因此许多DNA损伤因素均可使重组频率增加。

(4) Recombination related proteins









#### 4.1.3 RecBCD

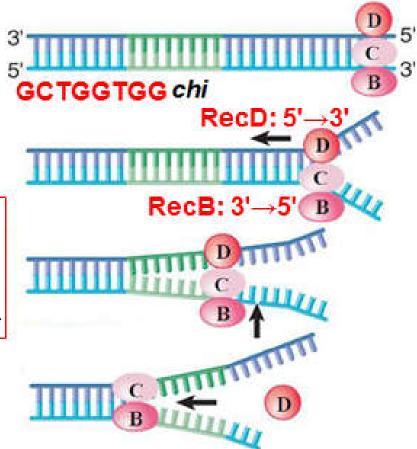
The RecBCD complex has helicase and nuclease activities.

RecBCD binds a double-strand break.

RecBCD unwinds the duplex.

RecBCD cleaves single strand at a position 4-6 bases to the right of the *du* site (重组频率较高的部位).

RecD dissociates and nuclease activity losses.



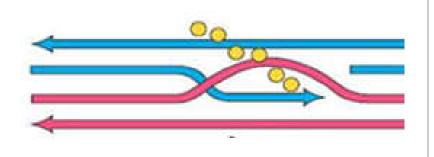
RecBCD provides a single stranded region with a free 3' end.



• In the presynapsis (联会前) step of recombination, RecA coats a single-stranded DNA that is participating in recombination.

 In synapsis, RecA promotes invasion (侵入) of another DNA duplex, forming a D-loop.

 RecA helps the invading strand scan for a region of homology in the recipient (受体) DNA duplex.

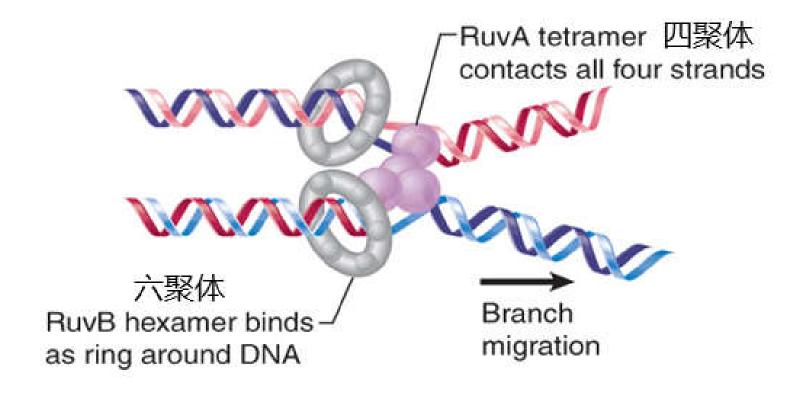


RecA



#### 4.1.5 RuvA and RuvB

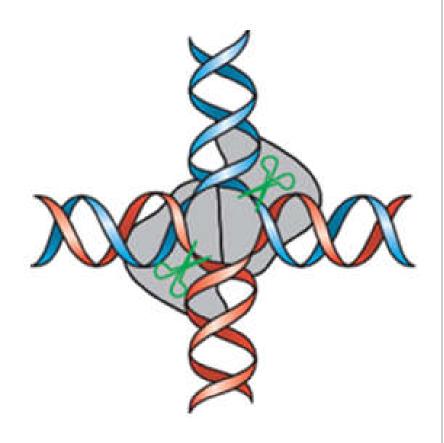
- RuvA and RuvB form a DNA helicase that can drive branch migration.
- Need ATP. RuvB is a ATPase.

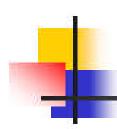




#### 4.1.6 RuvC

- Resolution of Holliday junctions.
- The clipping occurs preferentially (优先地) at the consensus sequence (A/T)TT↓(G/C).
- Branch migration is essential for efficient resolution of Holliday junctions.





### 4.2 Site-specific recombination

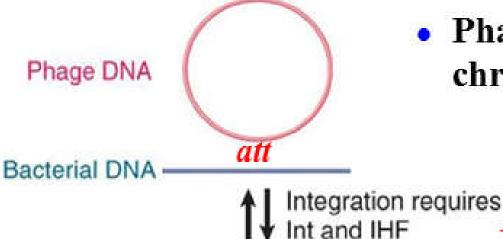
 Site-specific recombination involves the exchange of nonhomologous but specific pieces of DNA and is mediated by integrase that recognize specific DNA sequences.

位点特异性重组是由能识别特定DNA序列的整合酶介导的非同源但序列特定的DNA交换。

It does not require RecA or ssDNA.



### 4.2.1 Bacteriophage λ integration



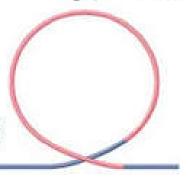
• Phage integrated in E.coli chromosome at att sites.

λ-encoded integrase (Int) 整合酶

Bacteria-encoded IHF (integration host factor) 整合宿主因子

λ噬菌体DNA通过重组 作用整合进E. coli染色 体的特异位点,成为 原噬菌体(prophage)。

λ-encoded excisionase (Xis) 切除酶



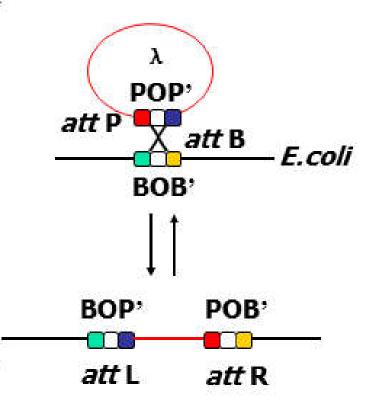
Excision requires 11 Int, Xis, and IHF

Prophage

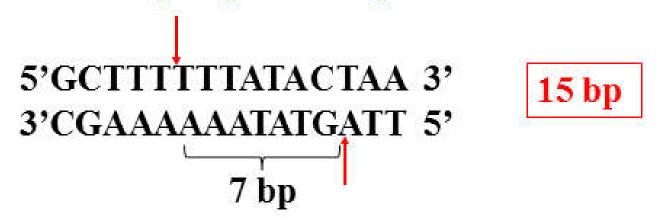


### 4.2.2 The integration/attachment (att) site

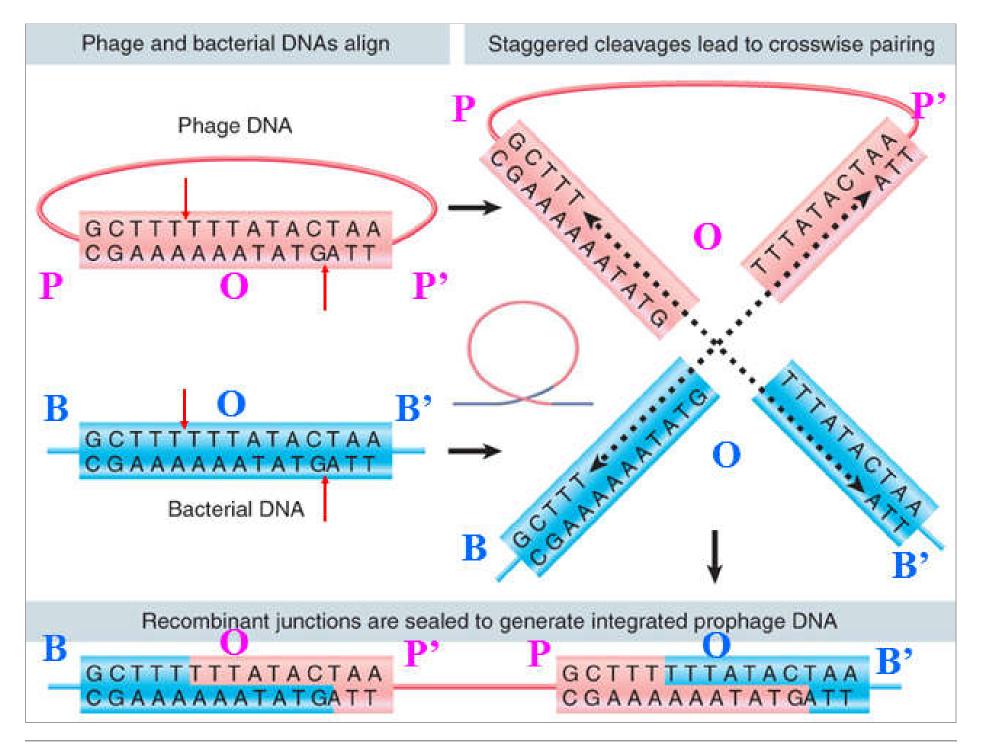
- attB由称为BOB'的序列组成, 而attP由POP'组成,O是核 心序列,是attB和attP所共 同的。
- 噬菌体DNA是环状的。重组 时被整合到细菌染色体中, 成为线性序列。原噬菌体的 两侧是两个新的杂种att位点, 左侧为attL,由BOP'组成, 而右侧为attR,由POB'组成。

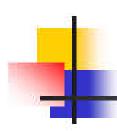


- > Core sequence the segment of DNA that is common to the attachment sites on both the λ phage and bacterial genomes.
- > It is the location of the recombination event that allows λ phage to integrate.



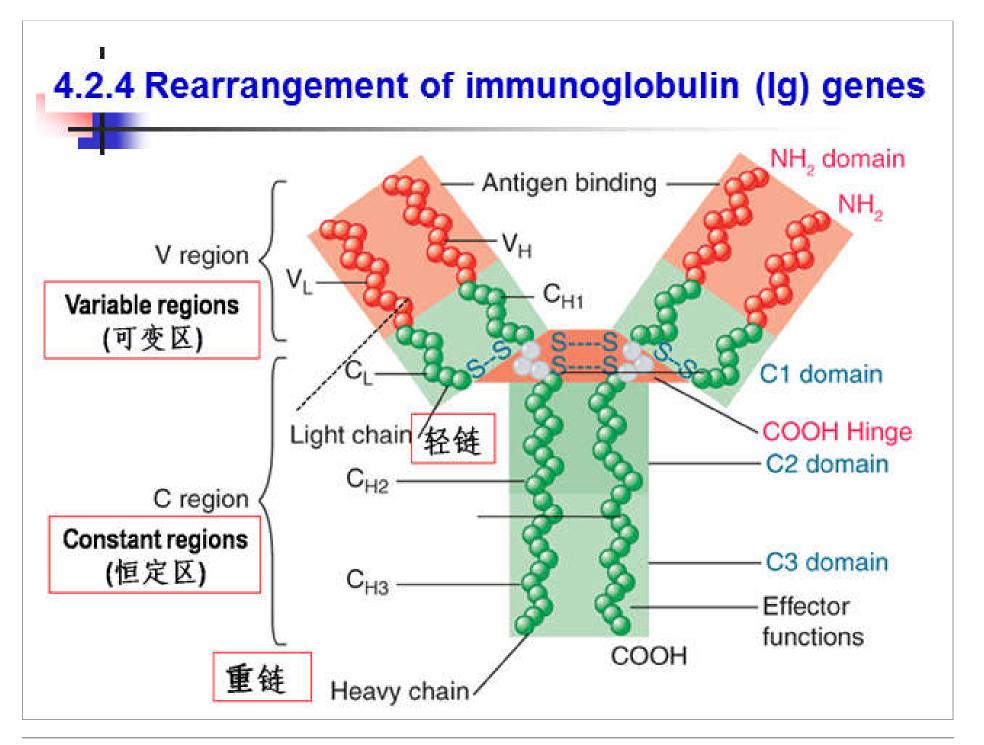
> Integrase makes staggered cuts (交错切) in the core sequence with 7 bp overhangs.





### 4.2.3 Features of λ phage integration

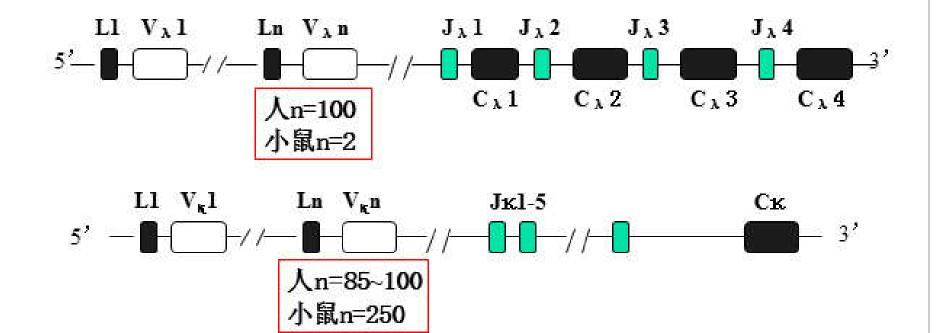
- (1) <mark>交換可逆</mark>,原先存在的DNA序列全部被保 留下来,并无丢失。
- (2) 噬菌体和细菌的DNA之间有一段很短的同源序列——核心序列(core sequence),重组交换必须通过这一段特定的核苷酸序列。
- (3) λ噬菌体编码整合酶(Int),该酶具有拓扑 异构酶性质,通过链的断裂和连接指导噬 菌体DNA插入*E. coli*染色体中。

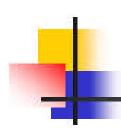




### (1) Gene structure of Ig light chain

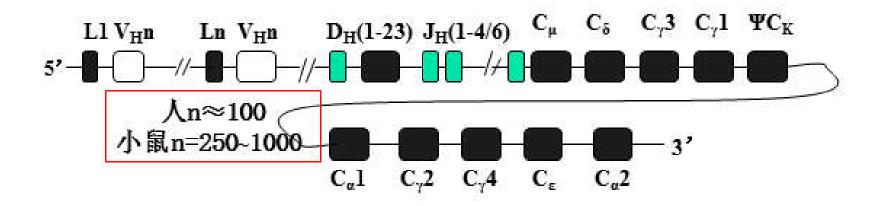
 抗体轻链有λ和κ两个基因家族,分别位于人类 22号和2号染色体上。λ和κ都由L(leader)、 V (variable)、J(joining)和C(constant)四种不同基因片段组成。



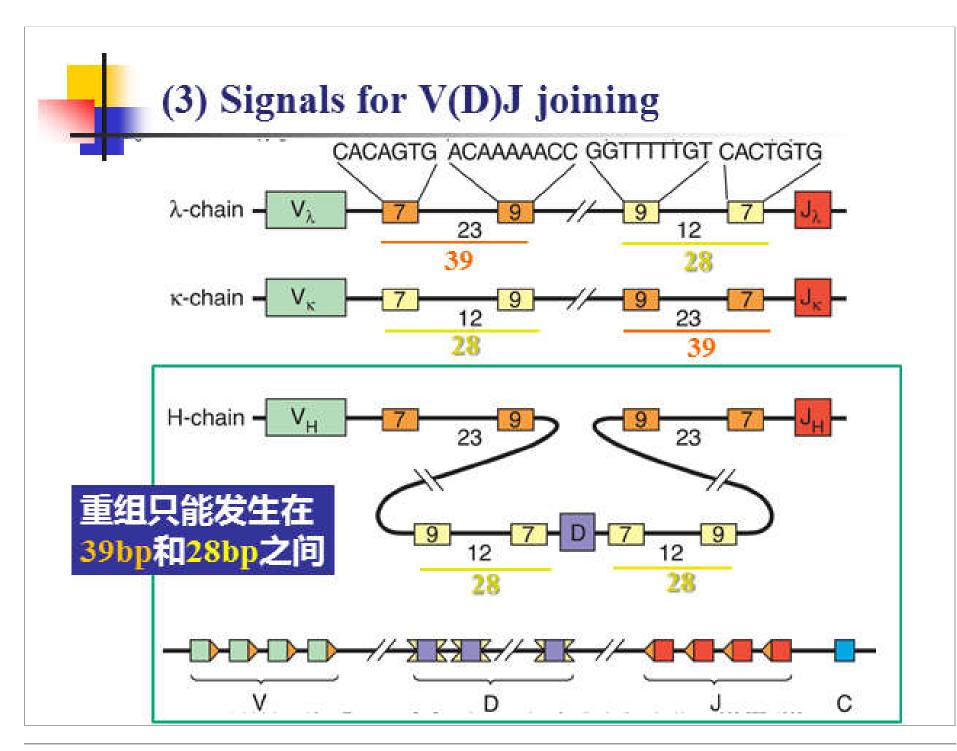


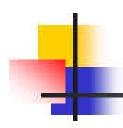
### (2) Gene structure of Ig heavy chain

 人类重链基因位于14号染色体。在重链基因中, DNA上除了有L、V、J、C片段外,在V<sub>H</sub>和J<sub>H</sub> 之间还有约20个多样性片段(D片段)。



IgG ( $\gamma$ ), IgM ( $\mu$ ), IgA ( $\alpha$ ), IgE ( $\epsilon$ ) and IgD ( $\delta$ )





### (4) Reasons for Ig diversity

V区和(D)J区不同片段在DNA分子水平上的各种排列组合是形成Ig分子多态性的根本原因。

L链: λ –V100 J4 , 有400种组合 , κ-V100 J5 , 有500种组合。

H链: V100 D20 J6,有100×20×6 = 1.2×10<sup>4</sup> H和L组合(400+500)×1.2×10<sup>4</sup> =1.08×10<sup>7</sup>



体细胞超突变:免疫细胞增殖过程中可变区的突变频率比一般细胞高出10<sup>4</sup>-10<sup>5</sup>
 倍。突变发生的部位大都在V-D-J外显子内或其附近。



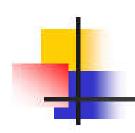
### 4.3 Transposition



Barbara McClintock (1902-1992) 20世纪40年代发现 1983 Nobel Prize in Physiology or Medicine

 In transposition, a transposon (转座子), or transposable element (转座元件, 转位因子), moves from one DNA address to another.  A transposon is a small DNA sequence that can move to virtually any position in a cell's genome.
 转座子是能够转移进细胞基因组内的几乎任何位置的短DNA序列。

- Requires no homology between sequences nor site-specific.
- Transposition has also been called illegitimate recombination (异常重组).
- Inefficient



Insertion sequence (IS元件, 插入序列)

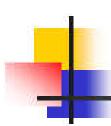
Composite transposon (复合型转座子, Tn)

Retrotransposon (反转录转座子) Mainly

Bacterial

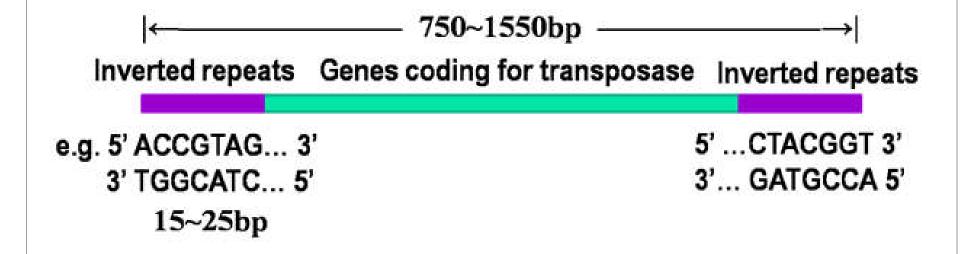
transposons

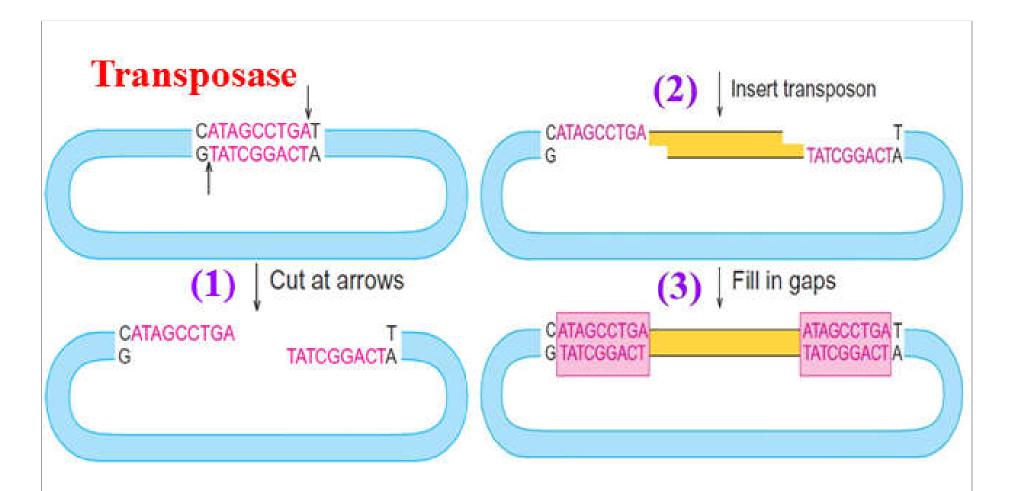
Eukaryotic transposons



### 4.3.1 Insertion sequences (IS elements)

- IS elements are the simplest transposons.
- Comprise transposase (转座酶) genes flanked by a short inverted terminal repeats (末端反向重复序列).





 The target site at which an insertion sequence is inserted is duplicated during the insertion process to form direct repeats at the ends of the transposon. 转座子插入处生成了宿主DNA的正向重复。

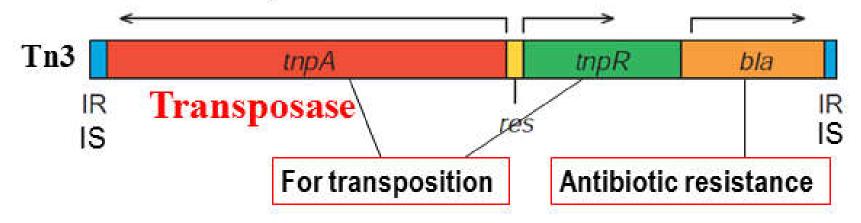


### 4.3.2 Composite transposon (Tn)

#### (1) Structure of Tn

### 复合型转座子

- Elements necessary for transposition
- Other genes (often conferring antibiotic resistance)

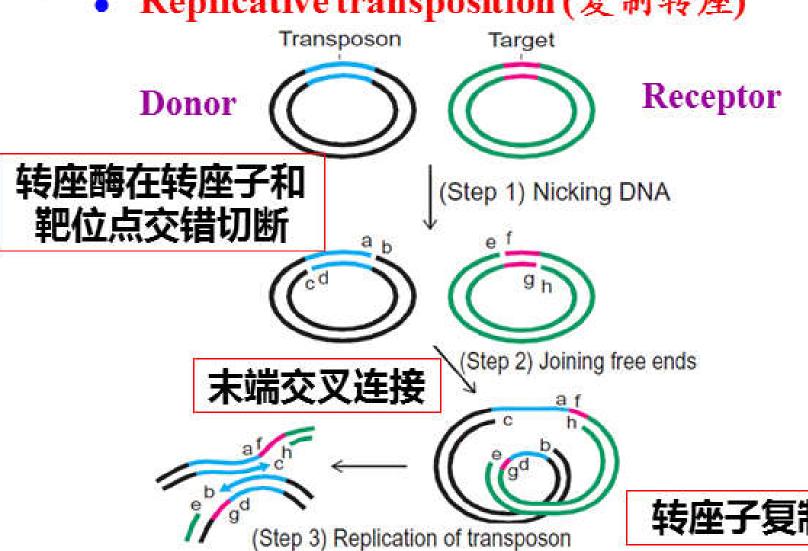


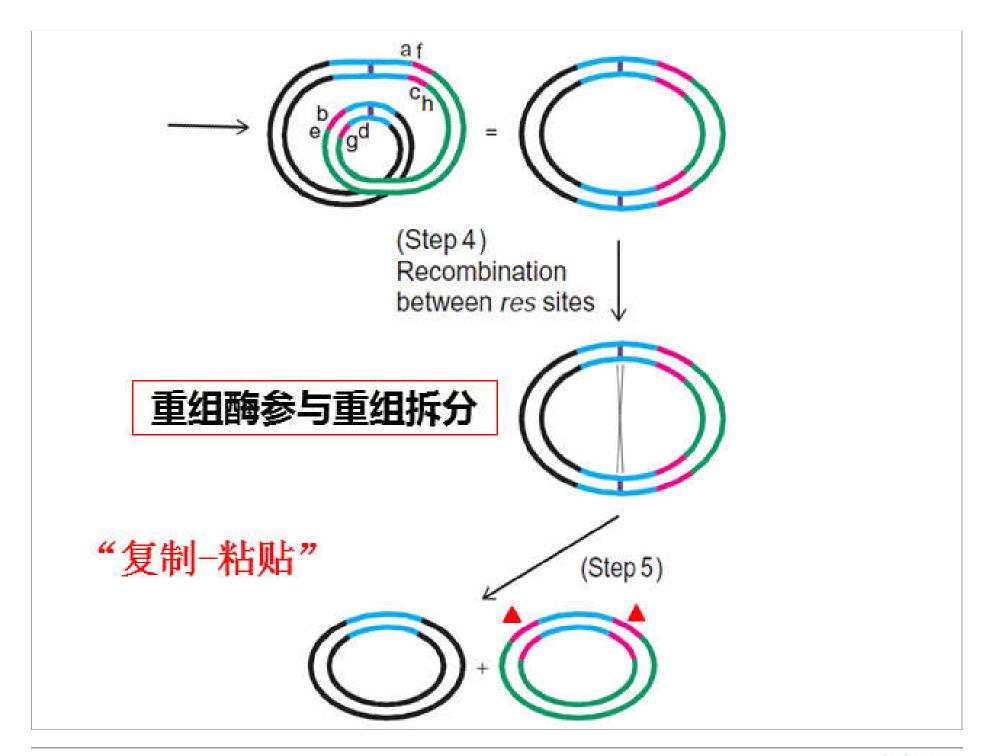
bla encodes β-lactamase (内酰胺酶), which protects bacteria against the antibiotic ampicillin (青霉素). This gene is also called *Amp*.



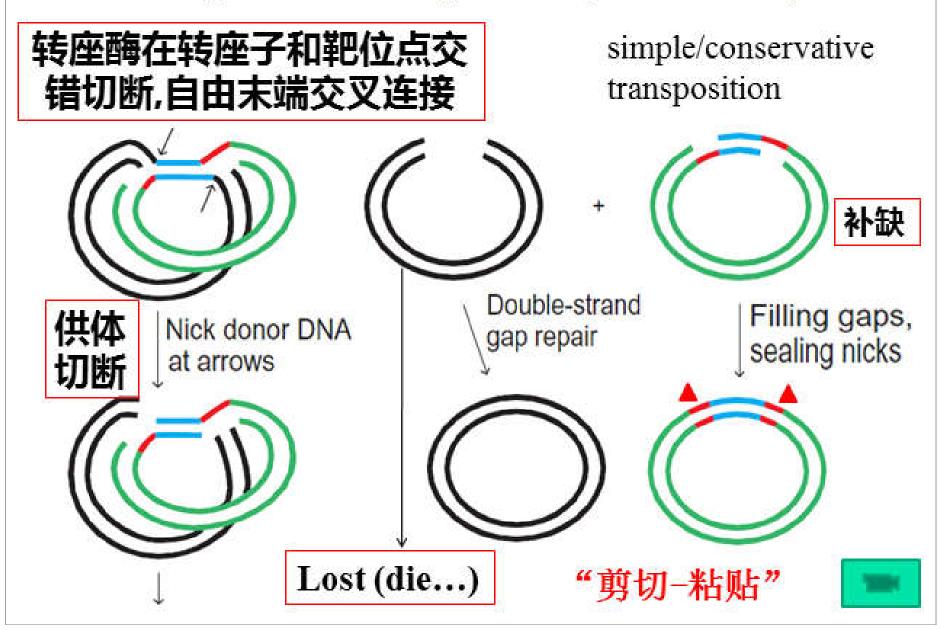
#### (2) Mechanisms of transposition

Replicative transposition (复制转座)





### • Nonreplicative transposition (非复制转座)

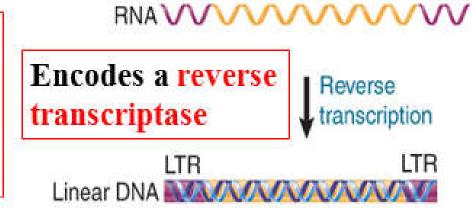




### 4.3.3 Retrotransposon

(1) LTR-containing retrotransposon

LTR (long terminal repeats, 长末端重复序列) is crucial for replication of most retrotransposons.



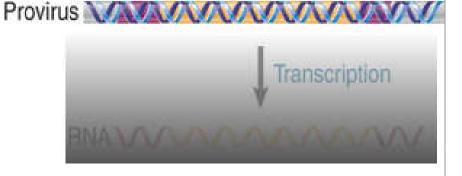
逆转录病毒(本身可看作转座子)

Eukaryotic transposons, many are retrotransposons.

e.g. Yest Ty element

\*\*Drosophila copia element

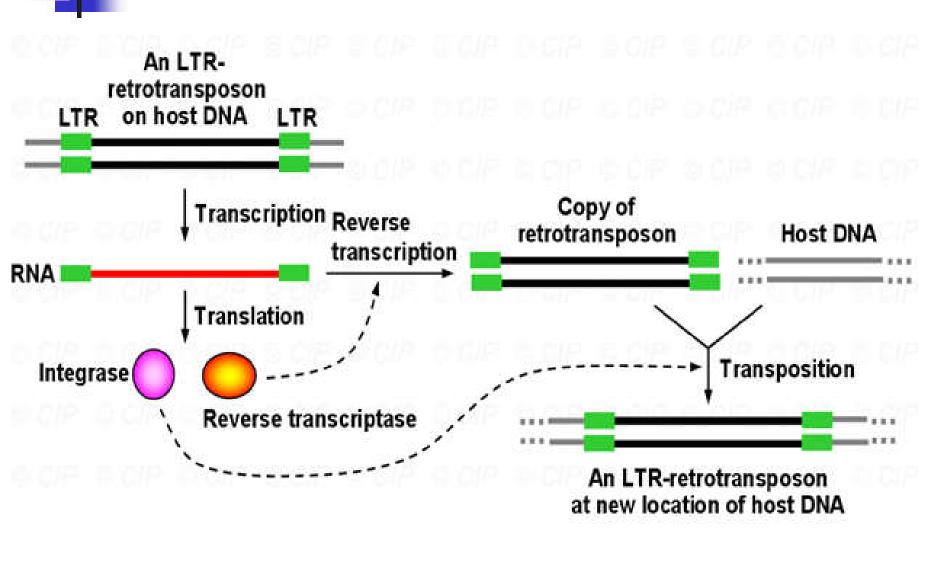
DNA→RNA→DNA



Integration



#### $DNA \rightarrow RNA \rightarrow DNA$





#### (2) LTR-free retrotransposons

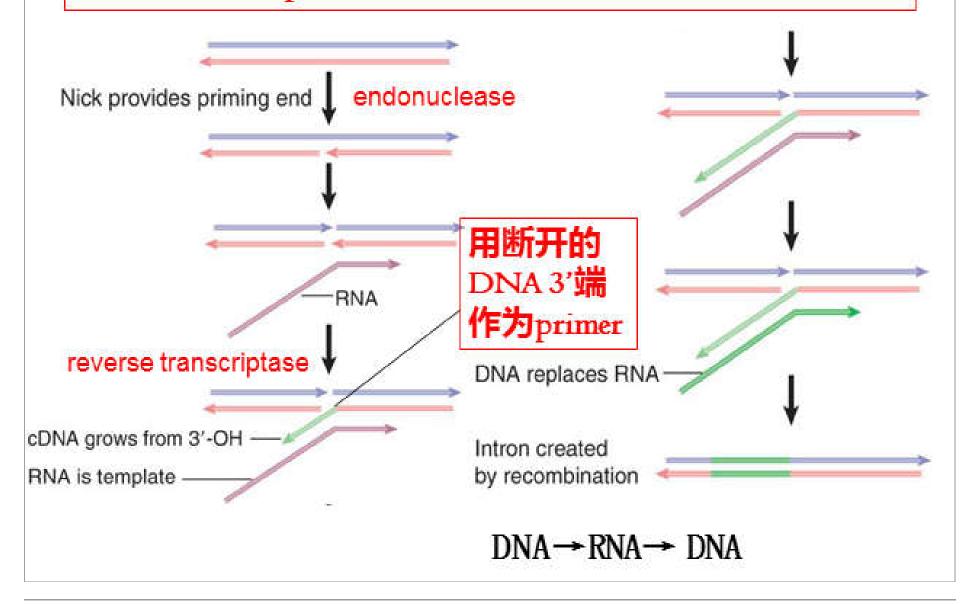
Retrotransposons that lack LTRs are much more abundant than those with LTRs, at least in mammals.

### 1 LINE

- The most abundant of LTR-free retrotransposons.
- They are autonomous (自主的) retrotransposons.

# LINEs codes for an endonuclease (内切核酸酶) and

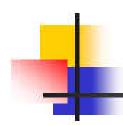
reverse transcriptase.





- nonautonomous retrotransposons.
- SINEs do not encode any proteins. Instead they take advantage of the retrotransposition machinery of other elements, such as LINEs.
- The transcripts of the *Alu* elements contain a domain that resembles (类似) the 7SL RNA component of signal recognition particle (信号识别颗粒).

7SLRNA结构域通常是帮助特定核糖体结合到粗面内质网。7SLRNA结构域帮助Alu的RNA到翻译LINEsRNA的核糖体,使Alu的RNA能和所需要的蛋白结合,从而被反转录插入到新的位点。



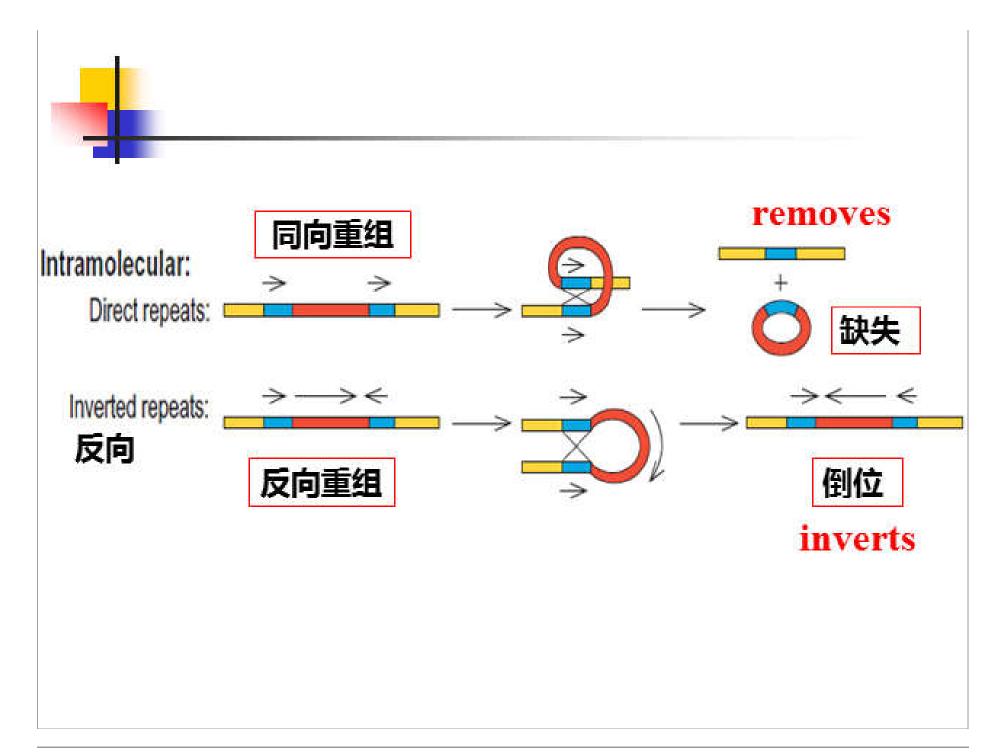
### 4.3.4 Genetic effects of transposition

### (1) Insertion mutation

The gene into which the transposon inserts is usually inactivated.

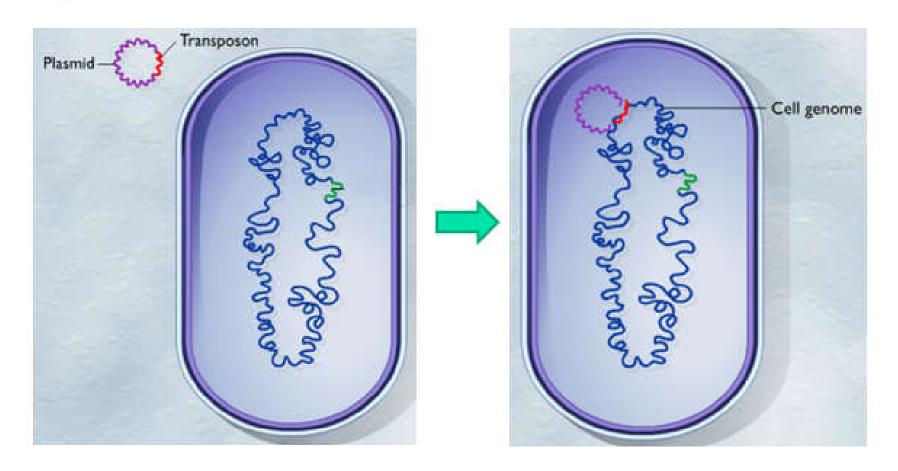
### (2) Deletion or inversion

Genes between two copies of a transposon can be deleted or inverted by recombination between them.





# (3) Introduction of new gene





# Summary

- 1. Types and phenotypic effects of mutation
- 2. Replication fidelity
- 3. Factors and mechanisms of DNA damage
- 4. Relationship between DNA damage and mutation
- 5. DNA repair mechanisms
- 6. Definition of homologous recombination, sitespecific recombination and transposition
- 7. Mechanisms of homologous recombination, site-specific recombination and transposition
- 8. Types and structural characteristics of transposons

