NAME	Recognition Sequences	Sticky/blunt	Source
ECORI	GAATTC G CTTAAG CTTAA	Sticky end	Escharichia coli
HindⅢ	AAGCTT, A TT CGA	Blunt non sticky	Haemophilus
BamI	9 G A T C C, G C C T A G, G CCTAG	1 - , 7 - 19	Bacillys 9mytoliquetans
EcoRV	GATATC GAT CTATAG CTA	Blunt	Escherichia (Oli

sphilus influence RE(disc) RE
Hind-III

GTCGAC GTC GAC

CAGCTG CAG CTG

+ GCTT A AGT Hind-III TTCGA

Nomenclature of Restriction Endo.

15t letter => Genus of Bacteria (italic) 2 8/3 Letter= Species of Barteria 4th 1etter => Strain of Bacteria (optional) Roman Numerical => Enzyme=> Isolateed from +St enzyme Escharichia coli Genes Species Isolated from bauteria

strain. Roman numbers following the names indicate the order in which the enzymes were isolated from that strain of bacteria.

Restriction enzymes belong to a larger class of enzymes called nucleases. These are of two kinds; exonucleases and endonucleases. Exonucleases remove nucleotides from the ends of the DNA whereas, endonucleases make cuts at specific positions within the DNA.

Each restriction endonuclease functions by 'inspecting' the length of a DNA sequence. Once it finds its specific recognition sequence, it will bind to the DNA and cut each of the two strands of the double helix at specific points in their sugar-phosphate backbones (Figure 11.1). Each restriction endonuclease recognises a specific mons (enclain dromic nucleotide sequences in the DNA.

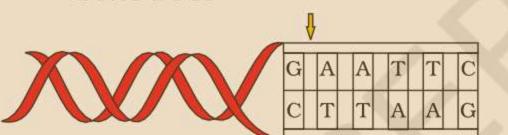
5'CCCGGG3'

Action of Restriction enzyme

Sma-I (Serratia

The enzyme cuts both DNA

Vector DNA



EcoRI cuts the DNA between bases G and A only when the sequence GAATTC is present in the DNA

Foreign DNA



