

R.E - Enzymes → Type I, Type II, III, IV

- Type I → 1 Enzymes have different subunits → Recognition, cleavage, modification units

- It requires energy → ATP molecule
- requires cofactor → S-adenosyl methionine
- Methylation at 1 base pair ← modification
- Recognition → 1000 bp away from the methylation
- This are not used more.

- Type II → 2 different enzymes  
↓  
cleavage      modification

- recognize the same target sequence
- They do not require ATP, Cofactor.
- Sequences are symmetrical.
- widely used in genetic engineering.

- Type III → Same as type I, 2 subunits

↓  
Recognition      Cleavage  
↓  
modification

Same sequence

- cleavage is done at 24-26 bp away from recognition site.

## • Expression vector

- not only carry the foreign DNA also expresses
  - have Regulatory elements - Prom, Ter
  - Ribosome binding site
  - Ori site, marker gene, multiple cloning site
- eg: PET.

- Vector: extrachromosomal, self replicating, autonomous
- recognition site, restriction enzymes, Ori site, marker gene, multiple cloning / cleavage site, Transformation, mobilisation, Selection, Cloning Vector, Expression Vector, gene of interest, Self regulatory elements, promoter, terminator, ribosomal binding site, DNA, mRNA, protein synthesis, plasmid, cosmid, BAC, YAC, PET, Ampicillin resistant, antibiotic Tetracycline, Streptomycin, Host cell, E. coli, Ligase
- express, fluorescence gene, green fluorescence protein, isolation & purification of plasmid

## • PBR322

- first vector developed by Boliver & Rodriguez → 1977

P - Plasmid

B - Boliver

R - Rodriguez

- molecular weight →  $2.83 \times 10^6$  Daltons
- have ori site

- marker gene → ampicillin, tetracycline

(AD) → recognition site → PstI, SalI, EcoRI, HindIII, EcoRV

- first artificial plasmid. (vector)

- Origin of replication in Plasmid PBR322 is known as PMB1