# POLYMERASE CHAIN REACTION (PCR)

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### Definition

It is a method of making multiple copies of a DNA sequence, involving repeated reactions with a polymerase.

(Oxford Languages)

It is a laboratory technique used to amplify DNA sequences. (NIH – National Human Genome Research Institute)

It is a method widely used to rapidly make millions to billions of copies (complete copies or partial copies) of a specific DNA sample, allowing scientists to take a very small sample of DNA and amplify it (or a part of it) to a large enough amount to study in detail.

(Wikipedia)



### Introduction

- By definition, it is a molecular biological technique used to amplify a single copy or a few copies of a specific DNA segment across several orders of magnitude, generating thousands to millions of copies of a particular DNA sequence.
- This revolutionary method was developed by Kary Mullis in the 1980s for which he was awarded with Nobel Prize in Chemistry in 1993.
- It is based on using the ability of DNA polymerase to synthesize new strand of DNA complementary to the offered template strand.
- It involves using short DNA sequences called primers to select the portion of the genome to be amplified.
- The temperature of the sample is repeatedly raised and lowered to help a DNA replication enzyme copy the target DNA sequence.
- The technique can produce a billion copies of the target sequence in just a few hours.



- Thermal cycler machine
- DNA template
- Primers
- DNA polymerase
- Nucleotides (dNTPs or deoxynucleotide triphosphate)
- Buffer solution
- Divalent cations





#### Thermal cycler machine

- Thermal cycler or thermocyclers, or PCR machine, are instruments used to amplify DNA and RNA samples by the polymerase chain reaction.
- It can be programmed to alter the temperature of the reaction every few minutes to allow DNA denaturing and synthesis.
- It has a *thermal block* with holes where tubes holding the PCR reaction mixtures can be inserted.
- The cycler then raises and lowers the temperature of the block in discrete, pre-programmed steps.



#### Thermal cycler.



### An older, three-temperature thermal cycler for PCR

### A 96-well plate modern thermal cycler for PCR





#### The Kary Mullis' Thermal Cycler





http://hpc.ilri.cgiar.org/beca/training/IMBB/lectures



### Thermal cycler machine today

- Standard PCR
- Nested PCR
- Touch down PCR
- Sequencing PCR
- Intersequence-specific PCR (ISSR)
- Many others
  - Real-time PCR machine
  - Digital PCR systme (dPCR)
  - Multi-block PCR thermal cyclers
  - 384-Well PCR thermal cyclers





### **DNA template**

- It is the DNA molecule that contains the segment of DNA to be amplified; it is basically a target sequence.
- Amount of DNA template is determined empirically; if too little template is used, primers may not be able to find their complementary sequences and if too much template is used, it may lead to increased mispriming events.





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- Generally, used concentration is 100-250ng for mammalian genomic DNA and 20ng for linearized plasmid DNA (circular plasmid DNA is slightly less efficiently amplified) per 50µl reaction.





### **DNA Primers**

- Primers are short synthetic DNA strands of about 18 to 25 nucleotides complementary to 3' end of the template strand.
- > DNA polymerase starts synthesizing new DNA from the 3' end of the primer.

#### Two types:

Forward primer: It is complimentary to the 3' end of antisense strand (3' - 5').

**Reverse primer:** It is complimentary to the 3' end of the sense strand (5' - 3').

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The concentration of each primer should be between 0.1 and 0.5 μm (0.2 μm is also said to produce satisfactory results.



### **DNA Primers**

It we consider the sense strand (5' – 3') of a gene, for designing primers, then forward primer is the beginning of the gene and the reverse primer is the reverse primer is the reverse-compliment of the 3' end of the gene.





### **DNA Primers – Designing**

### **Consideration**

- Typical primers are 18-28 nucleotides in length
- 50-60% GC compositon
- Have a balanced distributon of G/C and A/T domains
- No long strings of a single

#### <u>Good</u>

5' ATGCACTCAGACGTACAACGTGAC 3'

24 bases; AT: 12 GC: 12 (50% GC); Balanced distribution; Ta = 65°C

#### <u>Bad</u>

5' AAAACAAACGATTTTTT 3'

17 bases; AT: 14; GC: 3 (18 % GC); Unbalance distribution; Ta = 38°C





#### **DNA Primers – Designing**

#### **Characteristics**

Resources for primer design – Primer design tool

- CLC Workbench ◀ our focus
- Prime-BLAST
- PrimerQuest
- Primer3
- Primer3 plus
- PrimerZ
- PerlPrimer
- OligoPerfect
- OLIGO
- GenScript Real-time PCR (TaqMan) Primer Design
- AutoPrime
- RExPrimer
- BatchPrimer3
- Eurofins Genomics' Primer Design Tools





#### **DNA polymerase**

It sequentially adds nucleotides complimentary to template strand at 3'-OH of the bound primers and synthesizes new strnds of DNA complementary to the target sequence.

The most commonly used DNA polymerase is Taq DNA polymerase.

#### Taq DNA polymerase

- This enzyme is extremely heat resistant with a half-life of 40 minutes at 95°C (Optimal temp range is 75 80°C).
- It was originally isolated from thermophilic bacterium of the Deinococcus-Thermus group located near the Lower Geyser Basin of Yellowstone National Park by Thomas D. Brock and Hudson Freeze, in **1969**.
- Pfu DNA polymerase (isolated from Pyrococcus furiosus) is also used widely because of its higher fidelity (accuracy of adding complimentary nucleotide).





#### dNTPs or deoxynucleotide triphosphate (nucleotides)

All types of nucleotides are "building blocks" for new DNA strands and essential for reaction. It includes:

Deoxyadenosine triphosphate (dATP) Deoxyguonosine triphosphate (dGTF) Deoxycytidine triphosphate (dCTP) Deoxythymidine triphosphate (dTTP) Deoxyuridine triphosphate (dUTP)







### **Divalent cation**

- □ Typically magnesium (Mg) or manganese (Mn) ions are used.
- Mg<sup>2+</sup> is the most commonly used divalent cation Mn<sup>2+</sup> is used for PCR-mediated DNA mutagenesis (higher Mn<sup>2+</sup> con. Increases the error rate during DNA synthesis).

#### Magnesium chloride (MgCl<sub>2</sub>)

- It affects primer annealing and template denaturation, as well as enzyme activity.
- An excess of Mg<sup>2+</sup> gives non-specific amplification products, while low Mg<sup>2+</sup> yield lesser amount of desired product.
- It is also an essential cofactor for the synthesis and salvage of purine and pyrimidine nucleotides.
- It also affect their interaction with proteins and other ligands.



### **Polymerase buffer**

- All DNA polymerases are supplied with their own optimal polymerase buffer.
- However, buffer can also be prepared in lab.

#### Constituent of standard Polymerase buffer is:

- 10 mM Tris-HCl pH 8.3 (at room temperature)
- 50 mM KCl
- ✓ 1.5 mM MgCl<sub>2</sub>



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### Procedure of PCR

#### It is a multistep process which consists of following steps:

*Initialization:* It consists of heating the reaction chamber to a temperature of 94– 96 °C (94 °C) for 1 to 5 min. It causes separation of DNA double helix – *Initial denaturation.* 

**Denaturation:** It is the first regular cycling event which consists of heating the reaction chamber to a temperature of 94–98 °C for 20 – 30 seconds. It causes melting, or denaturation of DNA by breaking the hydrogen bonds between complimentary bases, yielding two single-stranded DNA molecules. Each strand acts as a template for synthesis of complimentary strand.

<u>Annealing</u>: Temp is lowered to 50 - 65 °C and kept the reaction mixture at this temp for 20 - 30 sec. This allows hybridization of primers to the primary strand.

**Extension/Elongation:** Reaction chamber is heated to 72 °C for 1 min. In this step, Taq DNA polymerase synthesize a new DNA stand complimentary to the DNA template strand by adding free dNTPs from the reaction mixture.

<u>Final extension</u>: This step is optional. It is performed at a temperature of 70 - 74 °C (72 °C) for 7 -10 min. It ensures that any remaining single-stranded DNA is fully elongated.

Store: Now the final product is stored for analysis. Generally stored at 4 - 15 °C



### Procedure of PCR







### Procedure of PCR



#### Temperature changes during each cycle of PCR





### Calculation of copies in PCR

The number of double stranded DNA pieces is doubled in each cycle, so that after n cycles you have 2<sup>n</sup> (2 to the n<sup>th</sup> power) copies of DNA.

Cycle	Calculation	Number of copies
1st	2 <sup>1</sup>	2
2 <sup>nd</sup>	2 <sup>2</sup>	4
3 <sup>rd</sup>	2 <sup>3</sup>	9
4 <sup>th</sup>	24	16
5 <sup>th</sup>	2 <sup>5</sup>	32
10 <sup>th</sup>	2 <sup>10</sup>	1,024
20 <sup>th</sup>	220	10,48,576

However it also depends on efficiency of PCR.





### **Overview of PCR**







# Further reading

- The History of PCR (RU 9577)
- http://siarchives.si.edu/research/videohistory\_catalog 9577.html
- PCR: A Revolutionary Invention <u>https://www.nature.com/scitable/blog/bio2.0/pcr\_a\_re</u> <u>volutionary\_invention/</u>
- How to design primers for PCR experiments Tips and considerations for sensitive PCR assays.
  https://www.zymoresearch.com/blogs/blog/how-todesign-primers-for-pcr-experiments

