

MOLECULAR CELL PHYSIOLOGY - Recombinant DNA Technology

- I. What is recombinant DNA technology?
 - A. A set of techniques used to combine genes from different sources and transfer those genes into an organism where they may be expressed
 - 1. A targeted and potentially permanent manipulation of organisms
- II. Why is recombinant DNA technology important?
 - A. It provides knowledge about how organisms "work"
 - B. It offers the potential to make organisms "new and improved"
 - C. It enables understanding and circumvention of disease
 - D. It provides a new (and "natural") way of making commercial products
- III. Technologies used to understand genes
 - A. Biochemical purification and analysis of a protein based on function (old)
 - B. Classical genetic analysis by mapping of genes defined by mutations (old)
 - C. Recombinant DNA technology (new)
 - 1. Enabled sequencing of human genome (6×10^9 base pairs)
- IV. From protein to gene
 - A. Isolate protein on basis of molecular function
 - B. Determine partial amino acid sequence of protein
 - C. Synthesize oligonucleotides that correspond to amino acid sequence
 - D. Use oligonucleotides as probes to screen genomic library
 - E. Sequence isolated gene
- V. From gene to protein
 - A. Isolate genomic clone corresponding to desired trait (often a mutant or altered gene)
 - B. Use genomic DNA to isolate cDNA for the mRNA coded by the gene
 - C. Sequence the cDNA and then determine amino acid sequence
 - D. Compare amino acid sequence to known similar proteins to learn about function
 - E. Use expression vector to produce the encoded protein
- VI. What is the basis of recombinant DNA technology?
 - A. Variation occurs naturally
 - 1. In meiosis, crossing over (prophase I) and recombination (metaphase I)
 - B. Bacterial genes can transfer from cell to cell
 - 1. Transformation - Incorporation of new DNA from stuff around the cell (Griffith)
 - 2. Transduction - Transfer of DNA into bacteria by virus (bacteriophage)
 - 3. Conjugation - Reproduction or "mating" of bacterial cells
- VII. Example of how small amounts of recombinant DNA are "manufactured" through use of plasmids
 - A. Plasmids carrying recombinant DNA can be used to infect bacteria such as *E. coli*
 - B. These plasmids contain a small, circular piece of DNA
 - 1. Plasmid can replicate itself within a cell or simply carry DNA (vector)
 - 2. Plasmid usually consists of replication origin, section that encodes for selection (like antibiotic resistance), and the section of DNA (20 kb) to be cloned
 - C. The plasmid can be modified by inserting a gene of interest into it
 - 1. Once modified, the plasmid is said to carry the recombinant DNA
 - a) Plasmids are modified by "cutting" and "pasting" pieces of DNA
 - 1) Restriction enzymes (endonucleases) cut old DNA at special sequences
 - 2) Polylinkers, with several restriction sites, can be inserted into plasmid to help make "sticky ends"
 - 3) New DNA with "sticky ends" (complementary DNA) "fits in"
 - 4) DNA ligase pastes the new DNA into the old DNA to make recombinant DNA in the plasmid
 - D. Once the recombinant plasmid is made, it can be taken up by transformation into a cell
 - E. The cell then integrates the DNA, replicates it, and continues to make copies of the recombinant DNA or protein products

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- VIII. Larger amounts of recombinant DNA are “manufactured” through use of other techniques
- A. **λ** phage: cut up genomic DNA (25 kb pieces) and allow modified **λ** phage to take up pieces; phage then infects *E. coli*; this colonies which, essentially, constitute the genomic library
 - B. Cosmids (45 kb pieces): generate pieces using **λ** phage and then insert, under special conditions into a plasmid
 - C. P1 phage (100 kb pieces): takes advantage of the *E. coli* virus, bacteriophage P1, whose head can accommodate larger DNA molecules than the **λ** phage
 - D. Bacterial artificial chromosome (300 kb pieces): make use of a large *E. coli* plasmid called the F-factor
 - E. Yeast artificial chromosome (1000 kb pieces): in essence, eucaryotic cells are bigger
- IX. How is a gene of interest located?
- A. The genomic "library" (two techniques)
 1. Start with DNA and cut it up into pieces by restriction enzymes
 2. Use reverse transcriptase to transform mRNA into DNA - then cut into pieces
 - a) Very useful for finding only the DNA that is expressed in a certain cell also referred to as complementary “cDNA”
 - B. Screen the library with a radioactive probe from another source
 1. Denature genomic DNA to make single-stranded DNA and dry on to a membrane such as nitrocellulose - this leaves most base pairs available for hybridization
 - a) Can use petri dishes containing recombinant **λ** virion present in plaques on A lawn of *E. coli* – place a piece of nitrocellulose on dish to allow some plaques to absorb
 - b) Store original petri dishes (marked for orientation) in refrigerator
 - c) Incubate nitrocellulose membrane in alkaline to disrupt virions and release encapsulated DNA, then dry filter
 2. Incubate the nitrocellulose with a (radioactive) probe containing unique DNA (or RNA) that encodes for the gene of interest – at least 20 nucleotides in length
 - a) Sometimes many probes are needed because the amino acid sequence in question can be encoded by numerous nucleotide sequences (this is called a degenerate “probe”)
 - b) A database (expressed sequence tag) is available that contains specific DNA sequences in other organisms that correspond to amino acid sequences
 3. The probe adheres to one or more pieces of genomic DNA that probably contains the gene of interest
 4. Go back to the original DNA source (petri dishes) to get more purification, sequencing, and further analysis
 5. Obtain DNA from vector using restriction enzymes and isolate / purify with electrophoresis (polyacrylamide for up to 2000 bp; agarose for 500 to 20,000 bp; pulse-field electrophoresis for 20,000 to 10,000,000 bp)
 6. Visualize using ethidium bromide or radioactivity
 7. Determine DNA sequence by Maxam-Gilbert (cut and incubate with G, A+G, C+T, C) or Sanger (dideoxynucleotides used) method – both followed by electrophoresis
- X. Analyzing specific nucleic acids
- A. Southern blotting (named after Edwin Southern)
 1. Digest DNA with restriction enzyme and then use gel electrophoresis to separate
 2. Denature with alkali and then transfer to nitrocellulose filter
 3. Incubate filter with specific radiolabeled DNA probe
 - B. Northern blotting (named as such to make it different from Southern blotting)
 1. Denature RNA to make an unfolded, linear molecule
 2. Use gel electrophoresis to separate and transfer to nitrocellulose filter
 3. Incubate filter with radiolabeled DNA probe (quantitative method for RNAs)
 - a) Can also use endonucleases that act only on single-stranded polynucleotides for quantifying specific RNAs

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- XI. "Mass" producing protein from cloned DNA
 - A. Plasmid expression vectors carrying a strong, regulated promoter
 - B. Plasmid expression vectors carrying the T7 late promoter (T7 RNA polymerase of bacteriophage T7)
 - C. Because eukaryotes may encode post-translational modifications, eukaryotic expression vectors are sometimes needed
 - D. Examples
 - 1. Vaccines
 - 2. Insulin
 - 3. Granulocyte colony-stimulating factor (stimulates production of phagocytic white blood cells)
 - 4. Human growth hormone
 - 5. Factor VIII for blood clotting (hemophiliacs need it)
- XII. Polymerase chain reaction
 - A. Directly amplifies specific DNA sequences in a complex mixture where the ends of the sequence are known
 - 1. Synthetic oligonucleotides complementary to desired DNA are added in excess to total denatured DNA and temperature is lowered
 - 2. Increase temperature (along with addition of *Thermus aquaticus* DNA polymerase) to 95 °C to melt strands and then cool to 60 °C to anneal primers
 - 3. Proceed with heating and cooling cycles to continue doubling desired DNA
 - B. Applications of polymerase chain reaction
 - 1. Allows recovery and rapid amplification of DNA
 - a) Newly synthesized DNA can then be used for analytical purposes
 - b) Newly synthesized DNA improves specificity of probes and screening of DNA libraries
- XIII. Are there problems or risks of recombinant DNA technology?
 - A. Special labs and precautions are used to prevent harmful organisms from getting into the environment
 - B. There are bioethics committees that oversee what is being done with recombinant DNA
 - C. Use of information derived from DNA analysis and gene mapping can be debated (insurance company policies, for example)