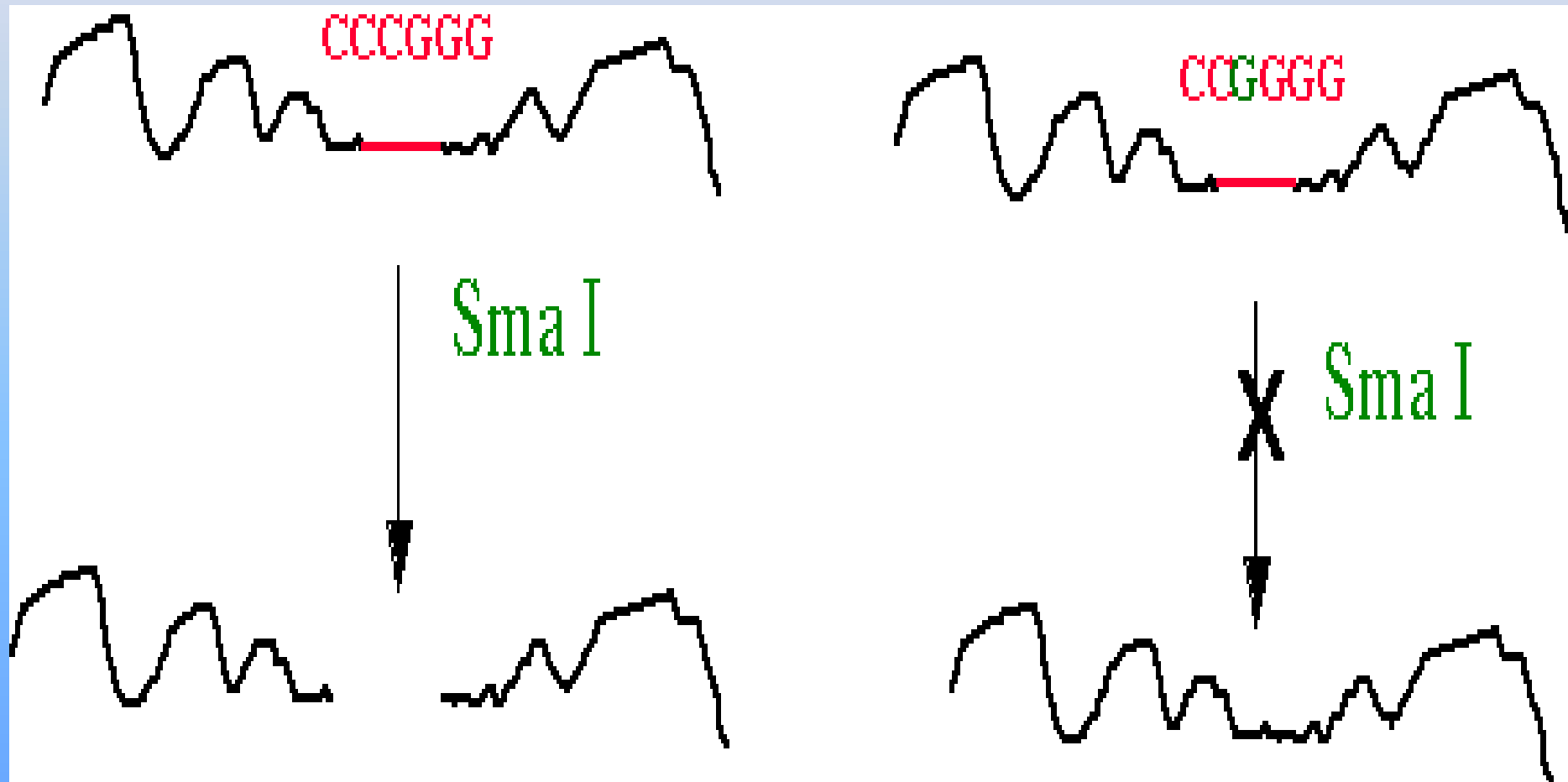


Enzim Endonuklease Restriksi:

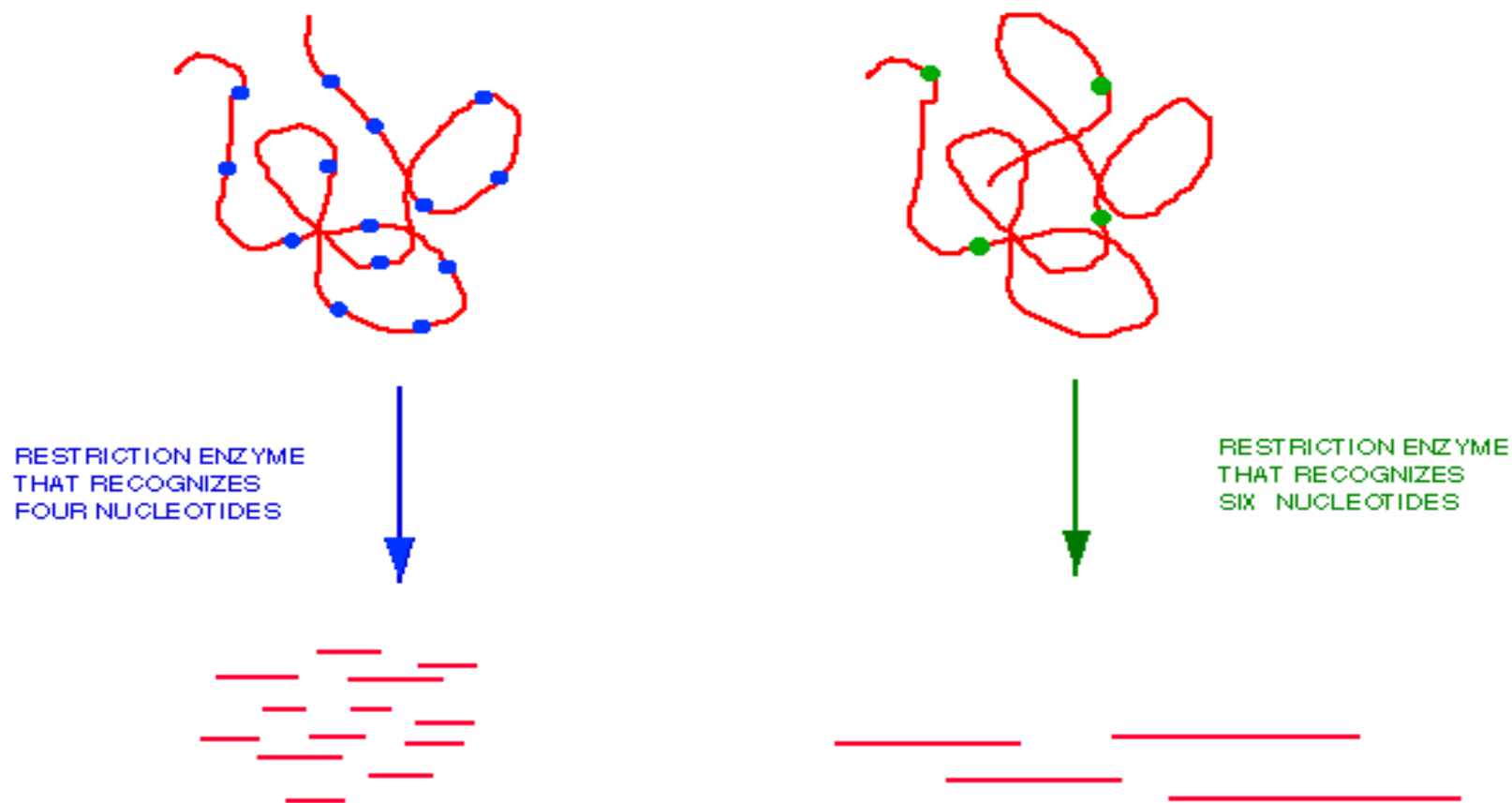
**MODAL DASAR REKAYASA GENETIKA**

**Genetika 4019**

Enzim Restriksi hanya memotong DNA pada urutan basa (nukleotida) yang “dikenalnya”




A restriction enzyme with a four nucleotide recognition sequence cuts long DNA more frequently and produce smaller DNA fragments than a restriction enzymes with a six nucleotide recognition sequence.



Any given **four nucleotide long recognition site** occurs in DNA, on average, at a distance of **256** ( $4^4$ ) nucleotides. Any given **six nucleotide long sequence** occurs, on average, at a distance of **4096** ( $4^6$ ) nucleotides. Any given **eight nucleotide long sequence** occurs, on average, at a distance of **65536** ( $4^8$ ) nucleotides

**RESTRICTION ENZYMES** often act as dimers; each restriction enzyme subunit recognizes **THE SAME (5'->3')** nucleotide sequence in complementary DNA strands:

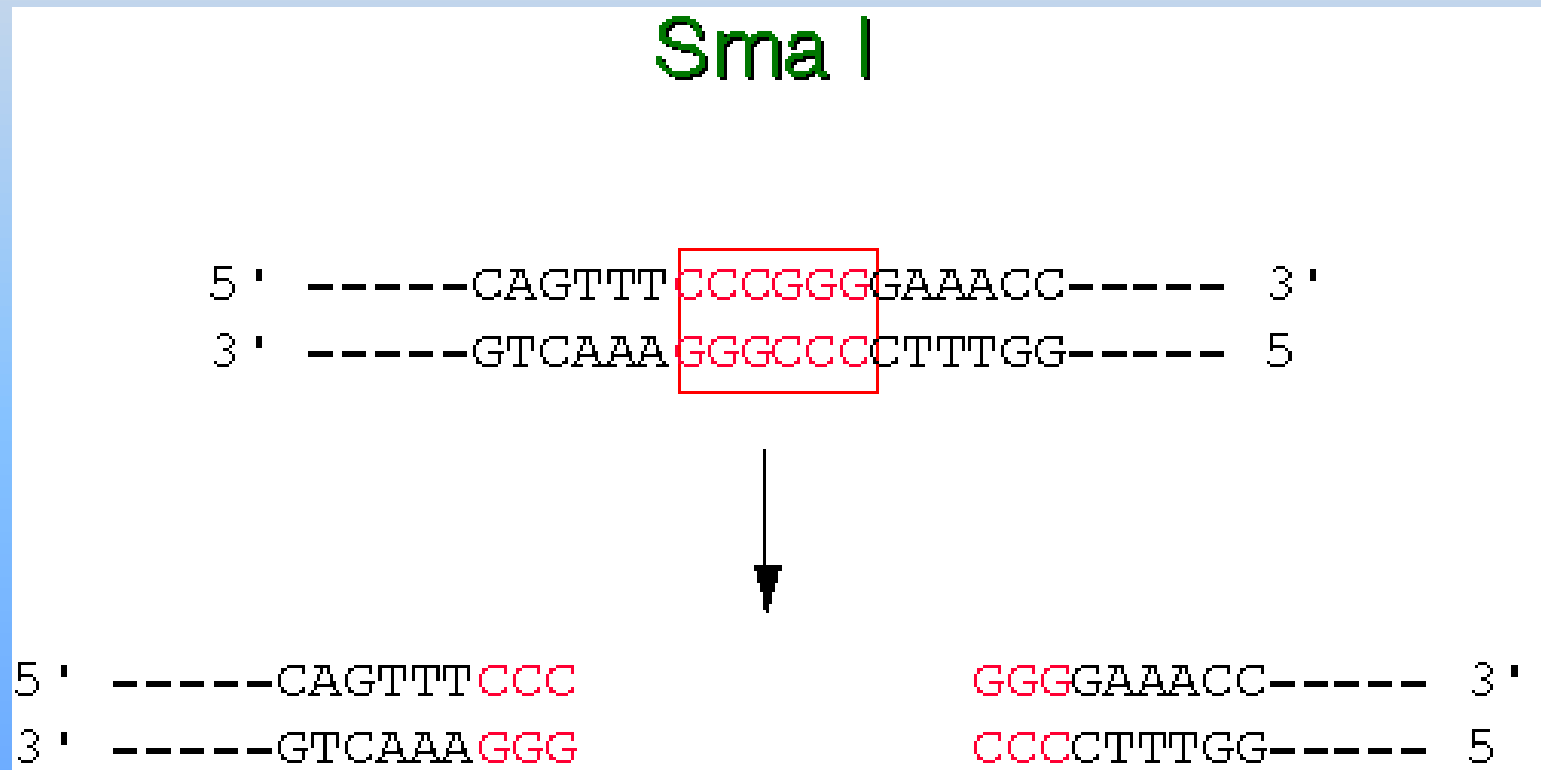


5 ' GCTAGGCATGAGTACCATT**AAGCTT**CGGATCGCATCGACTCAGC 3 '  
3 ' CGATCCGTACTCATGGTAA**ATTCGAA**GCSTAGCGTAGCTGAGTCG 5 '

As a result, restriction enzymes usually recognize palindromic sequences:

<b>EcoRI</b>	5 ' GAATTC 3 '	<i>EcoRI</i> = <i>E. coli</i> RY13 (R.N. Yoshimori).
	3 ' CTTAAG 5 '	
<b>HindIII</b>	5 ' AAGCTT 3 '	<a href="#"><i>Hind</i>=<i>Haemophilus influenzae</i></a>
	3 ' TTCGAA 5 '	
<b>SmaI</b>	5 ' CCCGGG 3 '	<i>Sma</i> = <i>Serratia marcescens</i>
	3 ' GGGCCC 5 '	
<b>TaqI</b>	5 ' TCGA 3 '	<i>Taq</i> = <i>Thermus aquaticus</i>
	3 ' AGCT 5 '	

Some restriction enzymes, for example, Sma I, cut both DNA strands in the middle of the recognition sequence and produce "blunt-end" DNA fragments:



Many restriction nucleases produce staggered cuts in DNA which leave short single stranded "tails" at the ends of DNA fragments:

(1)

**EcoRI**

```
5' -----CCTAGCGAATTCGTCTTA----- 3'
3' -----GGATCGCTTAAGCAGAAT----- 5'
```



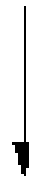
**EcoRI**

```
5' -----CCTAGCG           AATTCGTCTTA----- 3'
3' -----GGATCGCTTAA           GCAGAAT----- 5'
```

Many restriction nucleases produce staggered cuts in DNA which leave short single stranded "tails" at the ends of DNA fragments:  
(2)

## Hind III

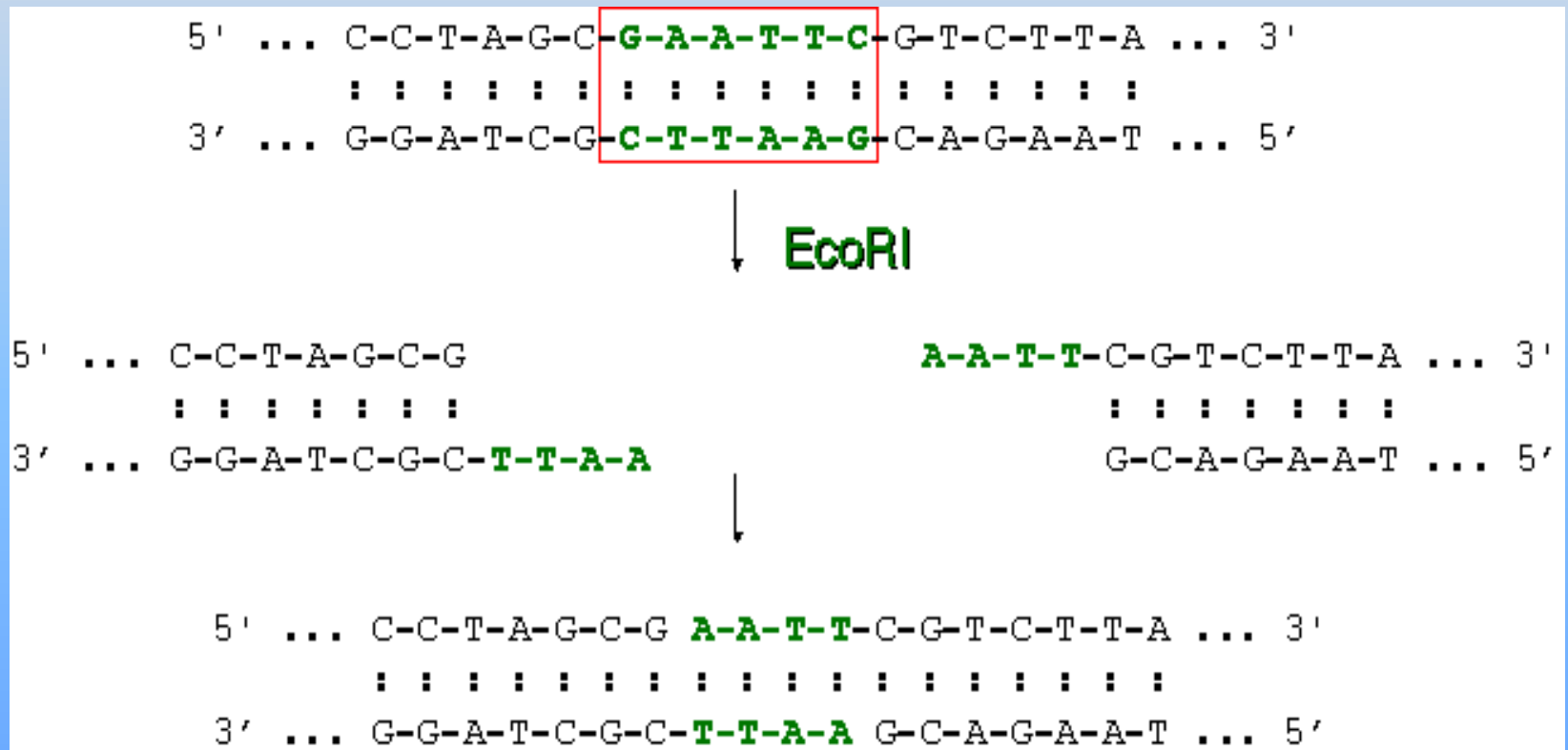
5' -----CAGTTTAAGCTTGAAACC----- 3'  
3' -----GTCAAATTCGAACTTTGG----- 5'



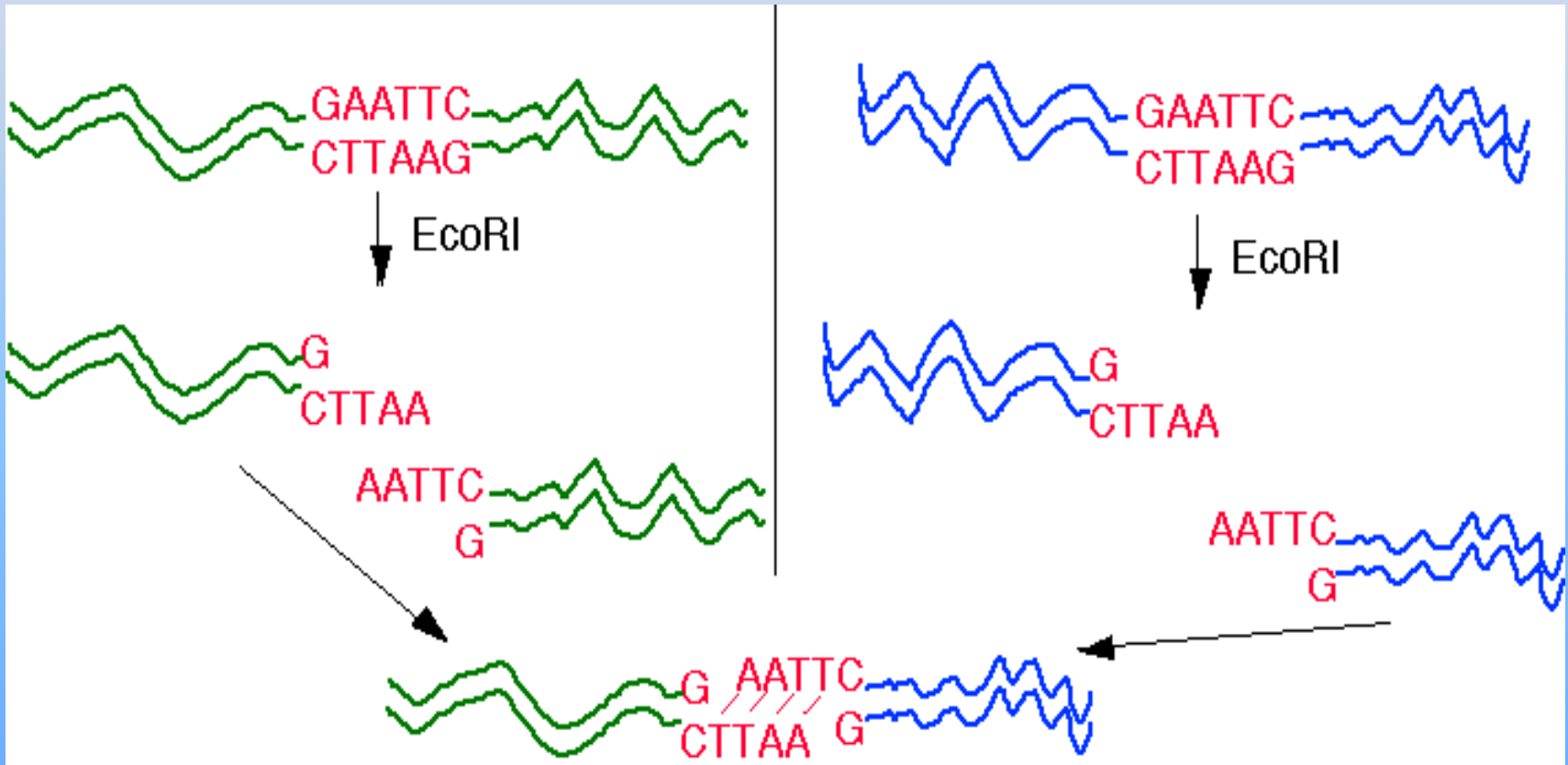
5' -----CAGTTTA **A** AGCTTGAAACC----- 3'  
3' -----GTCAAATTCGA **A** ACTTTGG----- 5'



Single stranded tails produced in the result of DNA cleavage by restriction nucleases are known as cohesive or "sticky" ends. They can form complementary base pairs with the DNA ends produced by the same restriction enzyme.

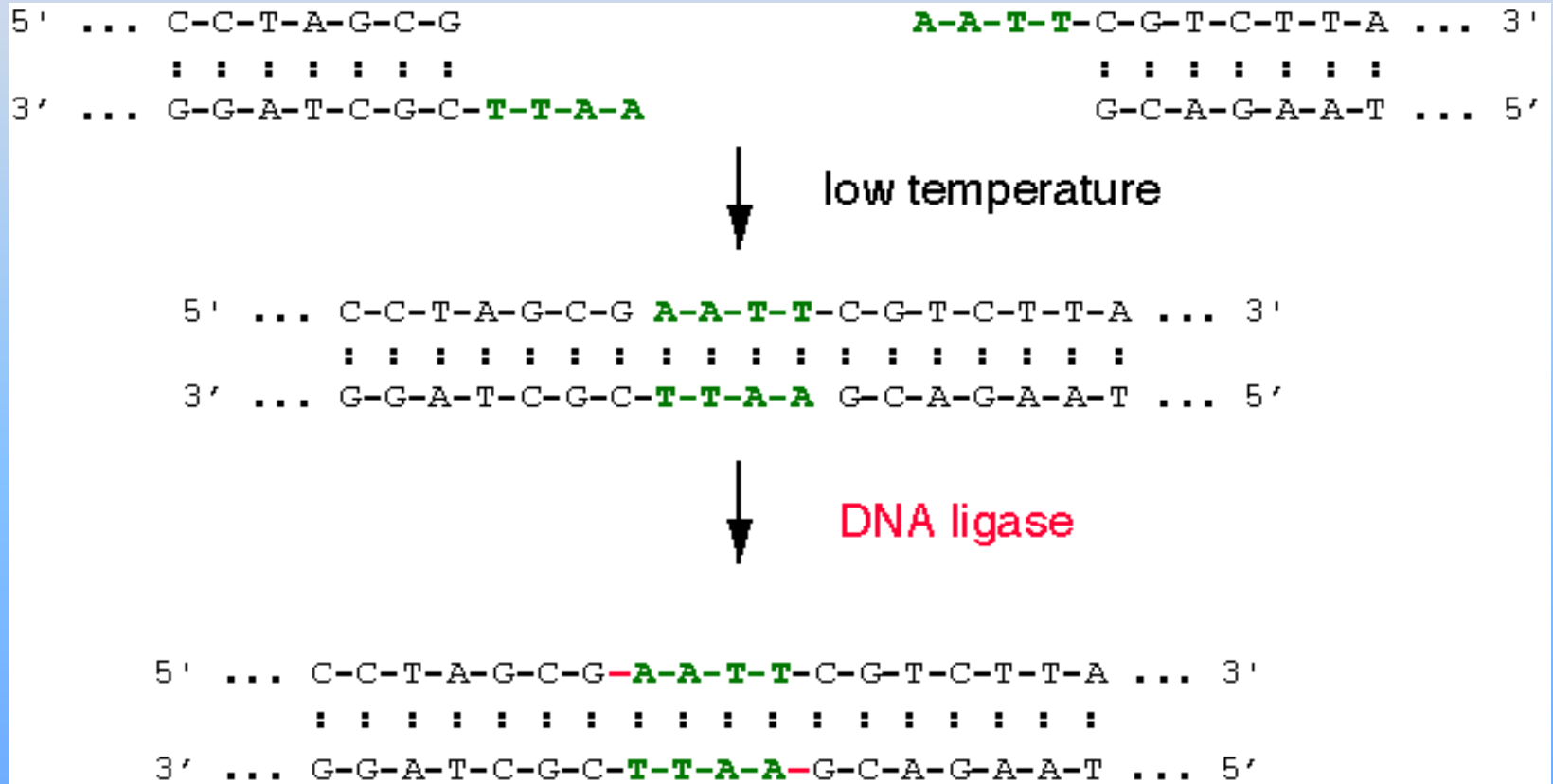


Sticky ends of different DNA fragments produced by the same restriction enzyme can base-pair to each other. Linking together such fragments will produce a recombinant DNA molecule.

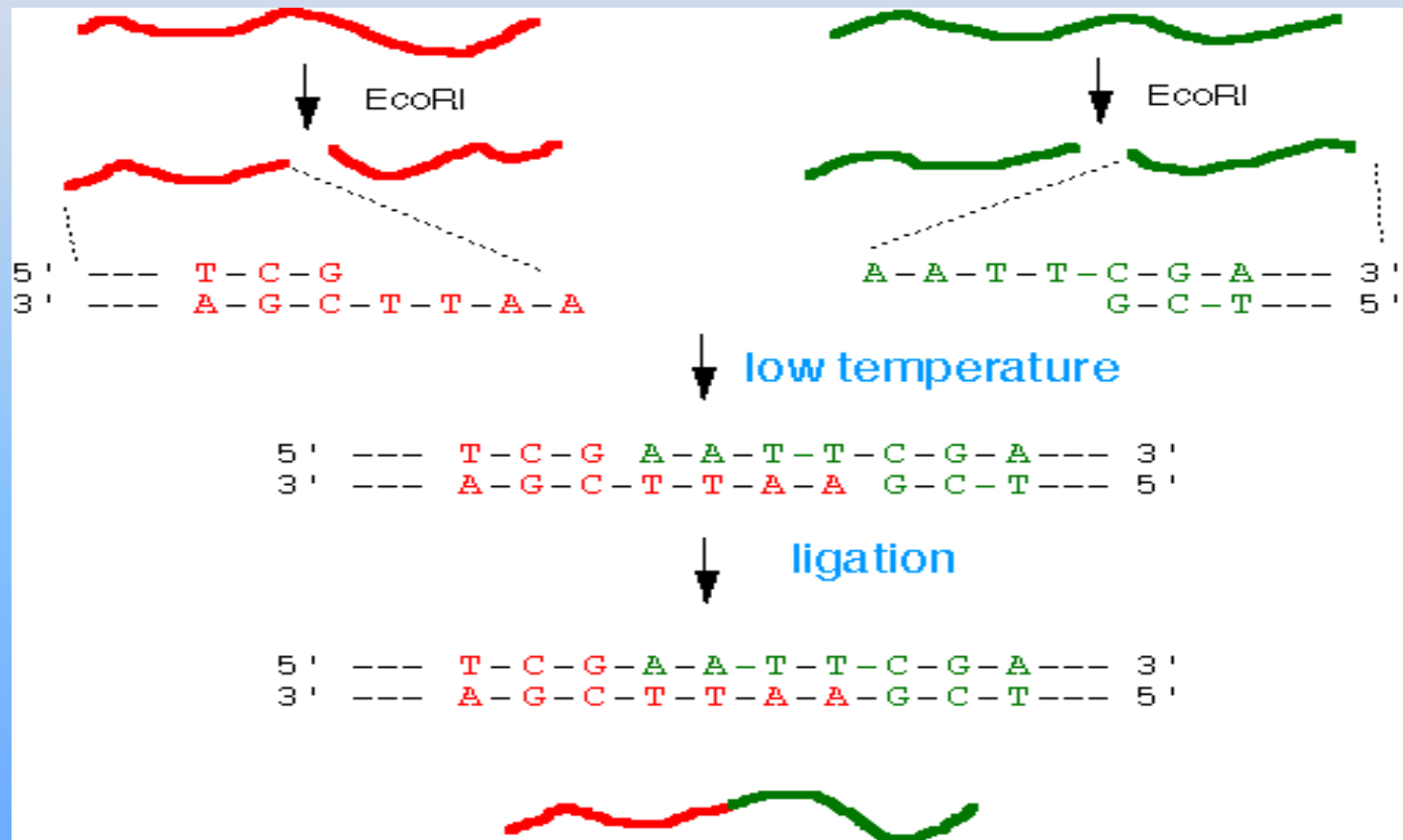


# DNA LIGASE

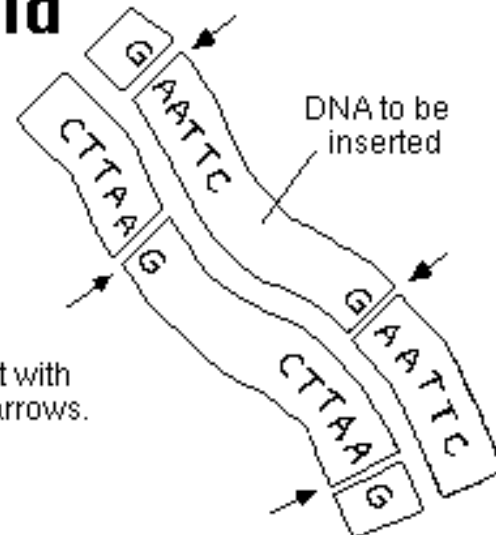
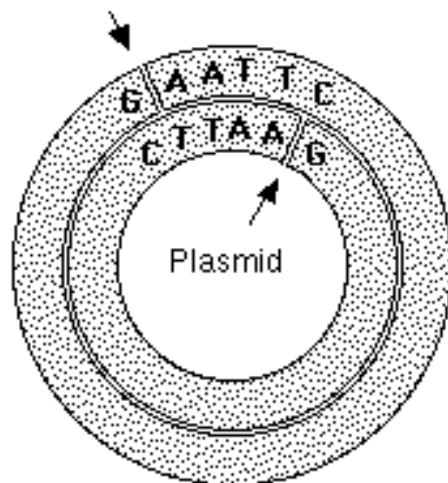
DNA ligase is an enzyme that can connect two DNA fragments together.



# Combining together fragments of different DNA molecules we can produce recombinant DNA

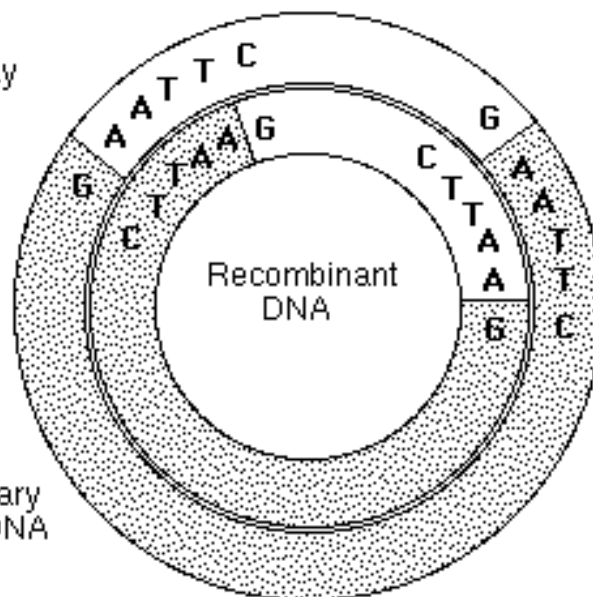


# Inserting a DNA Sample into a Plasmid



DNA is cut with *EcoRI* at arrows.

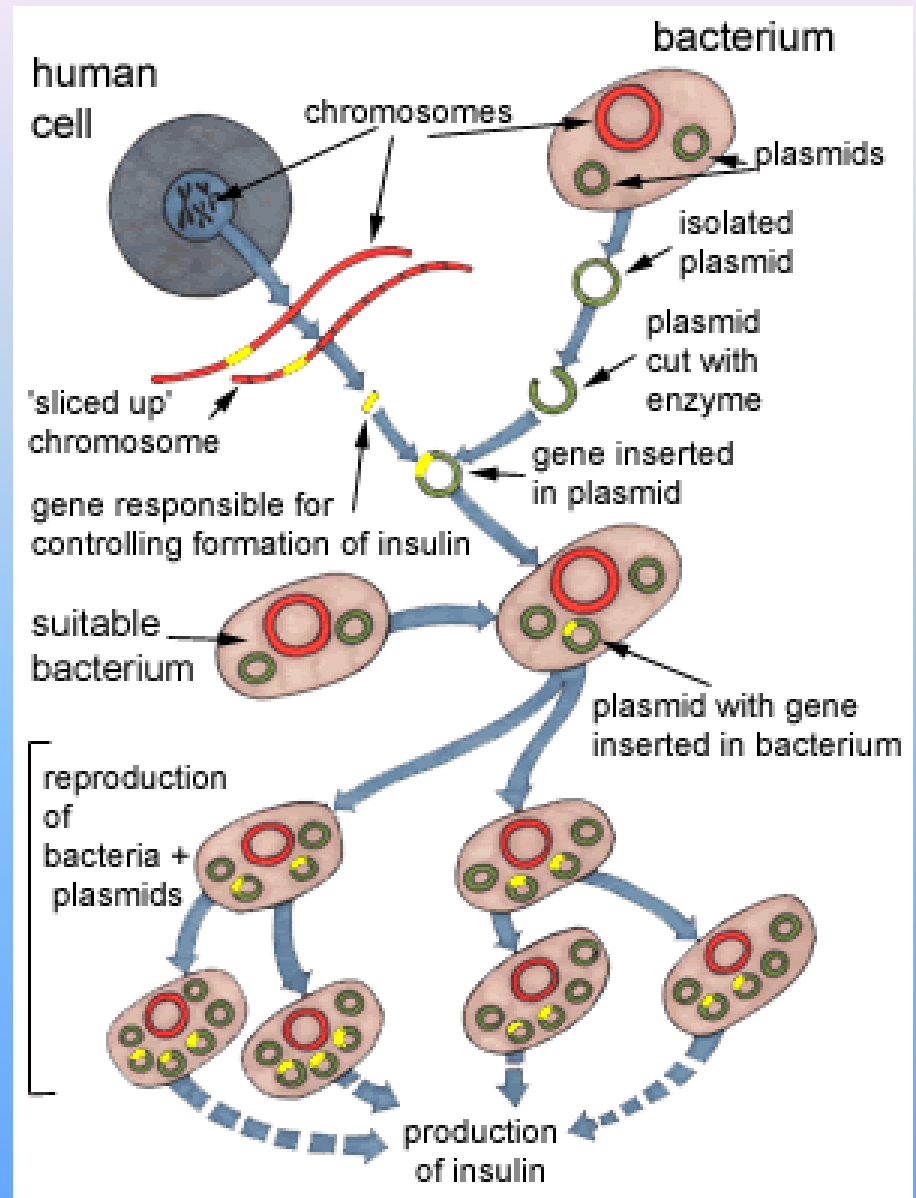
Resulting DNAs have sticky (complementary) ends.

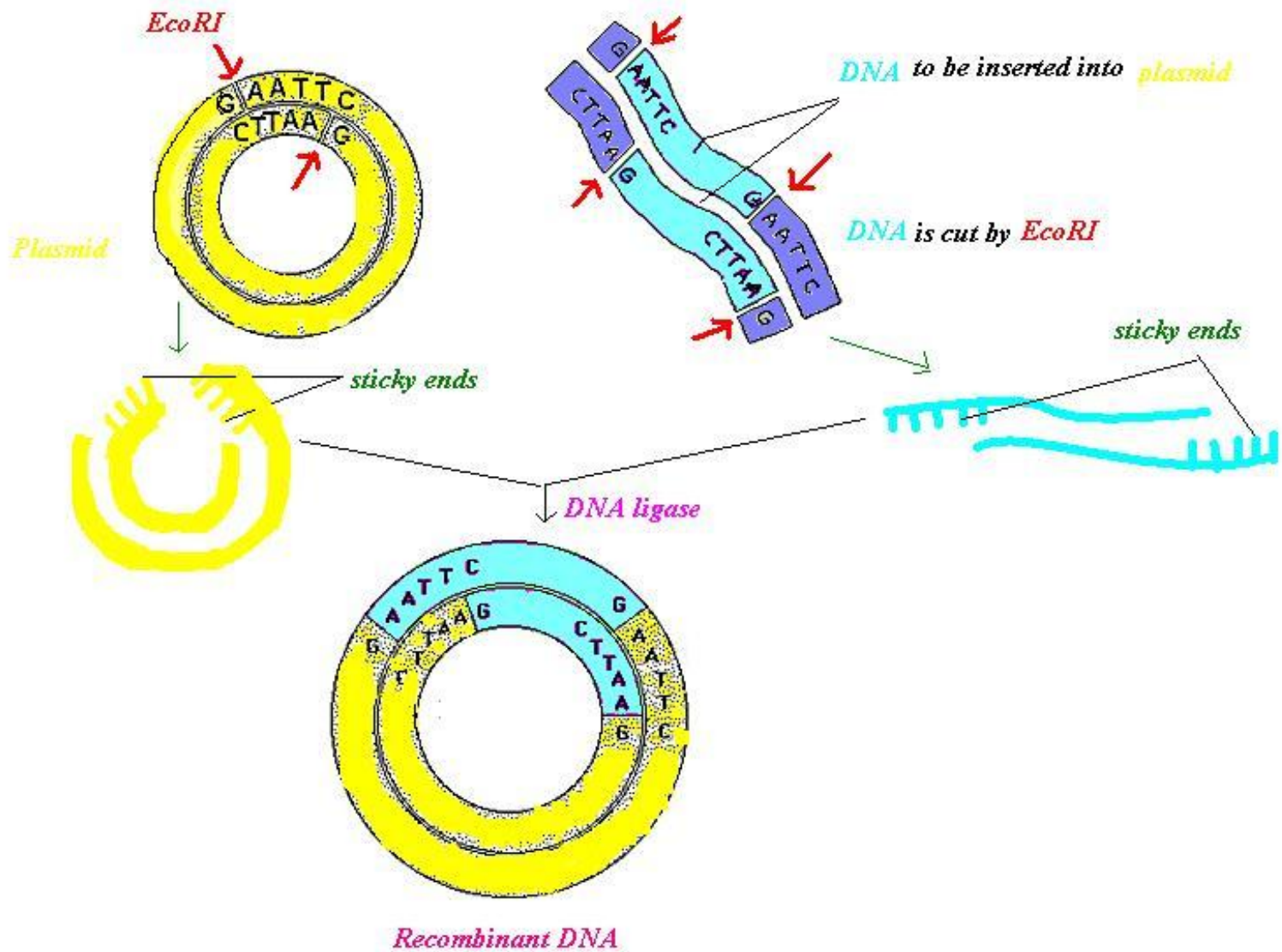


DNA is spliced by complementary base pairing and sealed with DNA ligase

# Contoh aplikasi pada produksi insulin

1. Gen manusia yang mengekspresikan insulin diisolasi
2. Plasmid dari bakteri diisolasi untuk kemudian diinkubasi dengan enzim restriksi
3. Gen manusia disisipkan kepada plasmid
4. Plasmid “dibiakkan” dalam sel bakteri (cloning gen)
5. Sel bakteri mengekspresikan gen insulin manusia, hasilnya (insulin manusia?)
6. Pada contoh di atas, apakah enzim restriksi yang digunakan pada sel manusia sama dengan yang digunakan pada plasmid?





# TUGAS INDIVIDUAL

1. Carilah sepuluh macam lagi endonuklease restriksi
2. Bagaimana urutan basa sebagai gugus palindrom untuk setiap enzim yang telah anda pilih