





6/22/2014

Enzymes and Restriction Digestion of DNA

B3 Summer Science Camp at Olympic High School

Dr. Jennifer Weller

Enzymes

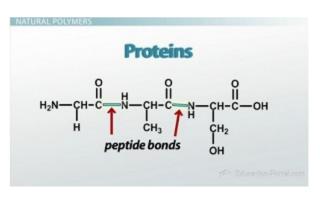
- Enzymes are proteins polymers of amino acids.
 - They carry out chemical reactions for cells.
 - They are chemical catalysts because they emerge unchanged from each reaction: they are neither reactants (substrates) nor products as usually defined.
 - They speed up the reactions cells need this because a lot of times the amount of reactant is tiny so the reaction would proceed very slowly without the enzyme.

Lysine (K)
Residue Mass 128

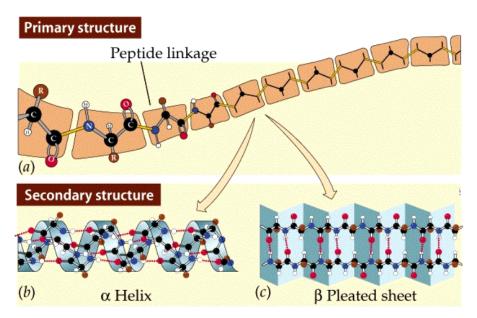
$$H_2N-CH_2-(CH_2)_3+C-COOH$$
 NH_2

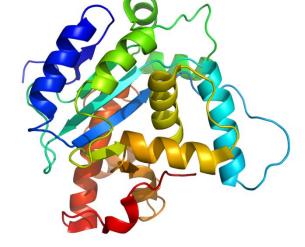
Arginine (R)
Residue Mass 156

 $NH_2-C-NH-(CH_2)_3+C-COOH$
 NH_2
 $NH_2-C-NH-(CH_2)_3+C-COOH$
 NH_2
 $NH_2-C-COOH$
 NH_2
 $NH_2-C-COOH$
 NH_2
 NH_2



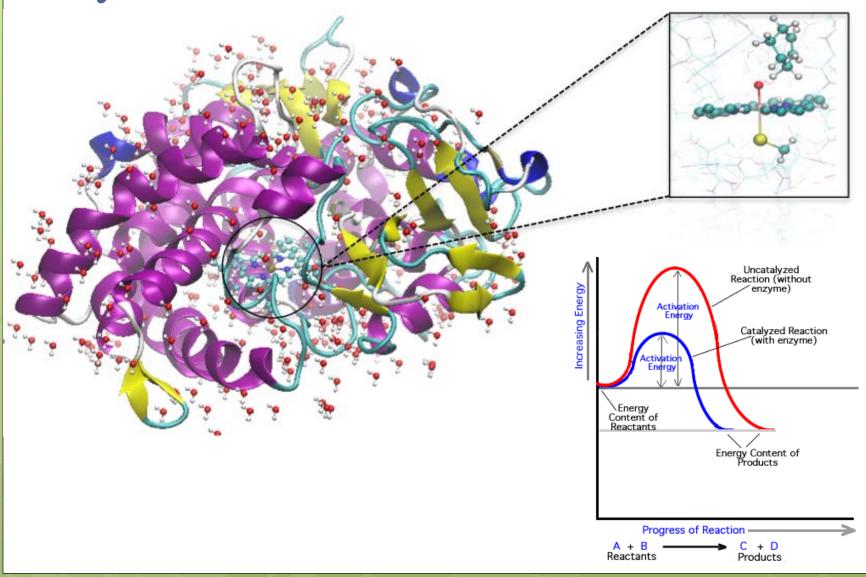
Enzymes





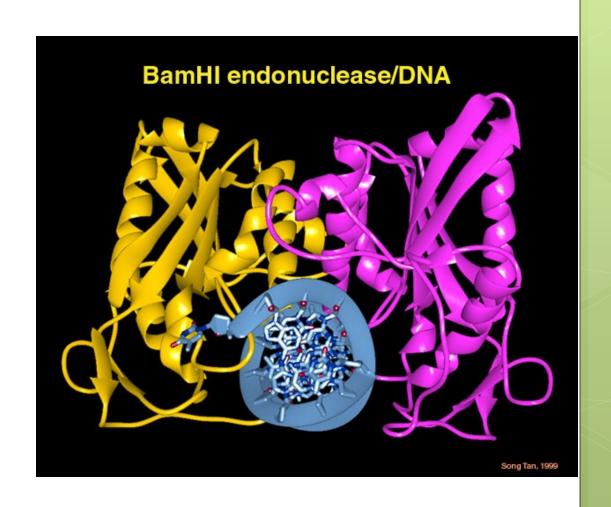
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Enzymatic Reaction



Restriction Endonucleases

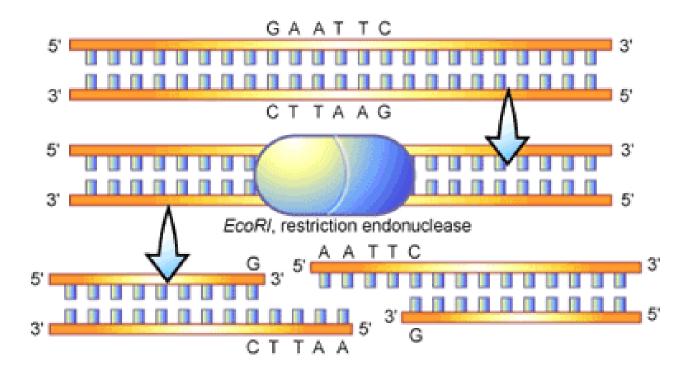
- Restriction endonucleases are enzymes.
 - The reaction they catalyze is to cut (cleave) the phosphate backbone of DNA so you end up with 2 pieces when you started with one



Restriction Endonuclease mechanism

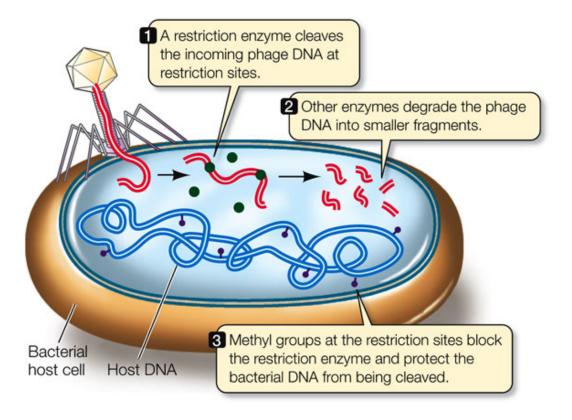
- Many Restriction Endonucleases require a <u>co-factor</u> (the Mg²⁺ above)
- Endonuclease means it cuts in the middle, not from the end of the DNA
- The enzymes are very accurate and can be very complete

Restriction Endonuclease binding site



- The enzyme *recognizes* a particular pattern of nucleotides
- For many enzymes the pattern reads the same on the opposite strands a <u>palindrome</u>
- They enzymes cut in the neighborhood of (within or just beyond) the recognition site and can give even (blunt) ends or asymmetric ends
 - In this case the overhang could be in either direction

Why did restriction endonucleases evolve?



LIFE 8e, Figure 16.1

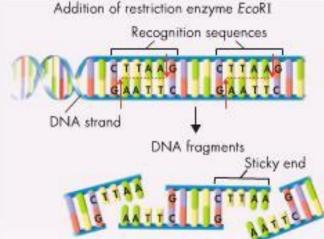
LIFE THE SCIENCE OF BIOLOGY, Flighth Edition, © 2007 Strauer Associates, Inc. and W. H. Freeman & Co.

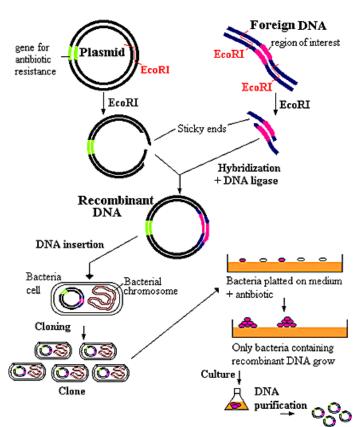
• Bacteria are also subject to viral infections (Called phage in this case)

Sticky ends are very useful in biotechnology

- Allow effective study of DNA variation for PCR and sequencing assays
- Can create recombinant products 'cloning'
 - Use bacteria to produce a gene of interest in large quantities
 - Human insulin
 - Protein to make cheese
 - Enzyme to fade blue jeans
 - Insect resistance (potato)

Pesticide resistance (corn, soybean, cotton)





Cloning into a plasmid

There are thousands of known REs and hundreds are available

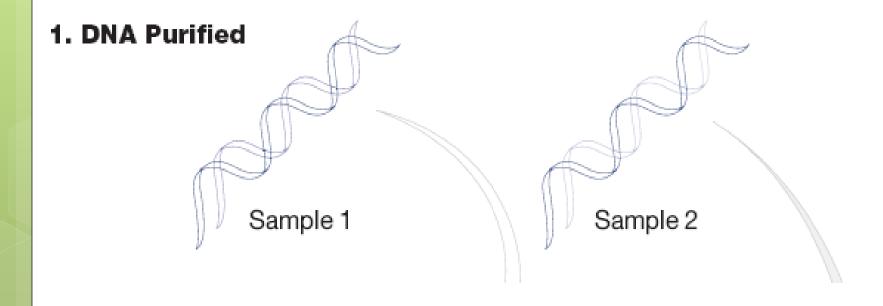
Palindromic sequence	Name of restriction enzyme that recognizes the palindrome
GAATTC CTTAAG	EcoRI
AAGCTT TTCGA <mark>A</mark>	HindIII
CTGCAG GACGTC	Pstl

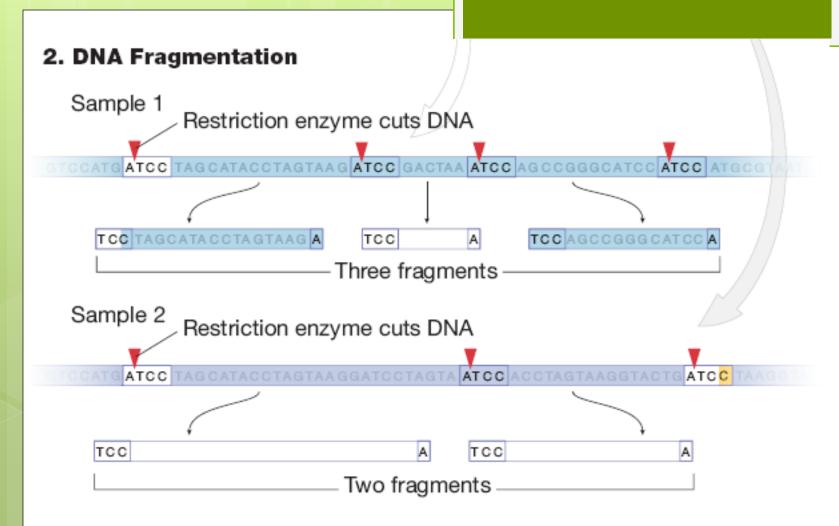
ENZYME	SEQUENCE
EcoRV	GAT/ATC
EcoRV-HF	GAT/ATC
EcoRV-HF RE-Mix	GAT/ATC
FatI	/CATG
BspDI	AT/CGAT
BspEI	T/CCGGA
BspHI	T/CATGA
PacI	TTAAT/TAA
PacI RE-Mix	TTAAT/TAA

For example, visit the New England Biolabs web site for a very large list.

The names are derived from the bacteria in which they were found: HinfI is from Haemophilus influenzae, I means the first purified.

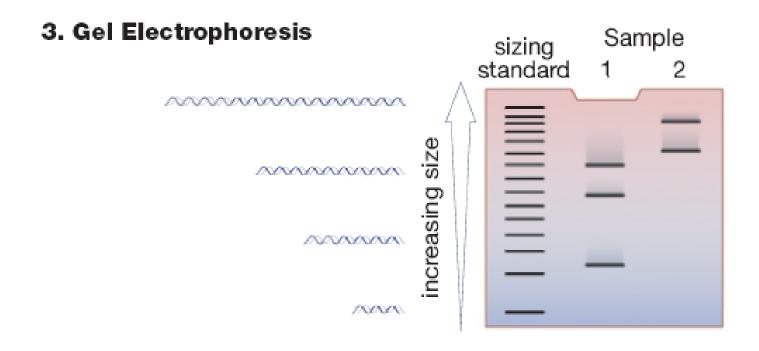
Example of using Restriction Endonucleases to characterize samples

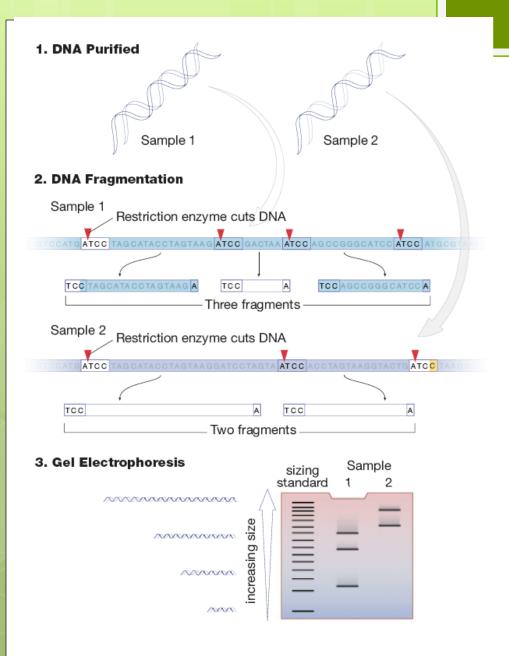




This difference in the two molecular sequence patterns is called a *polymorphism*. A nucleotide change is a <u>sequence polymorphism</u>, if it results in a pattern change then you could get a <u>length polymorphism</u>.

To detect a pattern or length polymorphism you can display the fragments on an agarose gel.





Animation – restriction enzymes

Restriction Fragment Length
 Polymorphisms