

## Topic C

### Dynamic DNA - Mobile DNA

Simple sequences (minisatellite)

Unique DNA (<5% in eukaryotes)

Gene family (globin)

Tandem genes (rRNA..)

### 9.3 Mobile DNA moderately repeated

Mostly in Eukaryotes

Process of moving - Transposition

"molecular parasites" **Selfish DNA**

Very slow process for elimination

## 9.3 Mobile DNA

First found in corn (1940, B. MaClintock)  
Weak similarity to bacterial elements

Transposition is in 2 modes:

DNA only  
RNA intermediate

### 9.3 The two classes of mobile elements

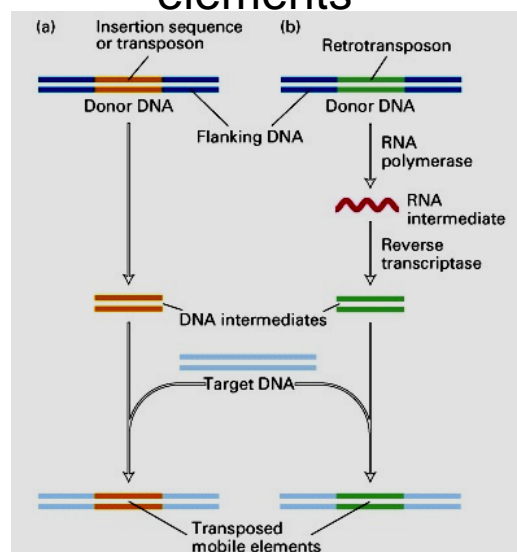


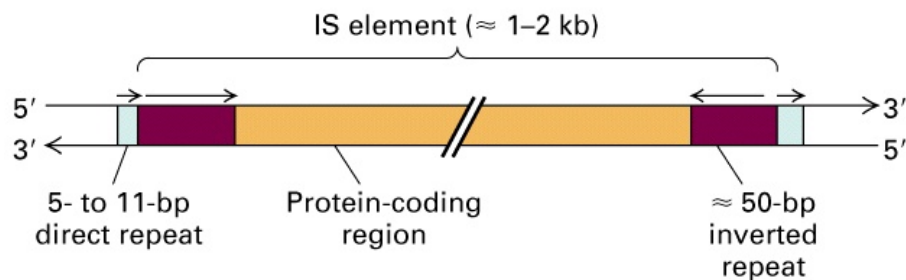
Figure 9-10

## 9.3 Mobile elements may move as

**TABLE 9-3** Major Types of Mobile DNA Elements

Type	Structural Features	Mechanism of Movement	Examples
<b>DNA-MEDIATED TRANSPOSITION</b>			
Bacterial insertion sequences (IS elements)	≈50-bp inverted repeats flanking region encoding transposase and, in some, resolvase	Excision or copying of DNA and its insertion at target site	IS1, IS10
Bacterial transposons	Central antibiotic-resistance gene flanked by IS elements	Copying of DNA and its insertion at target site	Tn9
Eukaryotic transposons	Inverted repeats flanking coding region with introns	Excision of DNA and its insertion at target site	P element ( <i>Drosophila</i> ) Ac and Ds elements (corn)
<b>RNA-MEDIATED TRANSPOSITION</b>			
Viral retrotransposons	≈250- to 600-bp direct terminal repeats (LTRs) flanking region encoding reverse transcriptase, integrase, and retroviral-like Gag protein	Transcription into RNA from promoter in left LTR by RNA polymerase II followed by reverse transcription and insertion at target site	Ty elements (yeast) <i>Copia</i> elements ( <i>Drosophila</i> )
Nonviral retrotransposons	Of variable length with a 3' A/T-rich region; full-length copy encodes a reverse transcriptase	Transcription into RNA from internal promoter; folding of transcript to provide primer for reverse transcription followed by insertion at target site	F and G elements ( <i>Drosophila</i> ) LINE and SINE elements (mammals) <i>Alu</i> sequences (humans)

## 9.3 General structure of bacterial IS elements

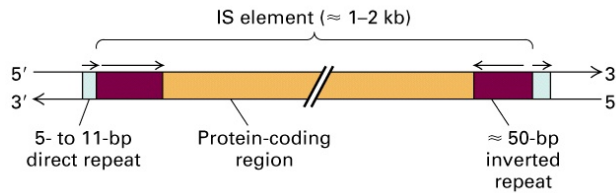


Transposase: excision and insertion

20 types in *E. coli*, seen in heteroduplex view

Figure 9-11

## 9.3 General structure of bacterial IS elements



Frequency of transposition:  
1/1000-1/10000 per generation  
In specific site -  $10^{-5}$  -  $10^{-7}$   
Reversion  $10^{-6}$  -  $10^{-10}$

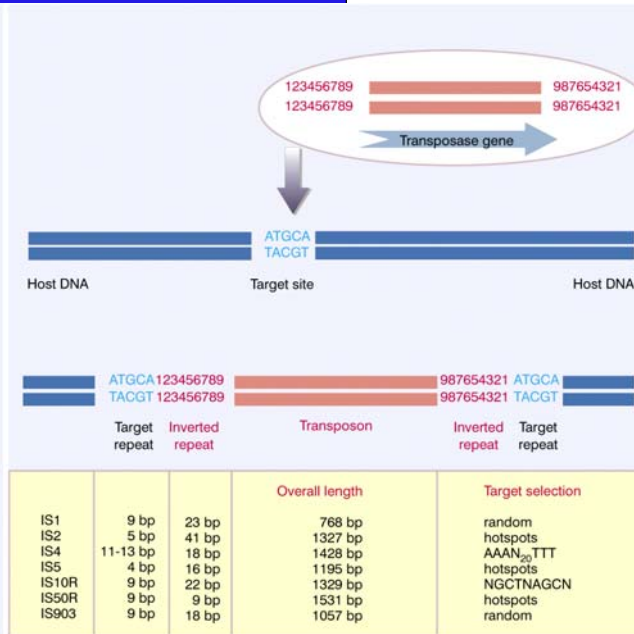
Figure 9-11

## 9.3 General structure of bacterial IS elements

Transposition is random site  
In lysogenic virus - insertion to new cell

# Transposon - basic

**Figure 15.1** Overview: transposons have inverted terminal repeats and generate direct repeats of flanking DNA at the target site. In this example, the target is a 5 bp sequence. The ends of the transposon consist of inverted repeats of 9 bp, where the numbers 1 through 9 indicate a sequence of base pairs.



## 9.3 General structure of bacterial transposons

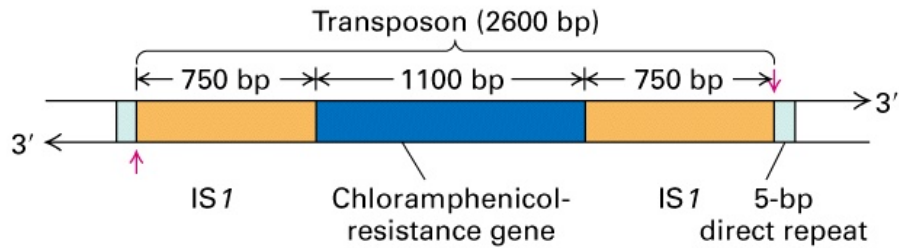
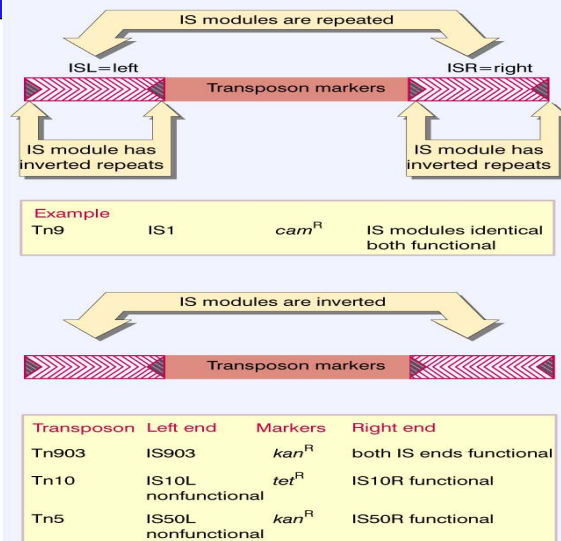


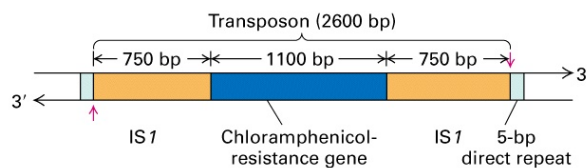
Figure 9-13

## Transposon - bacterial

**Figure 15.2** A composite transposon has a central region carrying markers (such as drug resistance) flanked by IS modules. The modules have short inverted terminal repeats. If the modules themselves are in inverted orientation (as drawn), the short inverted terminal repeats at the ends of the transposon are identical.



## 9.3 bacterial transposons - valuable tools



Easily selected (antibiotic resistant)  
 Act as a one site mutagen  
 Can be inserted into plasmid, viral etc

Figure 9-13

## 9.3 Eukaryotic transposons - valuable tools

In corn

In fly

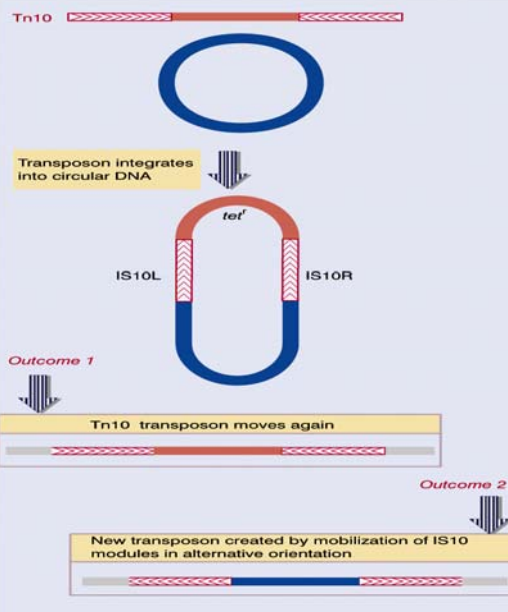
(50% of spontaneous mutation are mobile DNA)

Drosophila - P element, non replicative mechanism  
mode to introduce mutations

Figure 9-13

### Transposon - Creating a new one

**Figure 15.3** Two IS10 modules create a composite transposon that can mobilize any region of DNA that lies between them. When Tn10 is part of a small circular molecule, the IS10 repeats can transpose either side of the circle.



See you in few weeks....

Enjoy