



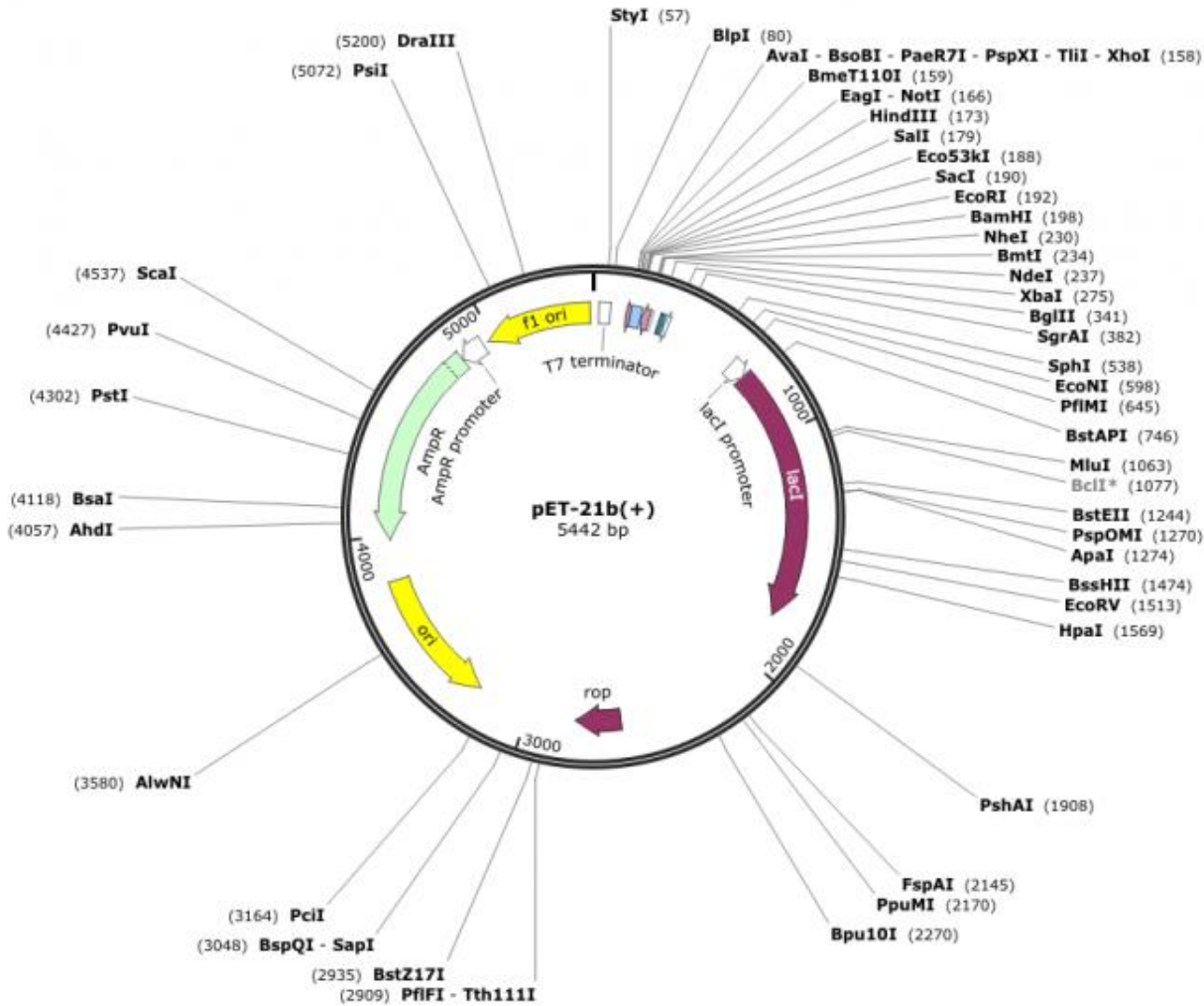
DNA synthesis & molecular cloning

Concepts

Methods

Applications

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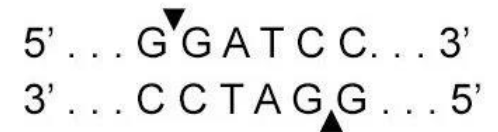
NdeI

Recognition Site:

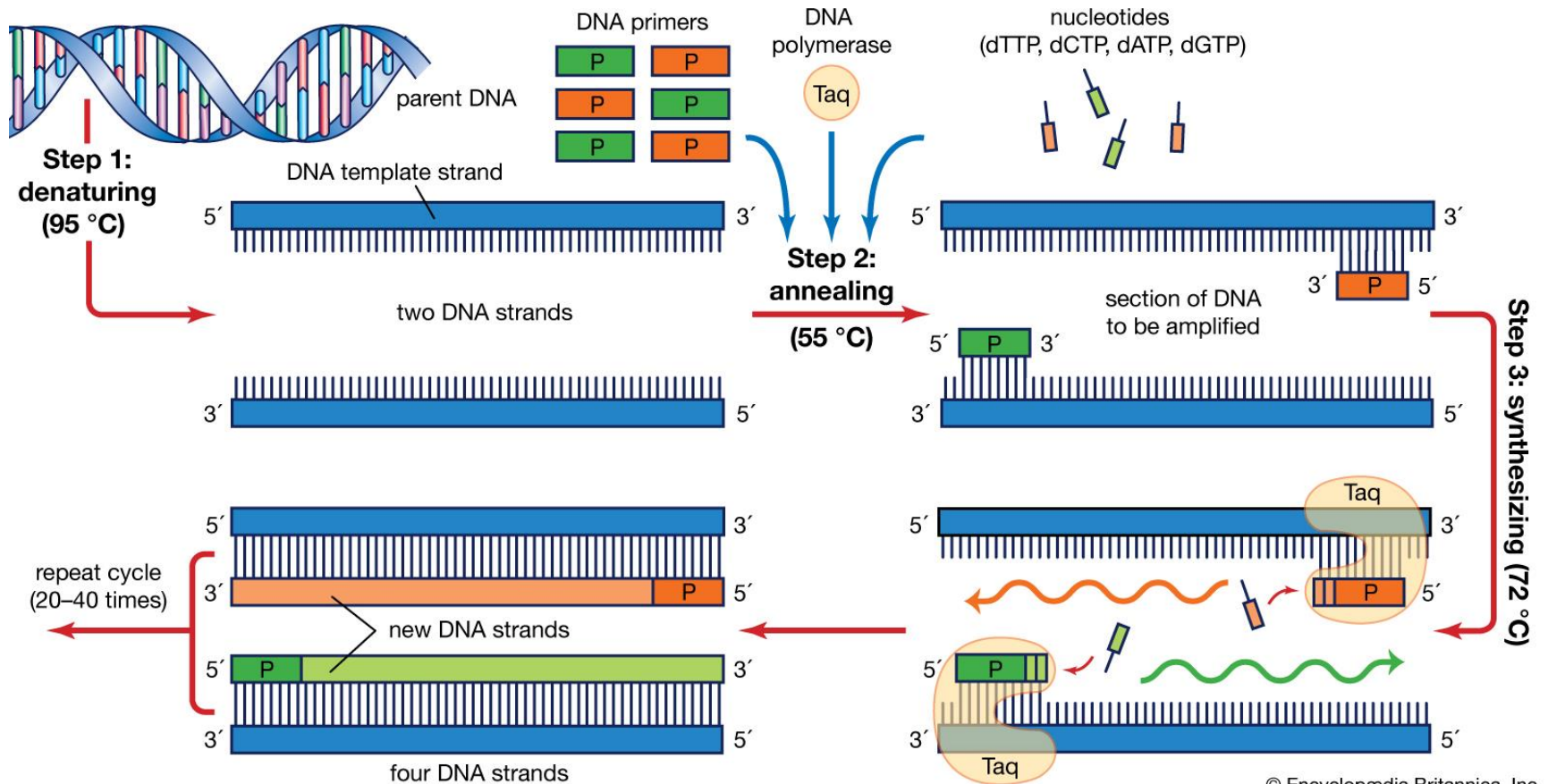


BamHI

Recognition Site:




Enzymatic DNA synthesis: polymerase chain reaction (PCR)



Amplification of up to 20 kbp DNA fragment from pre-existing template (genomic loci, cDNA library, cloned fragment etc.)

1 molekula DNA 



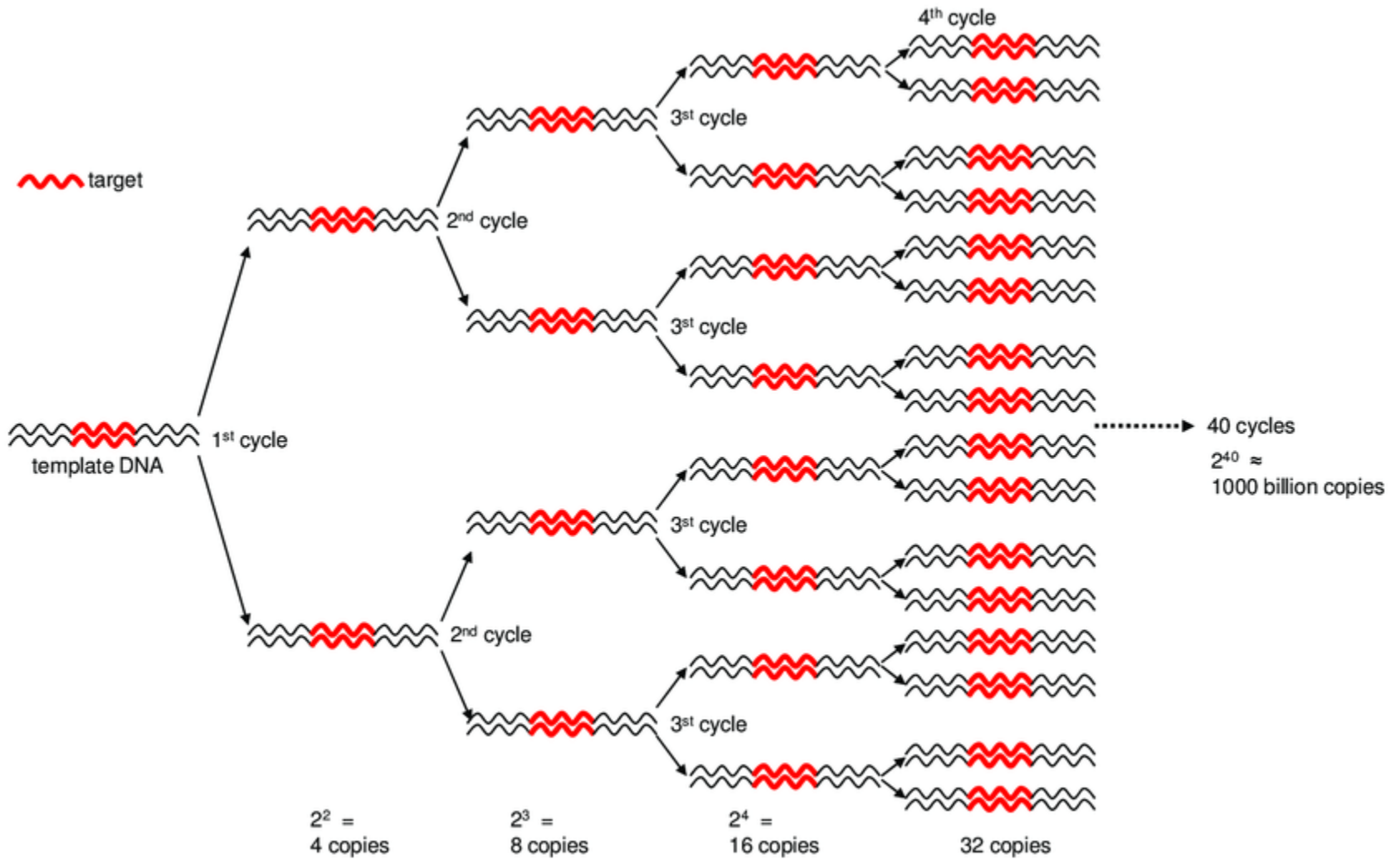
1. cyklus PCR: 2 molekuly DNA 

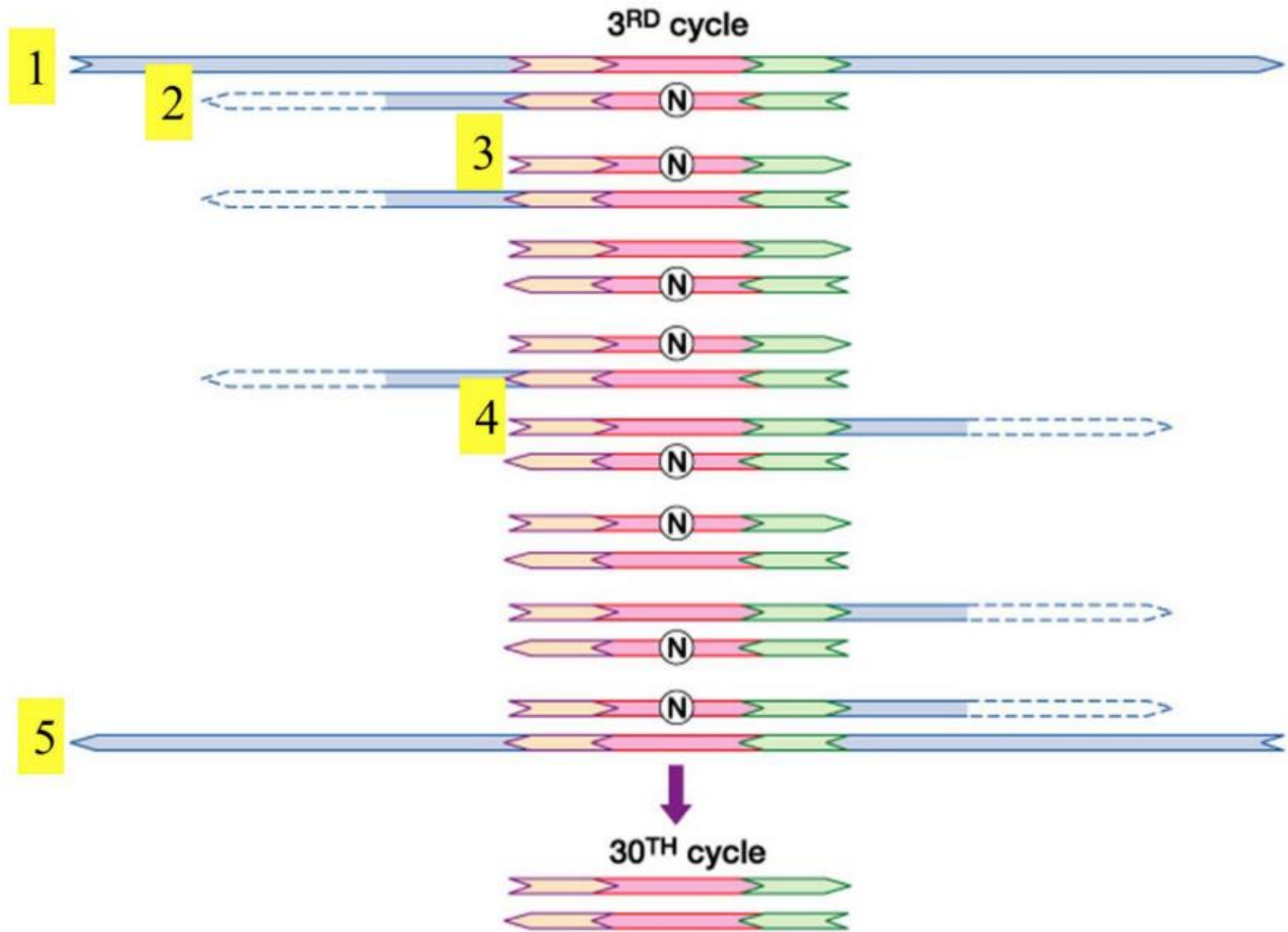




2. cyklus PCR: 4 molekuly DNA 

po n cyklech: 2^n molekul DNA





Predominant product because of exponential increase in copy number

Jaký je celkový počet polynukleotidových DNA řetězců po n cyklech?

Kolik je nově syntetizovaných polynukleotidových DNA řetězců po n cyklech?

Kolik je nově syntetizovaných cílových polynukleotidových DNA řetězců po n cyklech?

Kolik je cílových dvouvláknových DNA fragmentů po n cyklech?

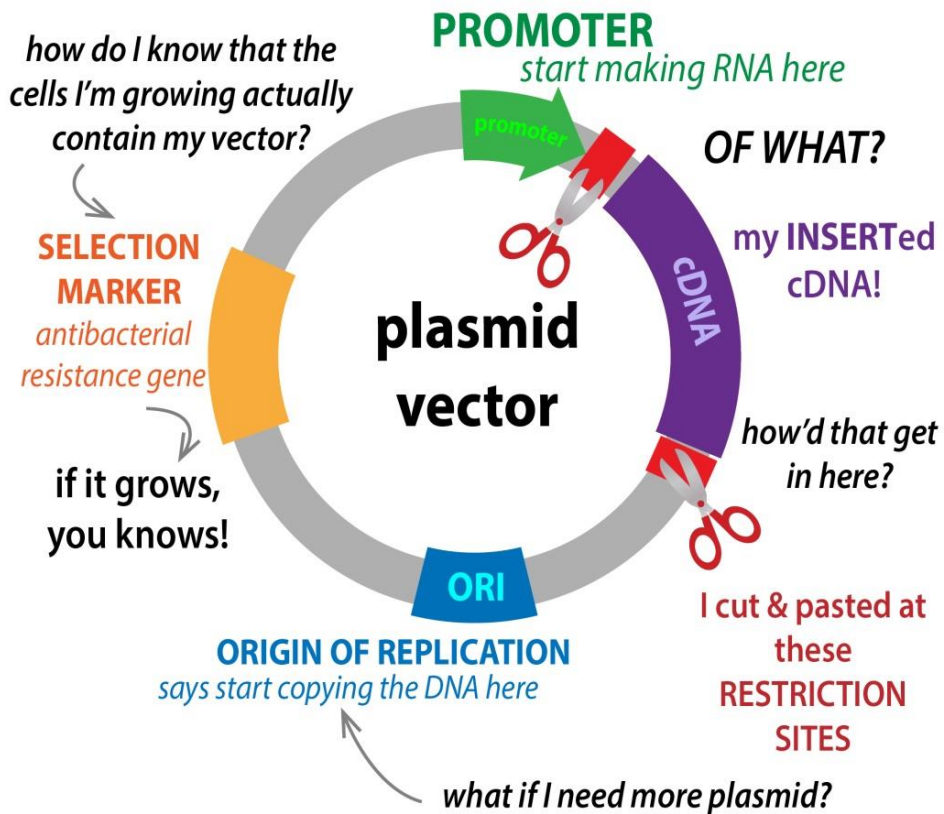
Kolik se spotřebuje primerů po n cyklech?

Kolik se zabuduje nukleotidů po n cyklech?

Kolik mikrogramů DNA se nasyntetizuje po n cyklech?

Questions

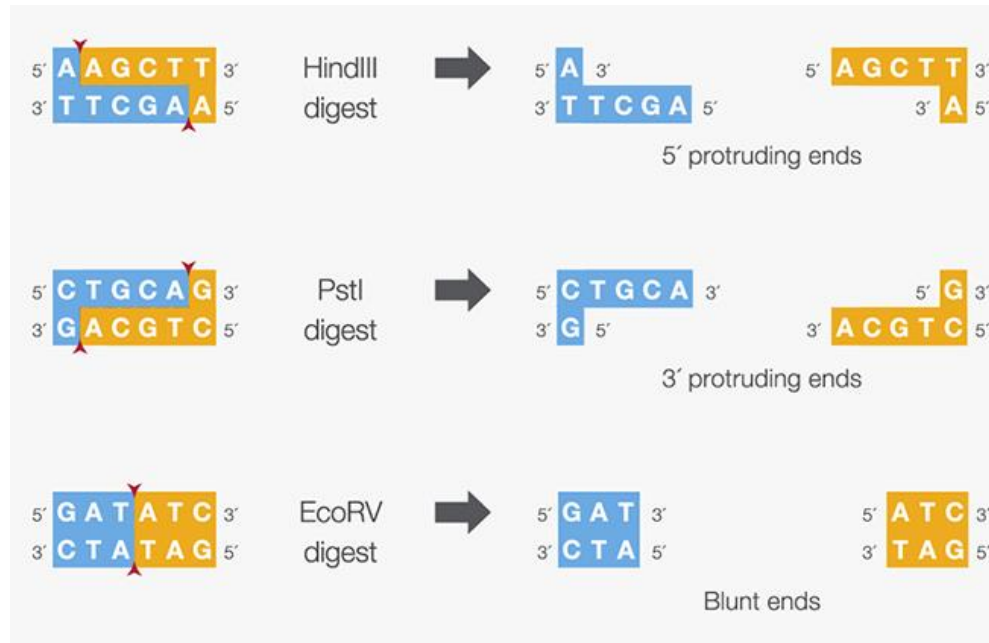
What is a plasmid?



- PLASMIDS are “extrachromosomal” (not part of the chromosomes), circular pieces of DNA.
- Similarly to chromosomes, they are double-stranded, which means they can easily be “unzipped” and copied (replicated).
- Plasmids use the host’s machinery (DNA polymerase), but they don’t have to wait for the host to divide to copy themselves → lots of copies of themselves.
- When the cell does divide, these copies will get split between the daughter cells, so they’ll inherit the plasmid as well.
- The plasmid can act as a VECTOR – a vehicle for taking genes we want to deliver into cells.

Always sequence your plasmid to double-check that the gene is correctly inserted!

Restriction endonucleases (restriction enzymes)



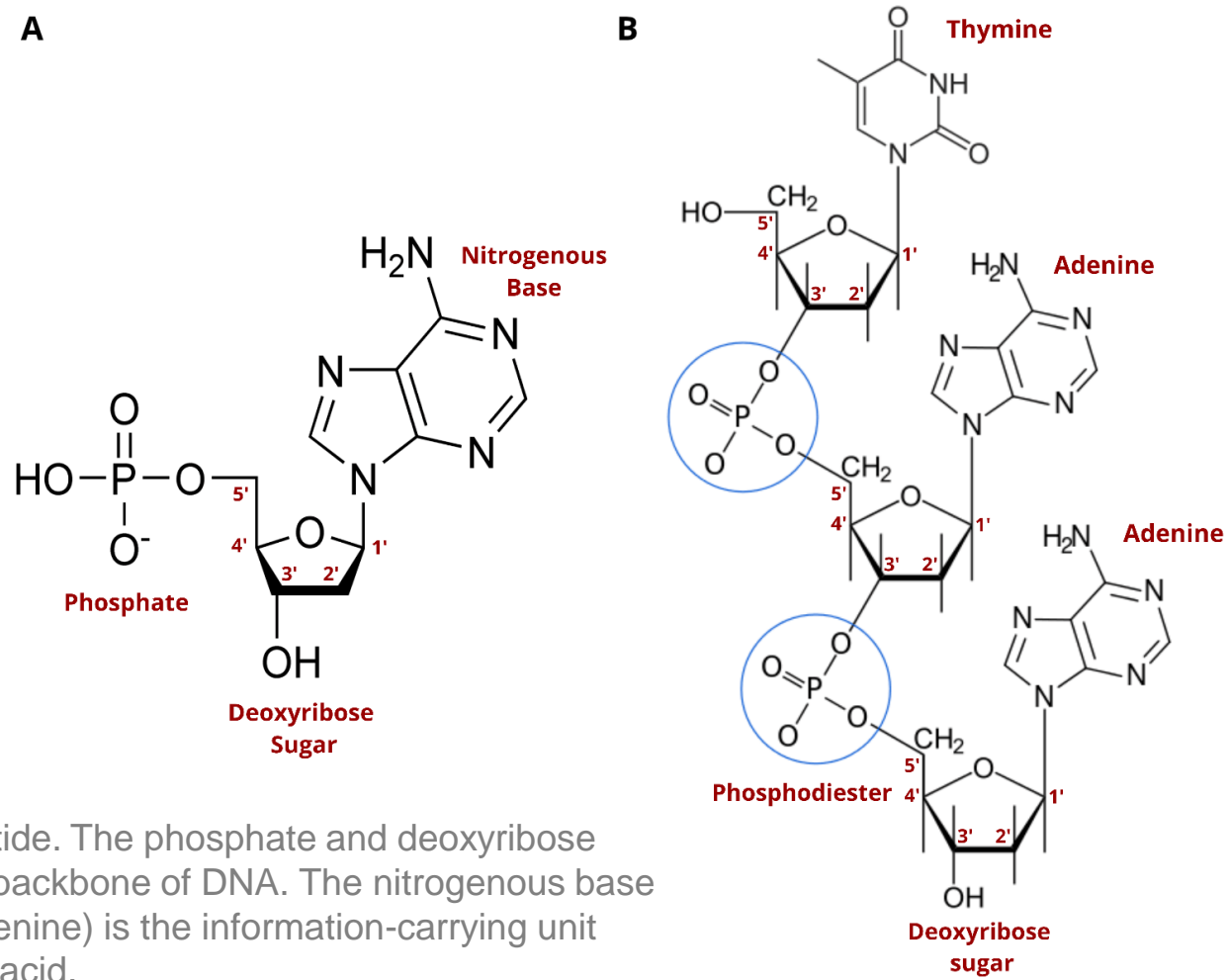
Type I enzymes cleave at sites remote from a recognition site; require both ATP and S-adenosyl-L-methionine to function; multifunctional protein with both restriction and methylase activities.

Type II enzymes cleave within or at short specific distances from a recognition site; most require magnesium; single function (restriction) enzymes independent of methylase.

Type III enzymes cleave at sites a short distance from a recognition site; require ATP (but do not hydrolyze it); S-adenosyl-L-methionine stimulates the reaction but is not required; it exists as part of a complex with a modification methylase.

Type IV enzymes target modified DNA, e.g. methylated, hydroxymethylated and glucosyl-hydroxymethylated DNA.

Structure of DNA

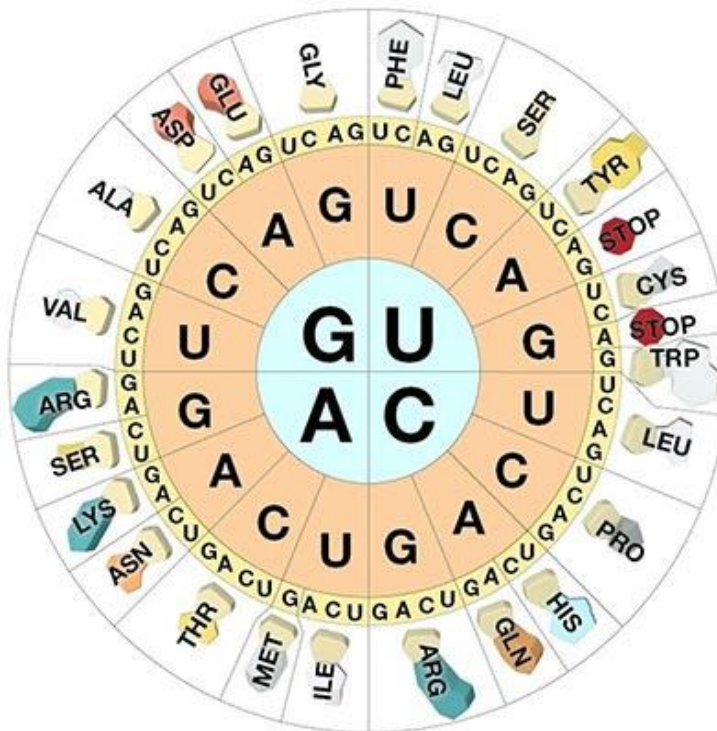


A) A single nucleotide. The phosphate and deoxyribose sugar form the backbone of DNA. The nitrogenous base (in this case adenine) is the information-carrying unit of each nucleic acid.

B) The structure of single-stranded DNA. In nature, enzymes form phosphodiester bonds (blue circles) that link the 5th position and 3rd position of adjacent deoxyribose sugars. Due to the modular nature of nucleotides, this chain can grow indefinitely.

Redundancy of the genetic code

- Degeneracy of codons is the redundancy of the genetic code, exhibited as the multiplicity of three-base pair codon combinations that specify an amino acid
- The genetic code is degenerate mainly at the third codon position
- The genetic code consists of 64 triplet codons specifying 20 canonical amino acids and 3 stop signals



		Second Letter				
		T	C	A	G	
First Letter	T	TTT } Phe TTC } TTA } Leu TTG }	TCT } TCC } Ser TCA } TCG }	TAT } Tyr TAC } TAA Stop TAG Stop	TGT } Cys TGC } TGA Stop TGG Trp	T C A G
	C	CTT } CTC } Leu CTA } CTG }	CCT } CCC } Pro CCA } CCG }	CAT } His CAC } CAA Gln CAG }	CGT } CGC } Arg CGA } CGG }	T C A G
	A	ATT } ATC } Ile ATA } ATG Met	ACT } ACC } Thr ACA } ACG }	AAT } Asn AAC } AAA Lys AAG }	AGT } Ser AGC } AGA } Arg AGG }	T C A G
	G	GTT } GTC } Val GTA } GTG }	GCT } GCC } Ala GCA } GCG }	GAT } Asp GAC } GAA Glu GAG }	GGT } GGC } Gly GGA } GGG }	T C A G

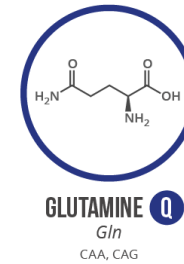
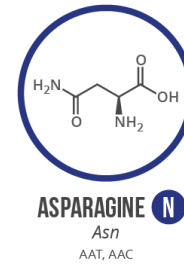
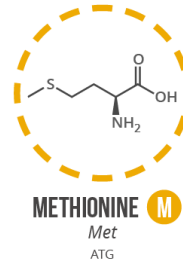
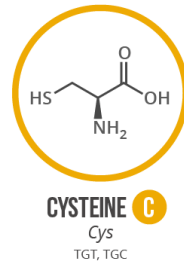
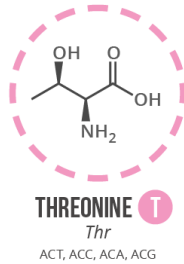
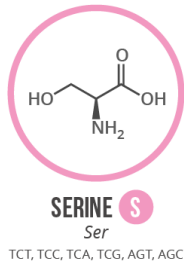
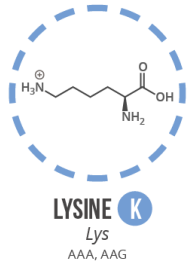
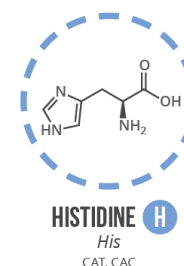
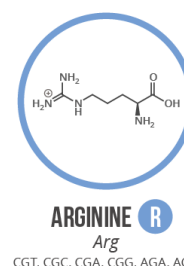
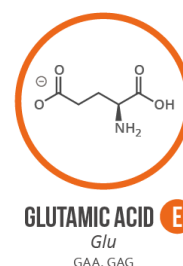
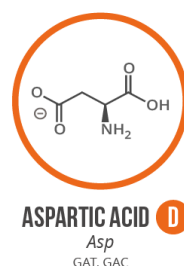
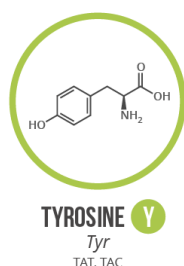
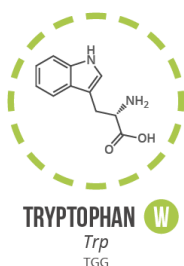
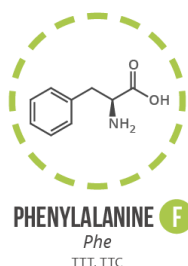
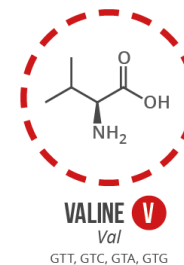
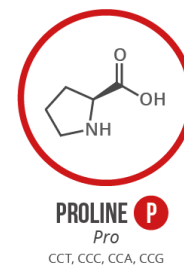
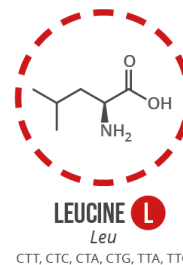
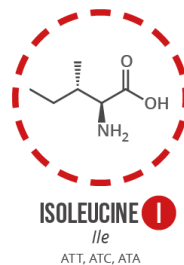
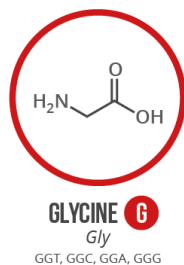
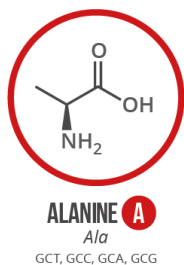
A GUIDE TO THE TWENTY COMMON AMINO ACIDS

AMINO ACIDS ARE THE BUILDING BLOCKS OF PROTEINS IN LIVING ORGANISMS. THERE ARE OVER 500 AMINO ACIDS FOUND IN NATURE - HOWEVER, THE HUMAN GENETIC CODE ONLY DIRECTLY ENCODES 20. 'ESSENTIAL' AMINO ACIDS MUST BE OBTAINED FROM THE DIET, WHILST NON-ESSENTIAL AMINO ACIDS CAN BE SYNTHESISED IN THE BODY.

Chart Key: ● ALIPHATIC ● AROMATIC ● ACIDIC ● BASIC ● HYDROXYLIC ● SULFUR-CONTAINING ● AMIDIC ○ NON-ESSENTIAL ○ ESSENTIAL

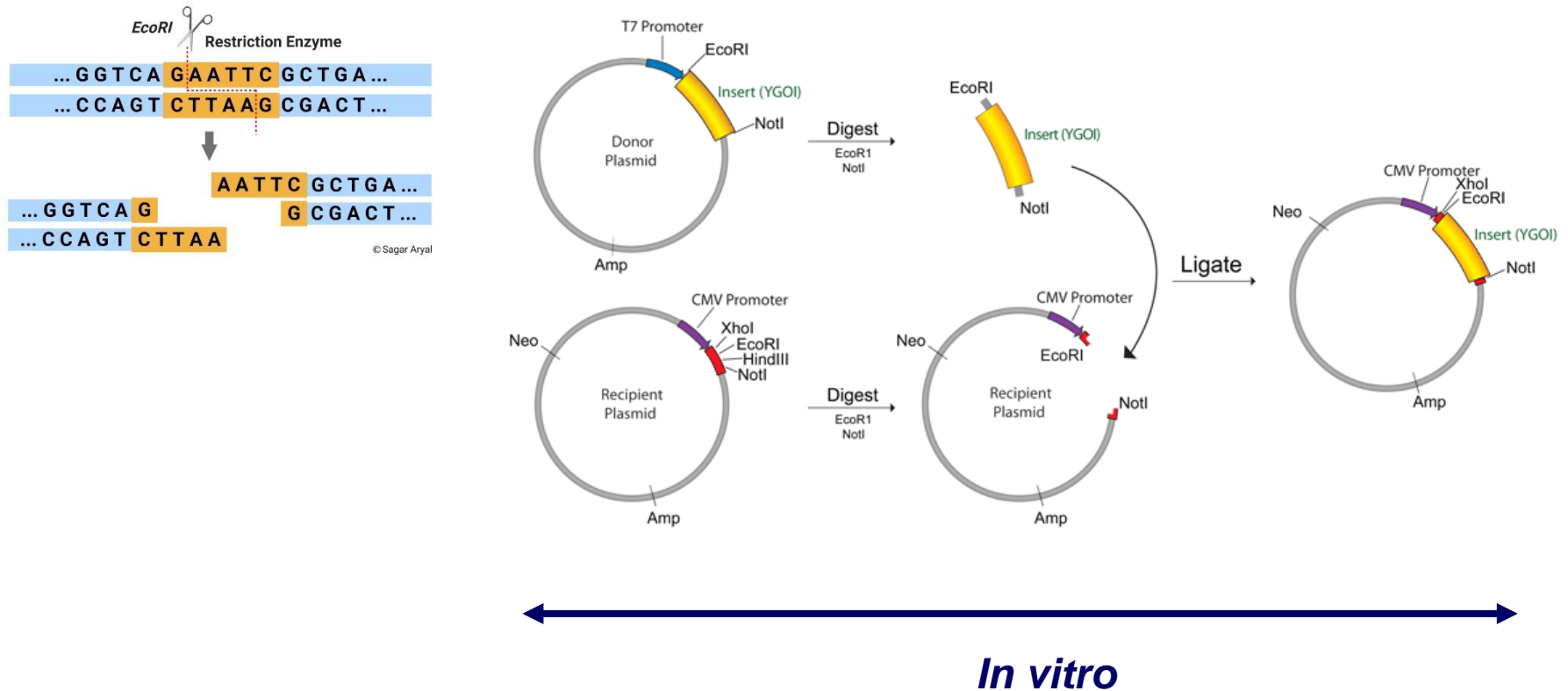
Chemical Structure
single letter code

NAME **A**
three letter code
DNA codons



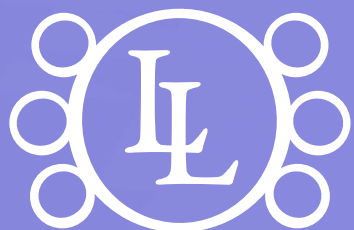
Note: This chart only shows those amino acids for which the human genetic code directly codes for. Selenocysteine is often referred to as the 21st amino acid, but is encoded in a special manner. In some cases, distinguishing between asparagine/aspartic acid and glutamine/glutamic acid is difficult. In these cases, the codes asx (B) and glx (Z) are respectively used.

Restriction cloning



The production of exact copies of a particular gene or DNA sequence using genetic engineering techniques is called gene cloning.

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