

Biological Chemistry Laboratory  
Biology 3515/Chemistry 3515  
Spring 2017

Lecture 21:

SDS Gel Electrophoresis

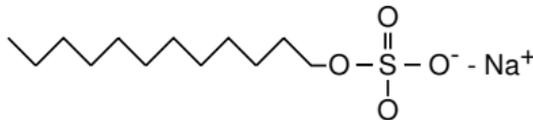
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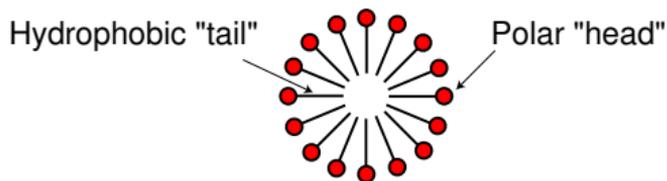
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# SDS - Sodium Dodecyl Sulfate



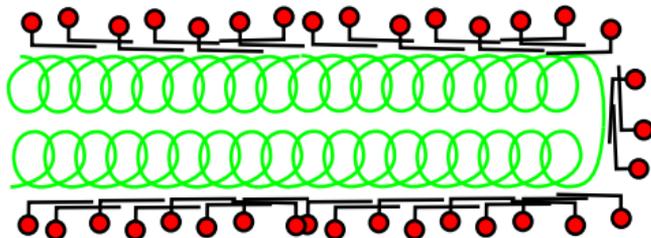
- Also called lauryl sulfate
- A common ingredient of shampoos
- Forms micelles in water



Micelles are three-dimensional, *i.e.*, roughly spherical.

# SDS Denatures Proteins and Binds to Them

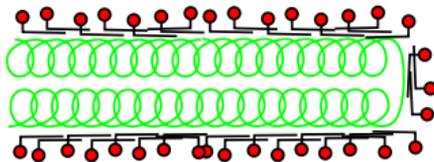
- Most proteins bind SDS at a constant ratio:  
≈ 1.4 g SDS per g protein.
- Complexes are rod shaped.
- Polypeptides form  $\alpha$ -helical structures in SDS.
- A possible structure of SDS-protein complexes:



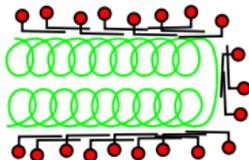
## Clicker Question from the Last Lecture

Which will have the higher electrophoretic mobility, in the absence of a gel?

1 A large protein with SDS bound:

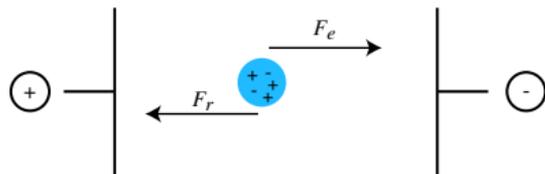


2 A small protein with SDS bound:



3 They will have the same mobility.

# Electrophoresis in the Absence of a Gel



- Electromotive force:  $F_e = z \cdot e \cdot E$

$z$  = average charge

$e$  = unit of electric charge, a constant

$E$  = electric field strength, proportional to voltage

- Resistive force:  $F_r = v \cdot f$

$v$  = velocity

$f$  = frictional coefficient, a molecular property

- Molecule accelerates until  $F_r = F_e$

## Velocity when $F_r = F_e$

- Electrophoretic and frictional forces:

$$F_e = z \cdot e \cdot E$$

$$F_r = v \cdot f$$

- When the forces are equal:

$$v \cdot f = z \cdot e \cdot E$$

$$v = \frac{z \cdot e \cdot E}{f}$$

- Define free mobility (mobility in absence of gel) as the velocity normalized by electric field:

$$M_0 = \frac{v}{E} = \frac{z \cdot e}{f}$$

- $M_0$  should be independent of applied voltage.

# Frictional Coefficient and Net Charge for SDS-Protein Complexes

## ■ Frictional coefficient

- Frictional coefficient for rod-shaped molecule:

$$f \propto \text{Rod Length}$$

$$f \propto \text{Molecular Weight}$$

$$f = C_f \cdot MW$$

- $C_f$  is a constant for SDS-protein complexes.

## ■ Net charge

- Charges from SDS generally overwhelm intrinsic charge of polypeptides.
- Since proteins bind a constant amount of SDS per g:

$$z \propto \text{Molecular Weight}$$

$$z = C_z \cdot MW$$

- $C_z$  is a constant for SDS-protein complexes.

# Free Electrophoretic Mobilities of SDS-Protein Complexes

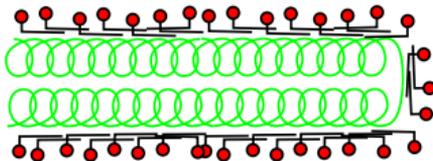
$$M_0 = \frac{z \cdot e}{f} = \frac{C_z \cdot MW \cdot e}{C_f \cdot MW}$$

$$M_0 = \frac{C_z \cdot e}{C_f}$$

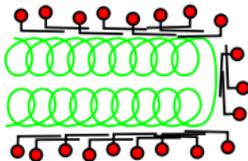
## Clicker Question #1

Which will have the higher electrophoretic mobility, in the absence of a gel?

1 A large protein with SDS bound:



2 A small protein with SDS bound:

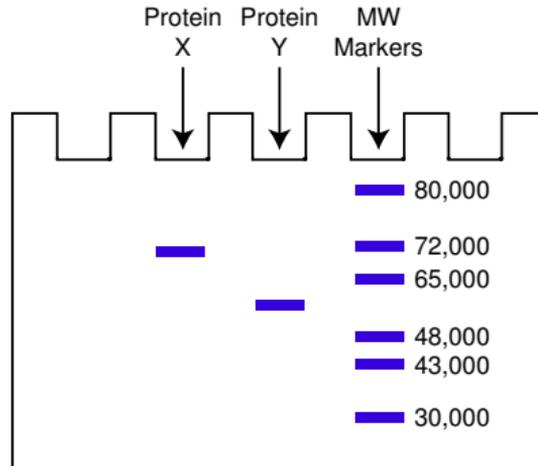


3 They will have the same mobility.

■ All SDS-protein complexes should have the same free mobility, irrespective of molecular weight!

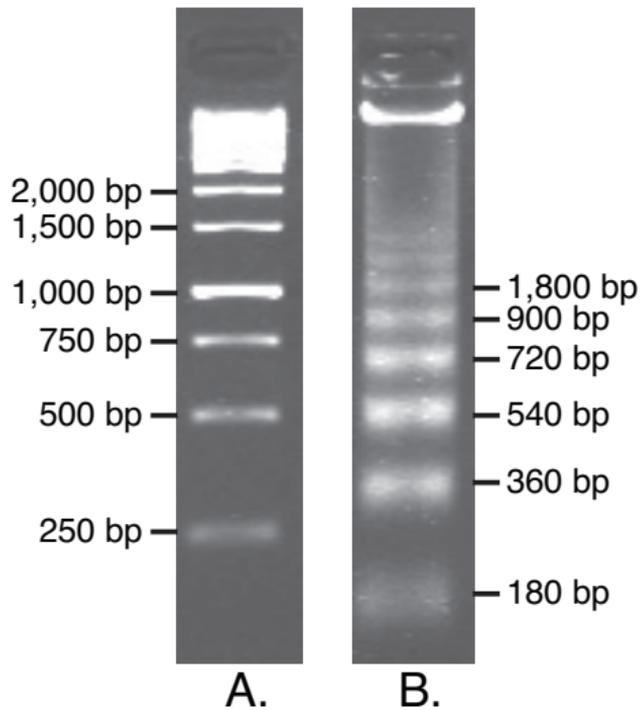
# Separation of Proteins on SDS Gels Depends Almost Exclusively on Sieving Effect of the Gel.

- Sieving effect depends only on size (since protein/SDS complexes have a common shape).
- SDS gels can be used to determine molecular weights of polypeptides



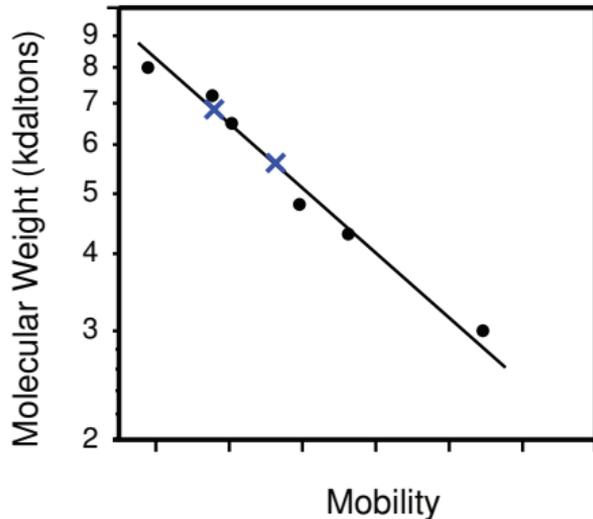
- What other class of molecules is expected to behave this way in electrophoresis?

# Gel Electrophoresis of DNA Fragments



- Electrophoresis through agarose gel.
- DNA stained by binding a fluorescent dye.
- A. Artificial DNA fragments.
- B. DNA fragments generated during cell death (apoptosis).

# Calibration Curve for SDS Gel Electrophoresis



- Measure mobilities of proteins with known molecular weights.
- Fit a line (or curve) to data for standards.
- Estimate molecular weights of other proteins from mobilities and empirical calibration curve.

# Cautions Regarding Determination of Molecular Weights by SDS Gel Electrophoresis

## 1 Molecular weights are for individual polypeptides.

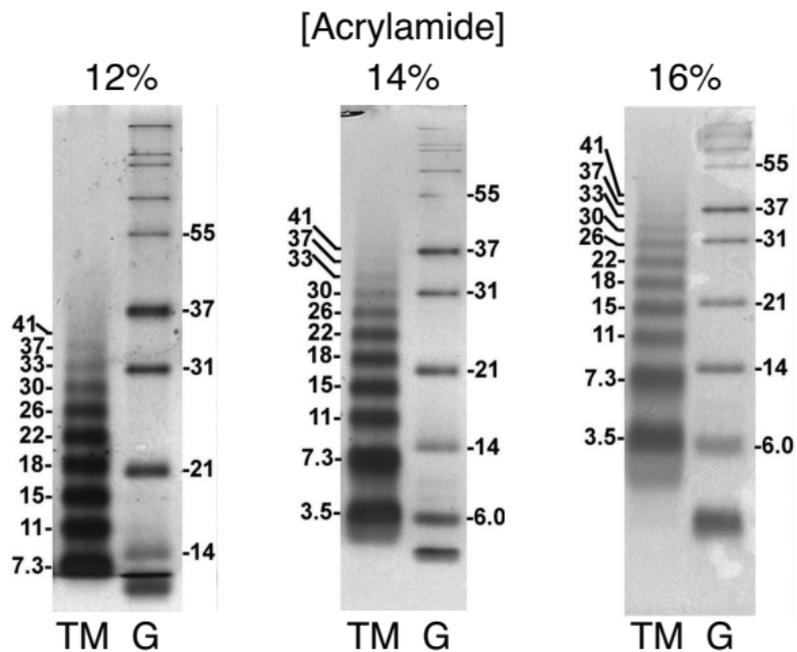
- Oligomeric proteins or other complexes are dissociated by SDS, and information about their structures or sizes is lost.

## 2 Deviations from expected mobility are common.

Possible explanations:

- Differences in amount of SDS bound to protein.
- Residual structure of polypeptide after binding SDS.
- Intrinsic charge of protein is not insignificant.

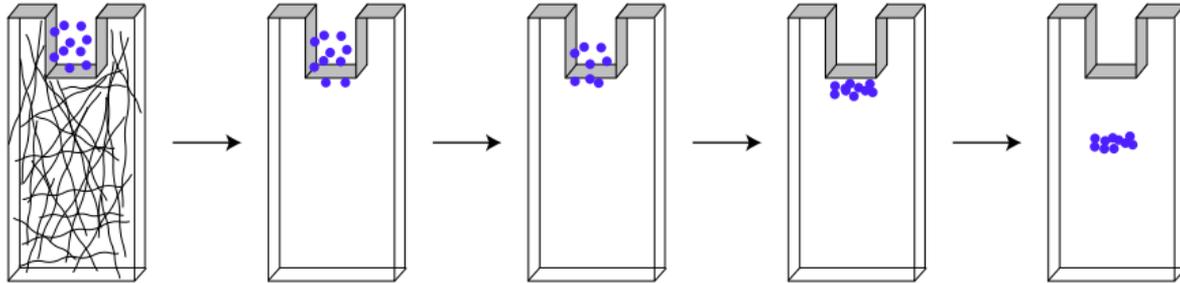
# Effects of Acrylamide Concentration on the Mobilities of Globular and Transmembrane Proteins in SDS Gels



- TM: Artificial mimics of transmembrane proteins.
- G: “Normal” globular proteins.
- Numbers indicate molecular weights, in kdalton.
- Mobilities decrease as acrylamide concentration is increased.
- Separation of smaller proteins increases as acrylamide concentration is increased.
- Separation of larger proteins is better at low acrylamide concentrations.
- Relative mobilities of TM and globular proteins change with acrylamide concentration.

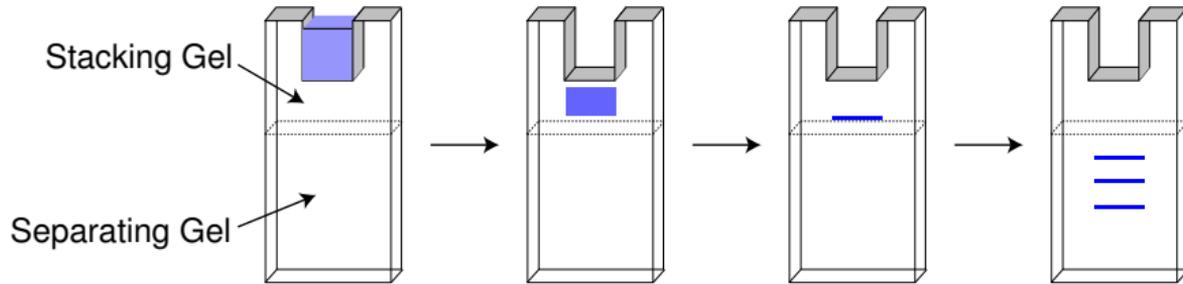
## Resolution and Sensitivity Depend on Narrow Protein Bands

- Natural “stacking” of protein molecules as they enter a gel:



- Final protein bands are often much narrower than the original sample applied in the well.

# Stacking and Separation in a Discontinuous Gel



How does it work?