

RESTRICTION ENZYMES, PALINDROMES & C-DNA

Presentation by

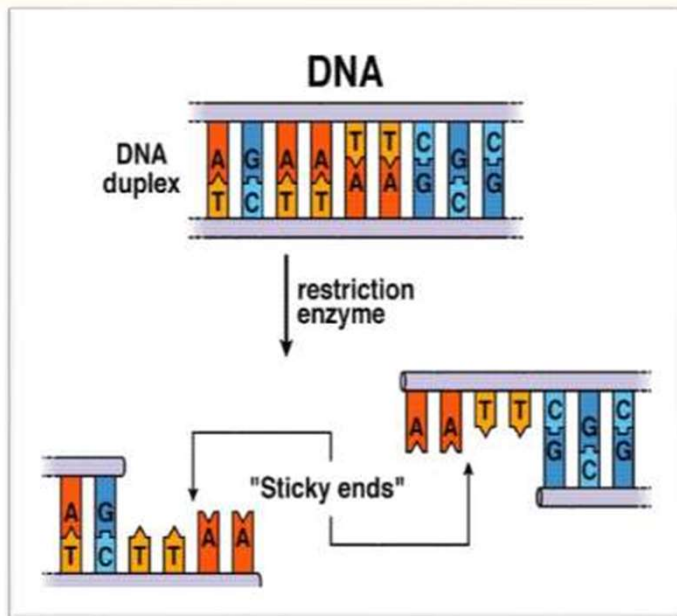
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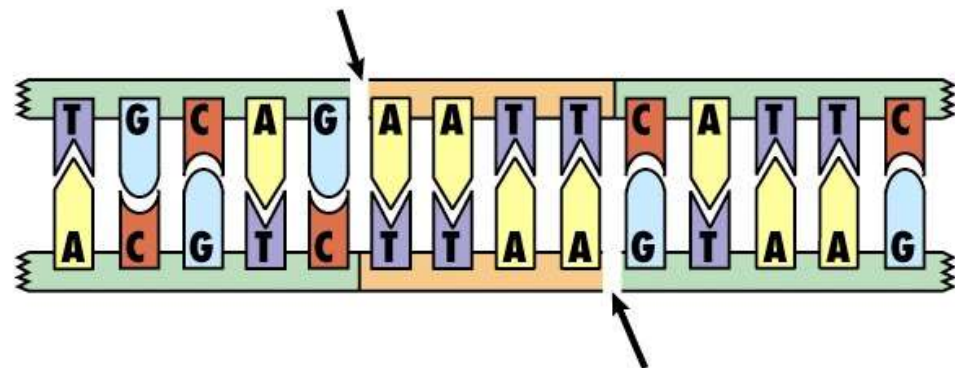
RESTRICTION ENZYME



- A special class of sequence-specific enzyme
- Found in bacteria
- Site-specific-cleave DNA molecules only at specific nucleotide sequence
- REases recognize DNA base sequence that are palindrome (MADAM)
- REase make staggered cuts with complementary base sequences for easy circulization

MOLECULAR SCISSORS/ RESTRICTION ENZYMES

Restriction
enzyme *EcoRI*
cuts here
(before AATT)

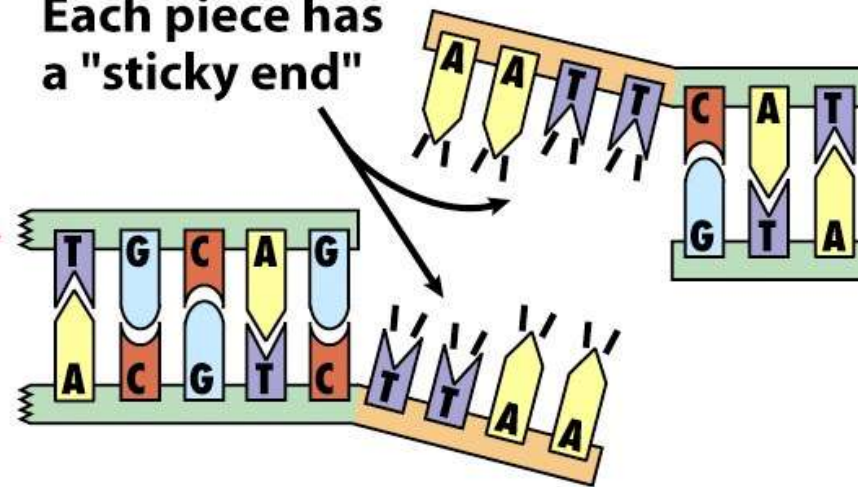


EcoRI
cuts here
(before AATT)

Pieces
separate

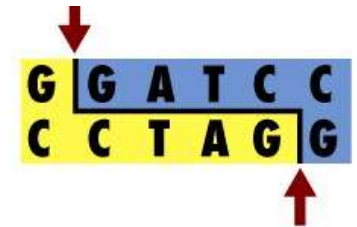


Each piece has
a "sticky end"



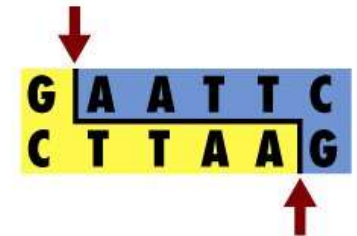
Bacillus amyloliquefaciens H

Bam H1



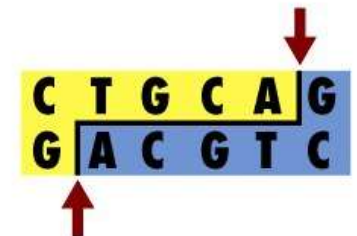
Escherichia coli Ry13

Eco R1



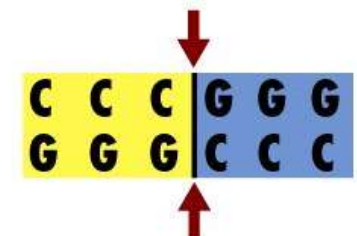
Providencia stuartii 164

Pst 1



Serratia marcescens SB

Sma H1



Rhodopseudomonas sphaeroides

Rsa 1

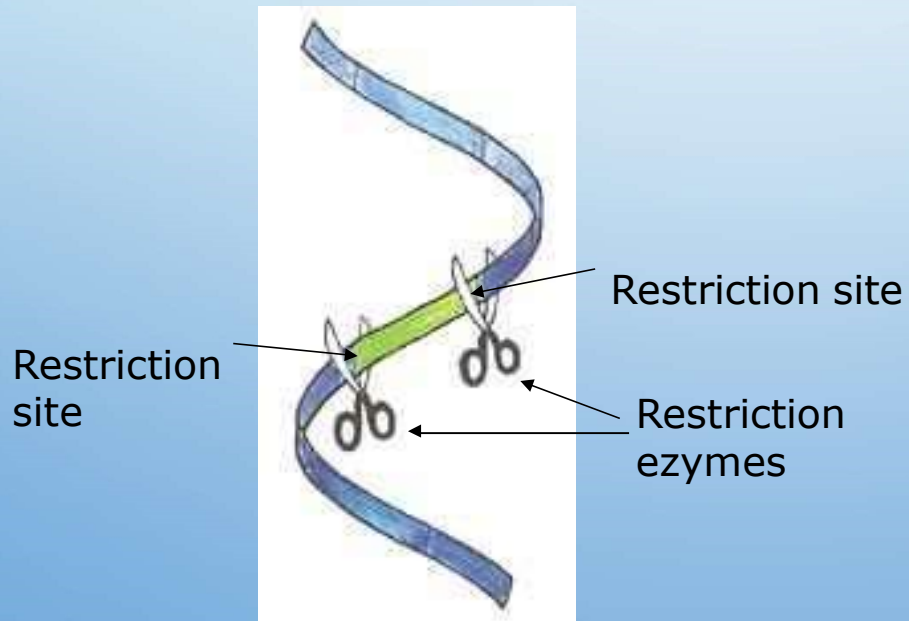


MOLECULAR TOOLS OF GENETIC ENGINEERING

- The genetic engineer's tool kit or molecular tool namely the enzymes are most commonly used in recombinant DNA experiments are
 - } Restriction endonucleases -DNA cutting Enzyme.
 - } DNA Ligases- DNA joining Enzyme.

RESTRICTION ENDONUCLEASES

Restriction enzymes act as molecular scissors and cut DNA at specific sites called restriction sites



RESTRICTION ENDONUCLEASES

- Named with particular reference to the bacteria from which they are isolated.

Eg. EcoR_I

Eco R_I, Eco R_{II}, Eco R_{III}

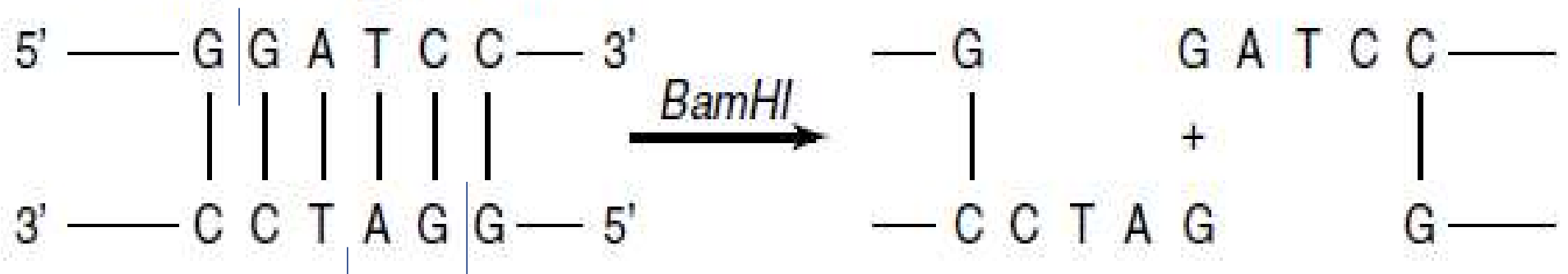
:Types 3

PALINDROME

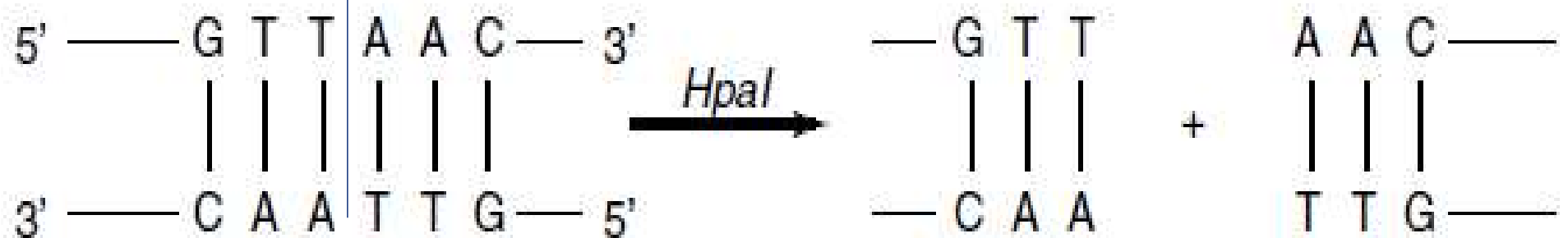
- MADAM
- EcoRI
- 5'GAATTC3' →
- ← 3'CTTAAG5'

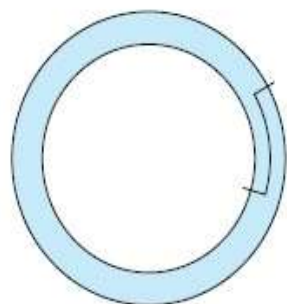
G|AATTC
CTTAAG|

A. Sticky or staggered ends



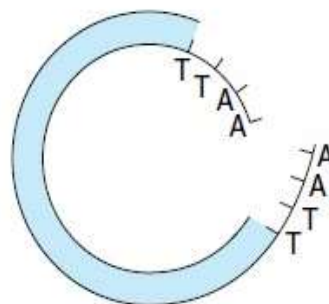
B. Blunt ends



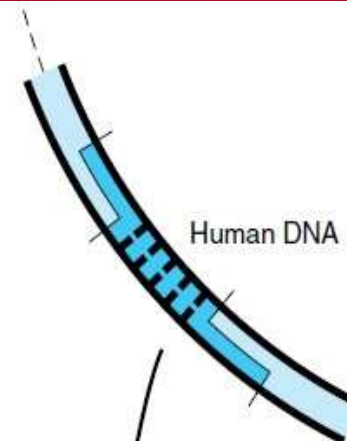


Circular plasmid DNA

EcoRI
restriction
endonuclease



Linear plasmid DNA
with sticky ends



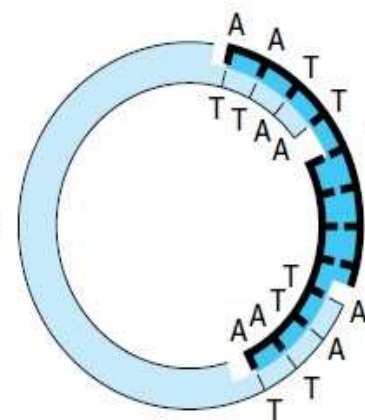
Human DNA

EcoRI restriction
endonuclease

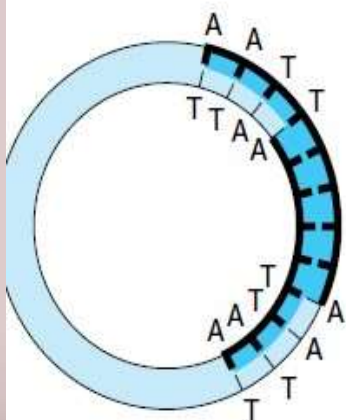


Piece of human DNA cut with
same restriction nuclease and
containing same sticky ends

Anneal



DNA
ligase

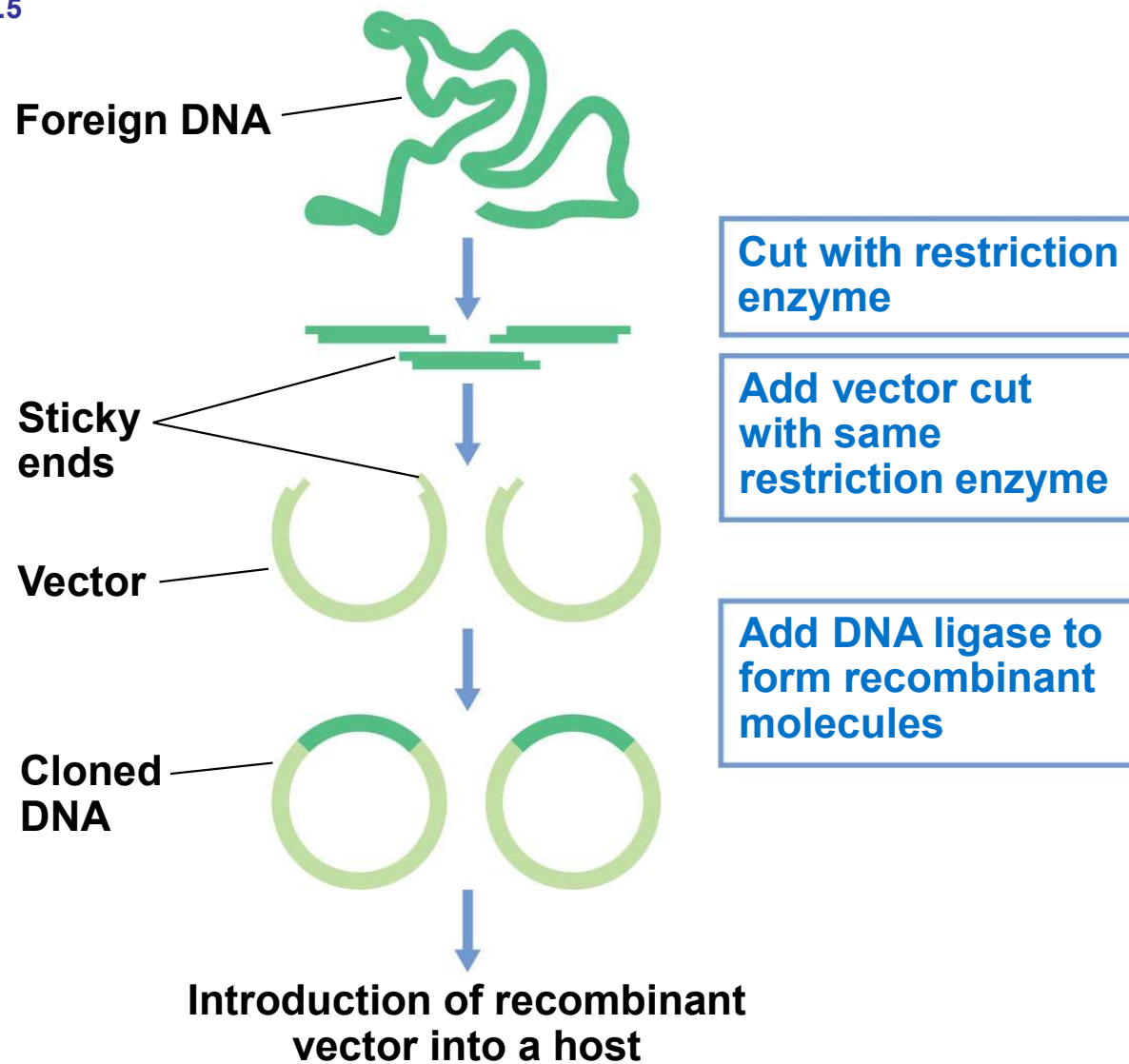


Plasmid DNA molecule with human DNA insert
(recombinant DNA molecule)

Endonuclease	Sequence Recognized Cleavage Sites Shown	Bacterial Source
<i>Bam</i> HI	↓ GGATCC CCTAGG ↑	<i>Bacillus amylo- liquefaciens</i> H
<i>Bgl</i> II	↓ AGATCT TCTAGA ↑	<i>Bacillus glolbigii</i>
<i>Eco</i> RI	↓ GAATTC CTTAAG ↑	<i>Escherichia coli</i> RY13
<i>Eco</i> RII	↓ CCTGG GGACC ↑	<i>Escherichia coli</i> R245
<i>Hind</i> III	↓ AAGCTT TTCGAA ↑	<i>Haemophilus influenzae</i> R _d

<i>HhaI</i>	↓ GCGC ↑ CGCG	<i>Haemophilus haemolyticus</i>
<i>HpaI</i>	↓ GTTAAC ↑ CAATTG	<i>Haemophilus parainfluenzae</i>
<i>MstII</i>	↓ CCTNAGG ↑ GGANTCC	<i>Microcoleus strain</i>
<i>PstI</i>	↓ CTGCAG ↑ GACGTC	<i>Providencia stuartii</i> 164
<i>TaqI</i>	↓ TCGA ↑ AGCT	<i>Thermus aquaticus</i> YTI

Figure 11.5



CDNA LIBRARY

COMPLEMENTARY DNA LIBRARY

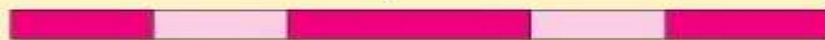
- cDNA libraries contain those DNA sequences that appear as mRNA molecules, and these differ from one cell type to another.
- This mRNA can be used as a template to make a complementary dsDNA (cDNA) molecule using the enzyme reverse transcriptase.

DNA of eukaryotic gene



1 Transcription in the cell

RNA transcript



2 Introns removed and exons spliced together in the cell

mRNA



3 Isolation of mRNA from cell and addition of reverse transcriptase
Synthesis of DNA strand

Reverse transcriptase



DNA strand being synthesized

4 Breakdown of mRNA



5 Synthesis of second DNA strand

DNA of gene without introns



Exons: Coding sequences; form part of protein

Introns: non coding sequences; removed from mRNA

Newly formed RNA = naïve RNA / RNA transcript

THANK YOU

