

Overview

- DNA microarrays
 Looking for gene mutations with DNA
 Studying gene expression with mRNA.
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 Polymerase chain reaction (PCR)
 Reverse Transcription PCR
 Proteomics
 From amino acids to proteins.
 Traditional protein experimentation
 Protein microarrays types.
 Fabrication
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 Appendix
 Eukaryotic gene regulation & post-translation modification (PTM).
 Protein studies.
 Examples of DNA lab-on-a-chip devices.

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Looking for Gene Mutations...

- Start with a sample of DNA from blood, and a control sample without the specific gene mutation. 1.
- 2. 3.
- 4.
- mutation. Denature the DNA into two complimentary single-stranded molecules. Cut the strands into smaller fragments and label the patient's sample with green dye and the control with red dye. A DNA chip with synthetic probes for both normal and mutated genes is utilized. Both samples are placed into the chip and allow to hybridize (bind) to the synthetic DNA probes on the chip. 5.

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Fabrication of the DNA Microarray





GeneChip probe arrays are manufactured through a combination of photolithography and combinatorial chemistry.

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Polymerase Chain Reaction (PCR)

- An in vitro method of replicating small DNA sequences into millions of copies over a short period of time.
 Nanomolar quantities of DNA can be replicated within a few hours.
 PCR may be used for genetic testing in disease diagnosis, monitoring response to treatment, and tissue typing.
 A typical PCR requires:
 Two oligonucleotide primers;
 A thermally stable DNA polymerase;
 Supply of free nucleotides; and
 A simult amount of DNA sample that contains the sequence of

 - A small amount of DNA sample that contains the sequence of interest.
- The DNA fragment of interest must be known, so that short DNA primer fragments can be synthesized in advance, and are complimentary to the 3' end of each sample stand.

Denatu	ring and Annealing Steps	
a)	y regen	Desired sequence or target region to amplify.
b)	, <mark>111111111111111111111111111111111111</mark>	Denaturing by heating 95 C
c)	furning <mark>uning</mark> garant f juning <mark>uning</mark> annug j	Annealing by cooling to 60 C, and binding of primers.
Steven S. Saliterman		Carey, FA, Advanced Organic Chemistry 5th Ed., 2002





Elongation and Sequence Amplification					
f)	,	Elongation of the primed polynucleotide fragments completes the second cycle and gives 4 DNAs.			
g)	5 3	Among the eight DNAs formed in the third cycle are two having the structure shown. This structure increases disproportionately in the succeeding cycles.			
Steven S. S	Saliterman	Carey, FA, Advanced Organic Chemistry 5th Ed., 2002			

DNA Am	olificatio	n Results		
	Cycle Number	Total Number of DNAs	Number of DNAs Containing Only the Target Region	
	0	1	0	
	1	2	0	
	2	4	0	
	3	8	2	
	4	16	8	
	5	32	22	
	10	1,024	1,004	
	20	1,048,566	1,048,526	
	30	1,073,741,824	1,073,741,746	
Steven S. Saliterman			Care	ry, FA, Advanced Organic Chemistry 5th Ed., 2002

Reverse Transcription PCR

- *mRNA rather than DNA* is used as the starting template.
 Useful for RNA pathogens like HIV and Hepatitis C, gene expression, and differentiating viable from nonviable cells.
- Step 1
 - The enzyme reverse transcriptase uses the mRNA template to produce a complementary single-stranded DNA strand called cDNA in a process known as reverse transcription.
- Step 2
 - DNA polymerase is used to convert the single-stranded cDNA into double-stranded DNA. These DNA molecules can now be used as templates for a PCR reaction.

Matt Carter, Jennifer Shieh, in Guide to Research Tech on), 2015

Proteomics

- The study of all proteins, including their:
 Relative abundance
 - Relative abund
 Distribution
 - Distribution
 - Post-translational modifications
 - Functions
 - Interactions with other macromolecules
- In a given *cell or organism* within a *given environment* and at a *specific stage in the cell cycle*.

Pardanani, A. et al., Mayo Clinic Proce

redings 77(11), 2002

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Proteome...

- The entire collection of proteins, estimated to be more than 100,000.
- More proteins comprise a proteome than genes a genome. Alternative gene splicing of mRNA, Posttranslational modification (PTM).
- There is neither a one to one correlation of gene to protein, nor mRNA levels to proteins levels.
- mRNA levels to proteins levels.
 PTM and signal transduction play a major role in cell transformation, such as tumor cells.
 Specific genes are turned on or off at the onset of initiation, development, and progression of diseases such as cancer.
 Signal transduction, via growth factors, proteins, and peptides, plays a major role in cell transformation (e.g. carcinogenesis).



Post-translational Modification...

- Post-translational modification (PTM):
 - Phosphorylation, glycosylation, acetylation, ubiquitination, methylation etc.
 - PTM of proteins, not detected through RNA analysis, may occur at different stages of tumor development indicative of early or late events of transformation.
 - High throughput techniques may useful for screening and surveillance.

Traditional Protein Experimentation...

- A. Protein purification may be performed by affinity chromatography, ion exchange, or subcellular fractionation. Then, 2-dimensional gel electrophoresis allows isolation of single proteins which may be digested for MS.
 B. Use of 1-dimensional gel allows partial electrophoretic separation, requiring further separation such as high-performance liquid chromatography (HPLC).
 C. The entire mixture is digested to peptides
- The entire mixture is digested to peptides and the peptide mixture is resolved by multi-dimensional chromatography. C.











Areas of Interest...

- 1) Abundance proteomics:
 - Relative abundance of specific proteins in a given tissue under different conditions of health and disease.
- 2) Cell-mapping:
 - Intracellular signaling pathways and regulatory networks mediated through protein-protein interactions.
- 3) Structural proteomics:
 - Study of active sites and functional domains of proteins to better understand processes such as enzyme catalysis, protein stability and interaction with multimolecular complexes.

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Protein Microarray Types

- Forward-phase microarrays (or Analytical):
 - Proteins and peptides are immobilized for capturing antibodies.
 Antibodies, sugars or aptamers are immobilized and labeled
 - proteins are captured.
 Sandwich mode a labeled secondary antibody is used for
 - detection.
- Reverse phase microarrays:
 - Complex samples such as serum, plasma, or even tissues are immobilized in an array format and probed with antibodies to determine the differential amount of protein molecules in the screened samples.

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Protein Microarray Uses...

- Protein expression profiling.
- Studying the following:
 - Posttranslational modifications.
 - Protein-protein binding.
 - Drug interaction.
 - Protein folding.
 - Substrate specificity.
 - Enzymatic activity.
 - Interaction between proteins and nucleic acids.

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Factors Affecting Performance...

- Appropriate surface for the immobilization of either protein or antibody samples.
- Microarray patterning technique.
- Protein conformational changes with expression, purification or immobilization may alter their function or render them inactive.
- Charged surfaces, temperature, pH and solvents may denature some proteins, and therefore surfaces must be biocompatible to minimize denaturation.
- Protein instability may lessen shelf-life.

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Summary DNA microarrays Looking for gene mutations with DNA Studying gene expression with mRNA. Studying gene expression with mRNA. Polymerase chain reaction (PCR) Polymerase chain reaction (PCR) Polymerase transcription PCR Poteromise Poteromise Poteromise Poteromise Poteromise experimentation Poteromise experimentation Poteromise Poteromise Poteromise Poteromise Poteromise Poteromise experimentation Poteromise experimentation Poteromise Poteromise Poteromise Poteromise Poteromise Poteromise Poteromise experimentation modification (PTM). Poteromise Poteromise



















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A miRNA Detector - Wang et. al.

MicroRNAs

- Class of *endogenous and noncoding* RNAs. Crucial to gene regulation, such as cell development, differentiation, metabolism, and apoptosis.
- 18-25 nucleotides long, low content and easily degraded.
- Photoelectrochemical (PEC) biosensor for microRNA detection.
 - MoS₂/g-C₃N₄/black TiO₂ ternary heterojunction as the photoactive material and gold nanoparticles carrying Histostar antibodies (Histostar@AuNPs) for signal amplification.
 - Deposited on indium tin oxide (ITO) electrode.

Wang MH, Yin HS, Zhou YL, et al. Photoelectrochemical biosensor for microRNA detection based on a MoS2/g-C3N4/blac TiO2 heterojunction with Histostar@AuNPs for signal amplification. Biosensors & Bioelectronics. 2019;128:137-143.

• PEC biosensors - low background noise, easy operation and excellent sensitivity.

• The challenge in PEC biosensors is suppressing the recombination of photo-generated electrons and holes in semiconductor materials.

• Suppression will increases photo-excited carrier lifetimes. • Ternary heterojunctions improve light absorption

efficiency, promote electron transfer and extend the lifetime of charge carriers,

Wang MH, Yin HS, Zhou YL, et al. Photoelectrochemical biosensor for microRNA detection based on a MoS2/g-C3N4/ TiO2 heterojunction with Histostar@AuNPs for signal amplification. Biosensors & Bioelectronics. 2019;128:137-143.





Rapid Detection of Pathogens - Hügle et. al.

• Fluorescent E. coli were preconcentrated from an initial volume of 1 ml down to 5 ml. Subsequent steps allowed for measurement of organism specific sfGFP mRNA.

• Preconcentration was followed by



Fluorescence image of E. coli courtesy of Flora V. Romeo

Preconcernication was followed by thermoelectric lysis and on-chip gelelectrophoresis of released nucleic acids (including mRNAs) to remove larger fractions (e.g. gDNA or plasmids) and residual cell debris.
Specific sfGFP mRNA was then quantified in the extracts by RT-qPCR.

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