

# **Proteomics**

**The process of identifying, characterizing, and quantifying all expressed proteins in an organism under one or several conditions.**

# **Electrophoresis**

- **Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE)**
- **Isoelectric Focusing (IEF)**
- **2 Dimensional Gel Electrophoresis**
- **Free-Flow Electrophoresis**
- **Capillary Electrophoresis**

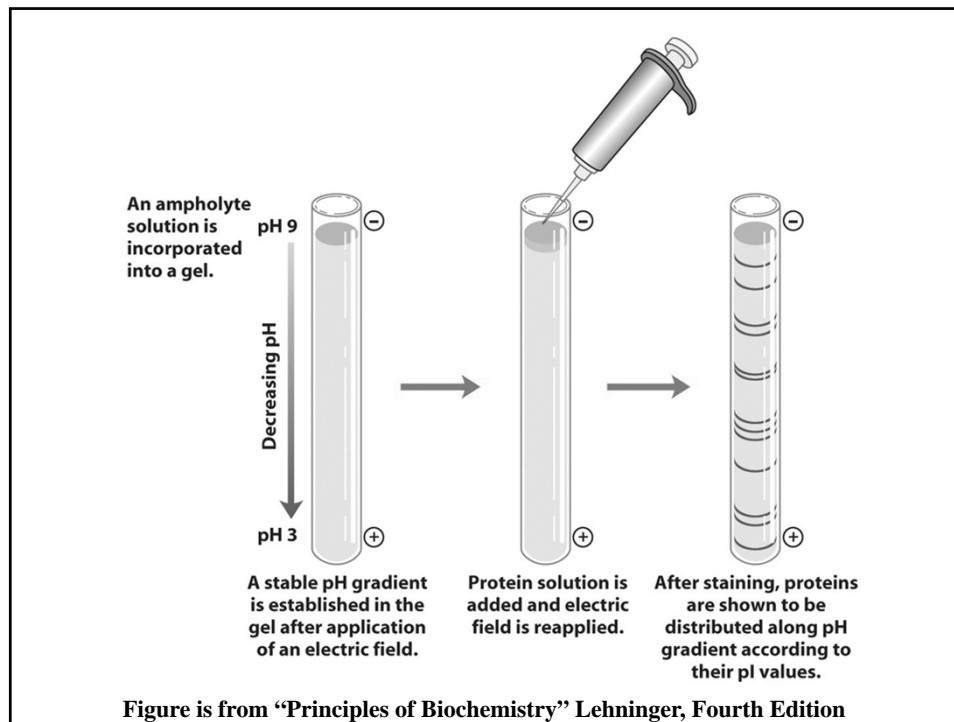
## SDS-PAGE

- **Separates proteins by size**
- **Denaturing gels**
- **Resolution dependent on**
  - Size of polyacrylamide gel
  - Concentration of acrylamide
    - one concentration or a gradient
  - Stacking of sample
- **Stain to visualize proteins**
  - Multiple stains available with varying sensitivity
    - Deep Purple, sypro ruby, sypro orange, silver

The diagram shows the chemical structures of acrylamide and bisacrylamide, and their polymerization into polyacrylamide and cross-linked polyacrylamide networks. Acrylamide is shown as  $\text{H}_2\text{C}=\text{CH}-\text{C}(=\text{O})\text{NH}_2$ . Bisacrylamide is shown as  $\text{H}_2\text{C}=\text{CH}-\text{C}(=\text{O})\text{NH}-\text{CH}_2-\text{NH}-\text{C}(=\text{O})\text{CH}=\text{CH}_2$ . The polymerization of acrylamide results in a linear chain of  $-\text{C}(\text{H}_2)-\text{C}(\text{H})(\text{C}(=\text{O})\text{NH}_2)-$  units. The polymerization of bisacrylamide results in a cross-linked network where the  $-\text{C}(\text{H}_2)-\text{C}(\text{H})(\text{C}(=\text{O})\text{NH}_2)-$  units are connected to each other via  $-\text{NH}-\text{CH}_2-\text{NH}-$  bridges.

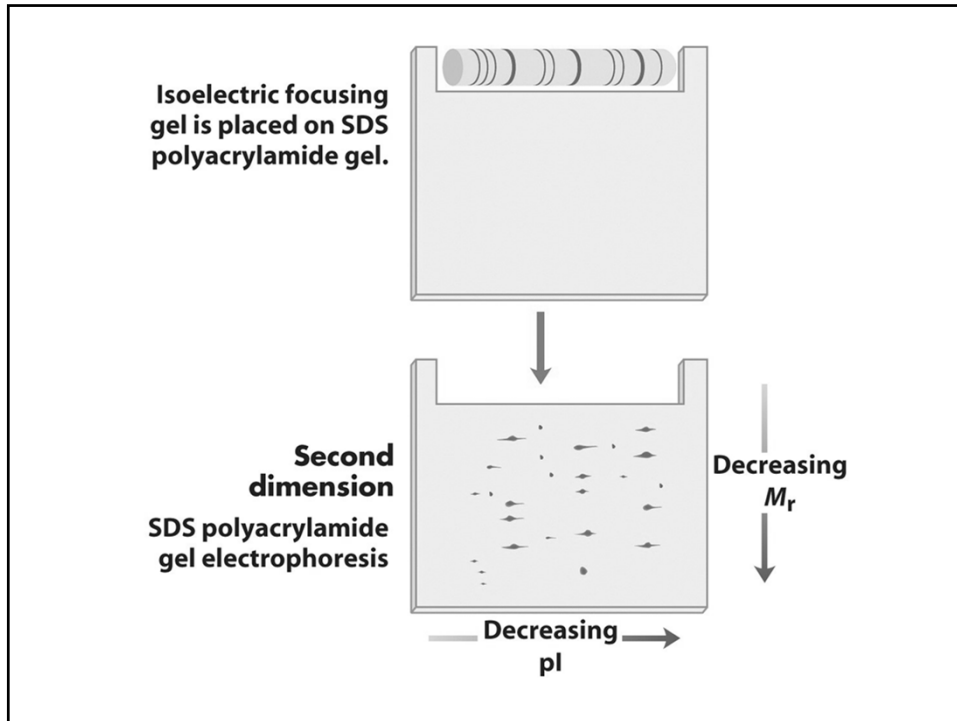
## IEF

- **Separation by charge**
- **pH gradient established by ampholytes**
- **Gel matrix**
  - polyacrylamide strips with immobilized pH gradient.
  - pH gradients in ranges from 3-11 NL to one pH unit (i.e. 6-7) in lengths of 7cm to 28 cm strips
  - Tube gels
- **Run times**
  - Immobilized IEF            6-24 hours
  - Vertical/tube gels IEF    2-6 hours

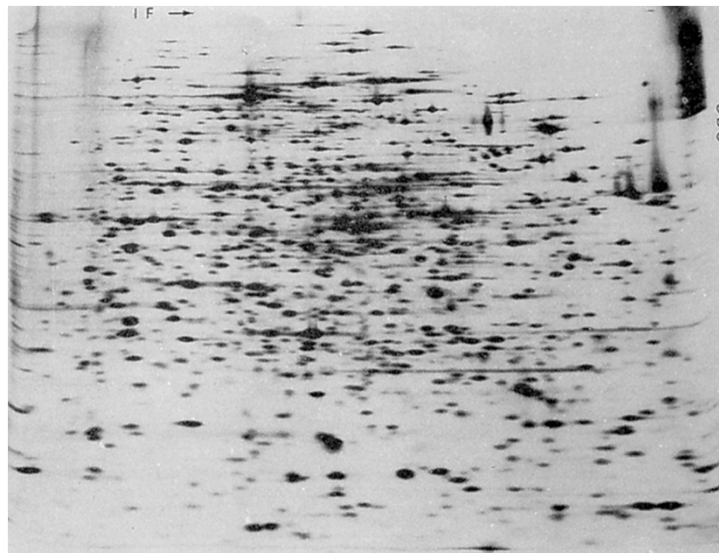


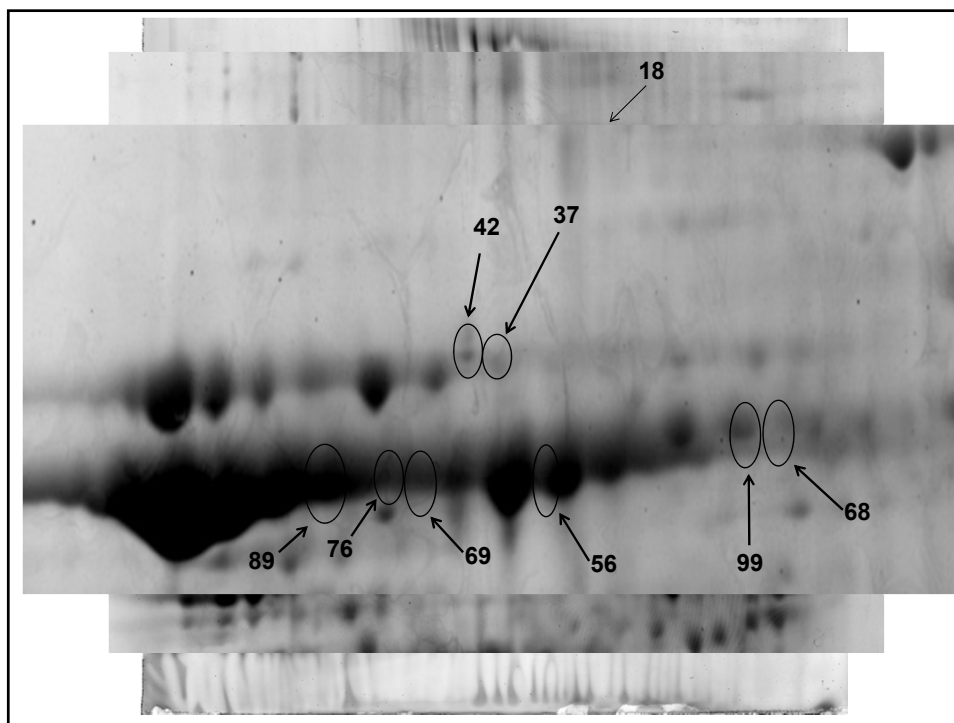
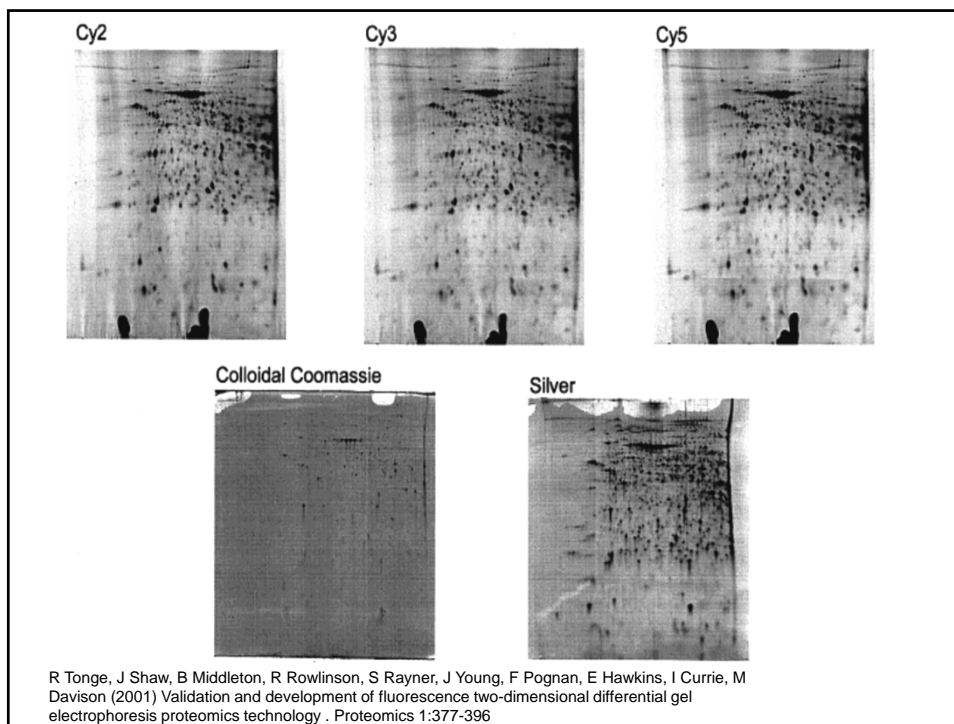
## 2D gel electrophoresis

- Perform IEF
- Place IEF gel in large well of SDS gel and perform electrophoresis
- Stain
- Cut out spots to identify by mass spectrometry



## 2D Gel Electrophoresis





Spot	Identification	Avg. ratio	T-test	% Cov	Accession #
31	Mitochondrial Chaperonin 60 (Zea Mays)	2.68	0.002	51	AAA33452.1
27	Similar to Heat-shock protein precursor	1.88	0.046	20	NP_001066882.1
242	Not analyzed	1.69	0.022	----	-----
18	Victorin Binding Protein, Avena sativa (glycine decarboxylase P subunit)	1.58	0.040	33	AAA63798.1
37	Dihydrolipoamide dehydrogenase family protein (glycine decarboxylase L subunit)	1.38	0.001	14	NP_001042918.1 AK330954
38	Not analyzed	1.35	0.014	----	-----
76	Serine hydroxymethyltransferase	1.31	0.002	25	AAA33687.1
69	Serine hydroxymethyltransferase	1.28	0.008	23	AAA33687.1
89	Serine hydroxymethyltransferase	1.28	0.044	30	AAA33687.1
42	Chloroplast ATP Synthase $\alpha$ -subunit T. aestivum	1.27	0.007	16	AAA84725.1
42	Dihydrolipoamide dehydrogenase family protein				AK330954
68	heat shock protein Hsp90	1.21	0.042	20	Os12g0514500
99	T-cytoplasm male sterility restorer factor 2 (mitochondrial aldehyde dehydrogenase 2)	1.15	0.031	23	AAG43988
56	Rubisco large sub unit	-1.39	0.022	32	ABR01438
56	Serine hydroxymethyltransferase			23	AAA33687.1

## Limitations

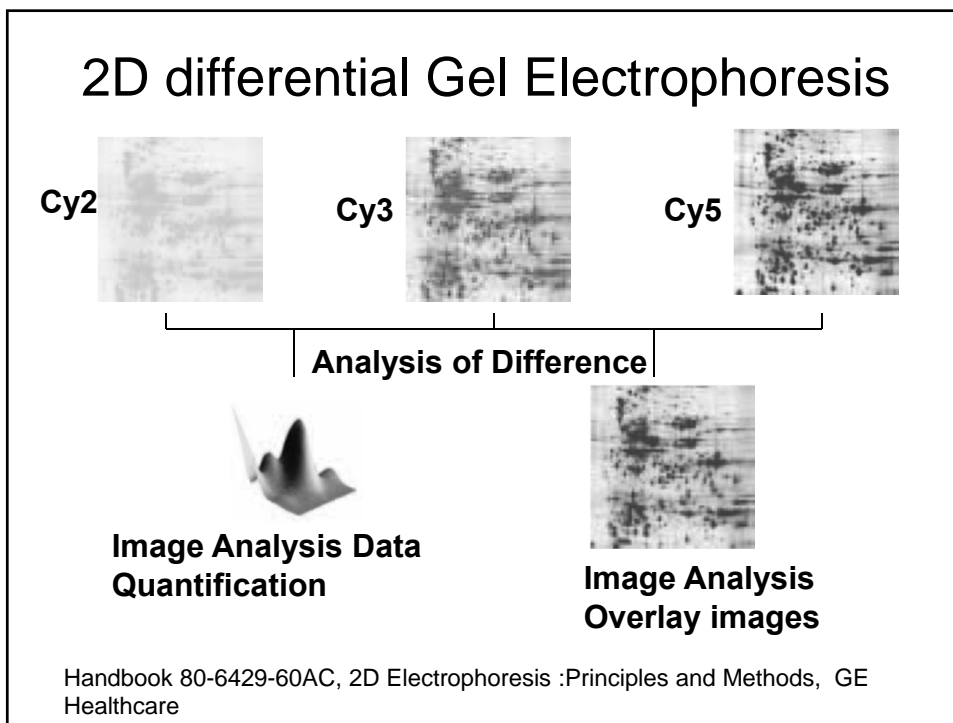
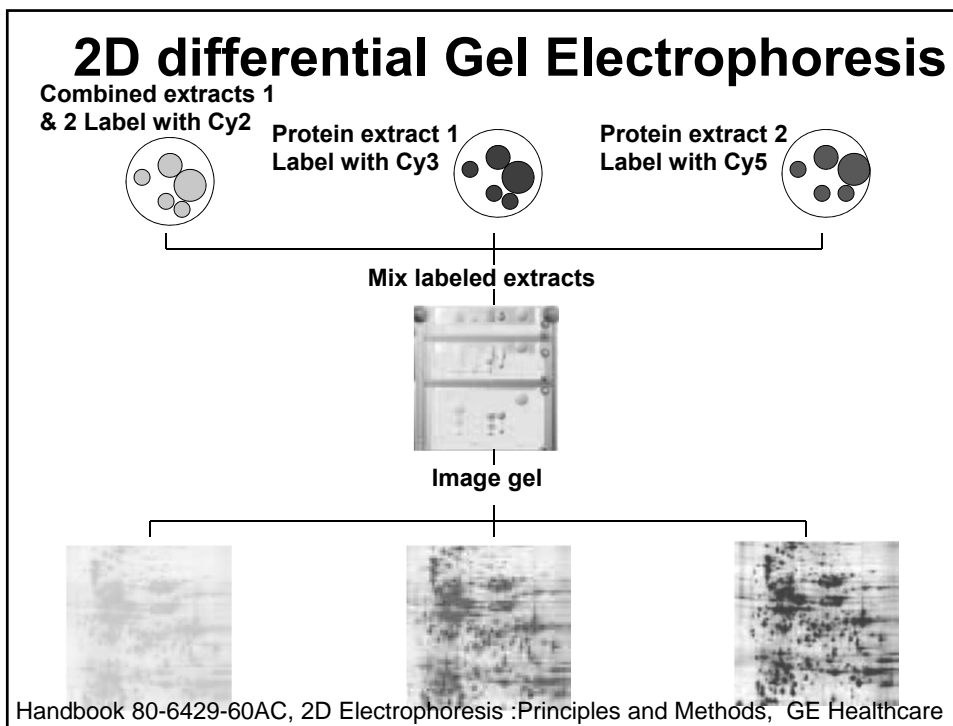
- A single protein can make multiple spots so number of proteins less than spots
- Usually see only most abundant proteins
- Separation limited by gel concentration and size
- Basic and membrane bound proteins are not well separated by 2D gel electrophoresis.

## Other 2D Gel Methods

- **Blue Native Gel followed by SDS gel**
  - Used for organelles such as mitochondria and chloroplasts
  - Keeps electron transport complexes together during native gel process
- **Non denaturing followed by denaturing**
  - Can allow for complexes to move together
  - Then separates subunits of complexes
- **Differential gel electrophoresis**

## Differential Gel Electrophoresis

- **Allows measurement of the relative concentration of proteins**
- **Method**
  - Isolate proteins from test and control
  - Label test proteins with one dye
  - Label control protein with second dye
  - Make third sample of mixed control and test and label with third dye.
  - Combine all three samples and separate by 2D gel electrophoresis
  - Analyze the intensity of the test and sample relative to the combined sample.

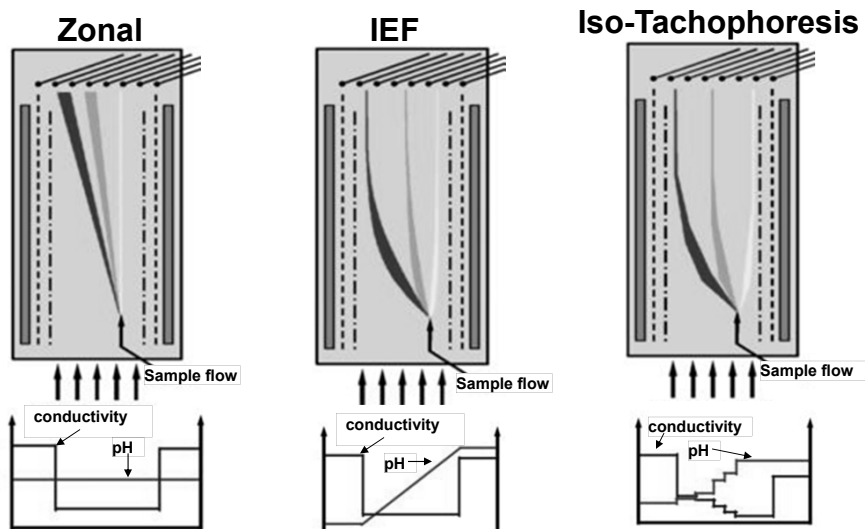




## Free Flow Electrophoresis



## Forms of Free-Flow Electrophoresis



<http://www.bd.com/proteomics/products/ffe/technology.asp>

## FFE

- **FFE can be used to separate any charged item that can be suspended in an aqueous solution.**
  - Cells (zonal, isotacho-)
  - Organelle (zonal, isotacho-)
  - Proteins (IEF)
  - Subcellular fragments (zonal, isotacho-)
  - Nanoparticles
- **Uses low molecular weight weak acids and bases to establish pH.**

## 2 Dimensional Chromatography

- Alternative means to reduced protein complexity.
- Consists of performing two or more usually orthogonal chromatographic steps prior to LC-MS/MS
- Process sometimes called Multidimensional protein identification technology (MuDPIT)

## 2D Chromatography

### Types of chromatography

Strong cation/anion exchange

SCX/SAX

Weak cation/anion exchange

WCX/WAX

Size exclusion chromatography

SEC

Hydroxyapatite chromatography

HA

Chromatofocusing

Hydrophobic interaction

HIC

Reverse Phase

RP

Mixed bed

## 2D Chromatography

- Advantages
  - Reduces complexity for LC-MS/MS and 2D gels
  - Can concentrate low abundance proteins
- Disadvantages
  - Typically up to 20% loss at each chromatographic step
  - Longer experiment times

# Gel Electrophoresis

## What you need to know

- Types of gel electrophoresis
  - Most common -- SDS-PAGE, IEF, 2D
  - Other methods (FFE, blue native, differential, etc.)
  - How differential gel electrophoresis works.
  - How each method separates proteins
  - Limitations
- 2 dimensional chromatography
  - How each method separates proteins
  - Limitations