


PV92 PCR


- Extract human genomic DNA and prepare samples for PCR
- PCR Cycle samples
 - Amplify PV92 locus of chromosome 16
- Agarose gel analysis
- Genotype individuals
- Hardy-Weinberg analysis of population genetics



What Is PCR?

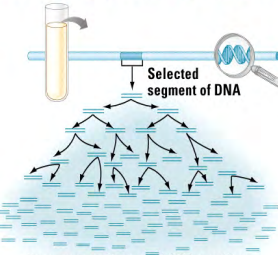
Polymerase Chain Reaction

- DNA replication gone crazy in a test tube!
- Makes millions of copies of a specific **target sequence** from template DNA
- Uses heat-resistant *Taq* polymerase from *Thermus aquaticus*



Polymerase Chain Reaction

With PCR, any specific segment—the target sequence—within a DNA sample can be copied many times (amplified) completely *in vitro*!

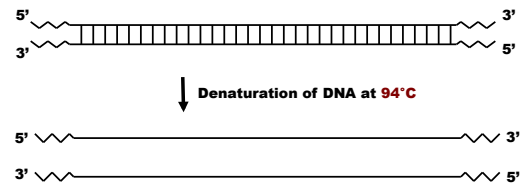


All you need:

- A heat-block that can rapidly and precisely change temperature (Thermocycler)
- Primers bracketing the sequence of interest
- A special heat-stable DNA-polymerase from a bacteria inhabiting hot-springs
- dNTPs
- Buffer & cofactors for the polymerase
- Source DNA as template

PCR Cycle – Step 1: Denaturing Template DNA

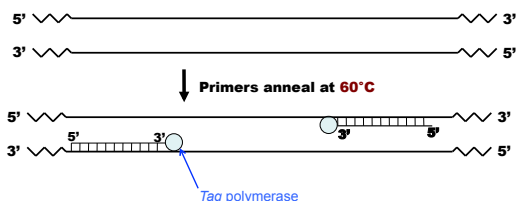
- Heat causes DNA strands to separate (“melt”)



↓ Denaturation of DNA at 94°C

PCR Cycle – Step 2: Annealing Primers

- Primers bind to the template sequence
Actual temp used depends upon primer specificity
- *Taq* polymerase binds to double-stranded substrate

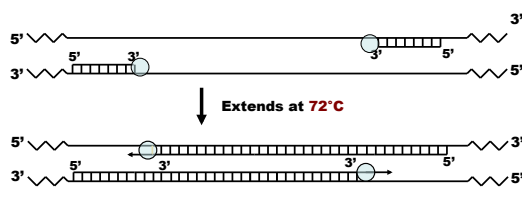


↓ Primers anneal at 60°C

Taq polymerase

PCR Cycle – Step 3: DNA Extension

- *Taq* polymerase adds dNTPs to extend DNA polymer from 3'-end of each primer
- DNA target sequence is replicated



↓ Extends at 72°C

Polymerase Chain Reaction

- The PCR procedure**

TECHNIQUE

Each temperature cycle:

- High heat (94°C)
 - “melt” DNA
 - dsDNA → ssDNA
- Low heat (45–60°C)
 - Allow primers to anneal to DNA
- Medium heat (72°C)
 - Reduce nonspecific binding
 - Taq polymerase polymerizes new DNA on template DNA
- Return to Step 1.
 - New DNA from Step 3 also used as template.

Polymerase Chain Reaction

- The PCR procedure**

TECHNIQUE The starting materials for PCR are double-stranded DNA containing the target nucleotide sequence to be copied, a heat-resistant DNA polymerase, all four nucleotides, and two short, single-stranded DNA molecules that serve as primers. One primer is complementary to one strand at one end of the target sequence; the second is complementary to the other strand at the other end of the sequence.

RESULTS During each PCR cycle, the target DNA sequence is doubled.

- Theoretical yield: 2ⁿ-fold (n = # of cycles)**
- By the end of the third cycle, one-fourth of the molecules correspond exactly to the target sequence, with both strands of the correct length. (See white boxes in Cycle 3.)
- After 20 or so cycles, the target sequence molecules outnumber all others by a billionfold or more.

Figure 20.7

POLYMERASE CHAIN REACTION

DNA region of interest.

- DNA is denatured. Primers attach to each strand. A new DNA strand is synthesized behind primers on each template strand.
- Another round: DNA is denatured, primers are attached, and the number of DNA strands are doubled.
- Another round: DNA is denatured, primers are attached, and the number of DNA strands are doubled.
- Another round: DNA is denatured, primers are attached, and the number of DNA strands are doubled.
- Continued rounds of amplification swiftly produce large numbers of identical fragments. Each fragment contains the DNA region of interest.

Polymerase Chain Reaction

- Animation from Harvey Lodish, et al *Molecular Cell Biology*
- http://bcs.whfreeman.com/lodish6/pages/bcs-main.asp?y=category&s=000108&n=05000&i=05010_06&o=005101_00520005300005400055000560005700058000590006000610000100002000030004000500060007001100020003000400050006000700080009001000110001200013000140001500016000170001800019000200002100022000230002400025000260002700028002900030003100032000330003400035000360003700038003900040004100042000430004400045000460004700048004900050005100052000530005400055000560005700058000590006000610006200063000640006500066000670006800069000700071000720007300074000750007600077000780079000800081000820008300084000850008600087000880008900090009100092000930009400095000960009700098009900100000

Chromosome 16 PV92 PCR Procedure Overview

- Obtain hair.
 - Milker cells for genomic DNA extraction
- Add sample to InstaGene™ matrix in micro test tube.
 - Or rinse mouth to obtain cheek cells for DNA extraction
- Incubate at 56°C for 10 minutes, then repeat vigorous agitation.
- Incubate at 100°C for 8 minutes, then repeat vigorous agitation.
- Centrifuge samples for 8 minutes to pellet matrix.

DNA template preparation

DNA amplification

The positive controls consist of the 96-well PCR plate containing the three positive control templates.

- DNA molecular mass that is greater than that of the InstaGene matrix.
- InstaGene matrix.
- Reaction buffer.
- Reaction primers (1-2).
- Reaction dNTPs (3-4).
- Reaction temperature.

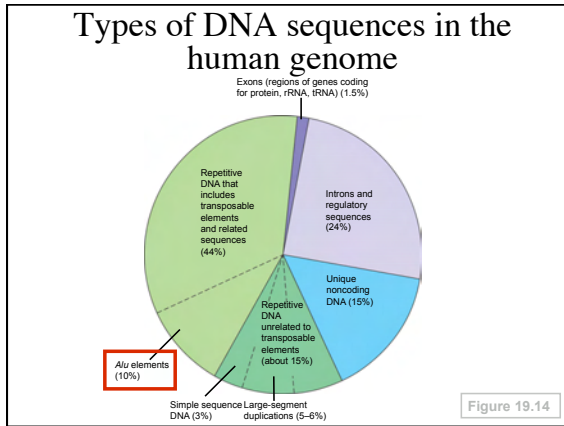
Electrotransfer PCR samples on agarose gels at 100V for 30 minutes. Stain with Fast Blue® DNA stain or ethidium bromide.

Determine student genotypes for A1u insertion and perform Hardy-Weinberg analysis on class results.

Protocol Highlights: Genomic DNA Extraction

- InstaGene™ - Chelex-cation exchange resin; binds cellular magnesium ions
- 56°C - loosens connective tissue and inactivates DNases
- 100°C - ruptures cell membranes and denatures proteins

InstaGene Extraction:
InstaGene matrix binds released cellular Mg²⁺



Alu Repeats

- Classified as SINEs (Short Interspersed Repetitive Element) — ~300 bp; highly conserved
- Mobilized by an RNA polymerase-derived intermediate (retroposition)
- Approx. 500,000 *Alu* copies per haploid genome, representing about 5% of the genome
- Named for the *Alu* I restriction site within the element

