

***Separation tools for
electrospray mass spectrometry
- Liquid Chromatography***

Introduction to Chromatography

Definition

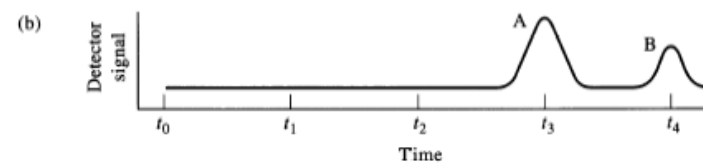
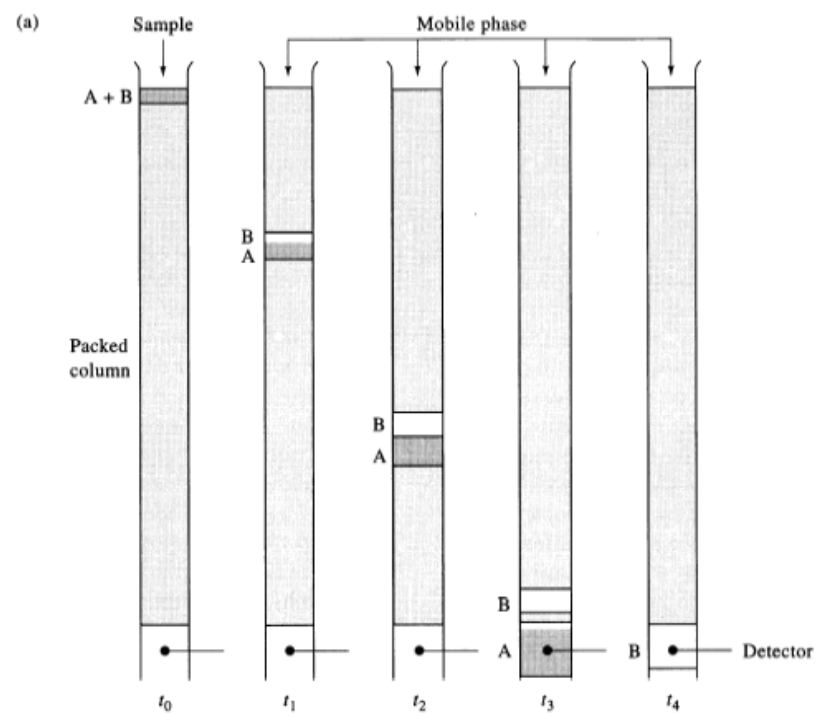
Chromatography is a separation technique based on the different interactions of compounds with two phases, a *mobile phase* and a *stationary phase*, as the compounds travel through a supporting medium.

Components:

mobile phase: a solvent that flows through the supporting medium

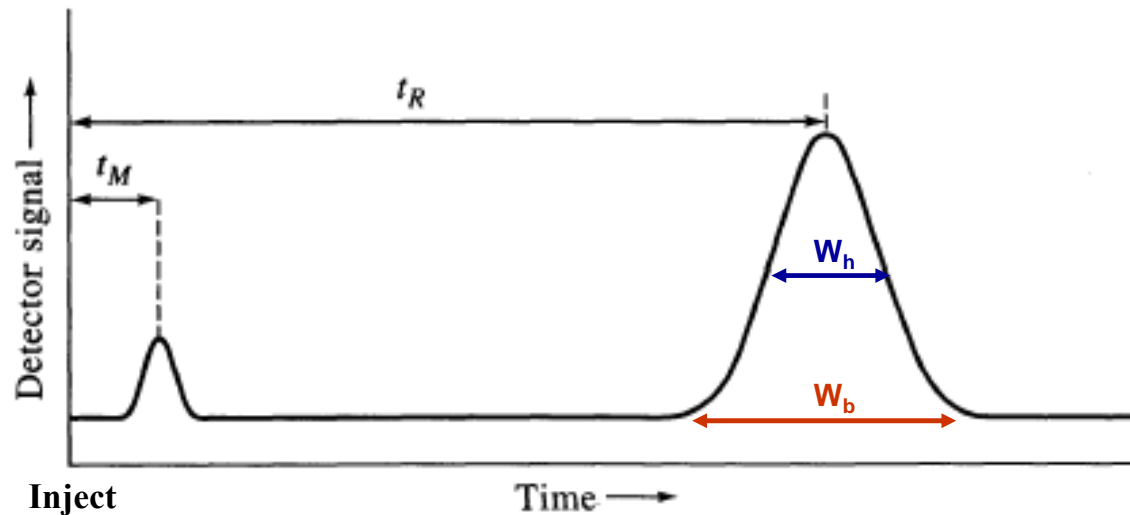
stationary phase: a layer or coating on the supporting medium that interacts with the analytes

supporting medium: a solid surface on which the stationary phase is bound or coated



Theory of Chromatography

I. Typical response obtained by chromatography (i.e., a chromatogram): chromatogram - intensity vs. elution time



Where:

t_R = retention time

t_M = void time

W_b = baseline width of the peak in time units

W_h = half-height width of the peak in time units

II. Solute Retention:

Capacity factor (k'): more universal measure of retention, determined from t_R or V_R .

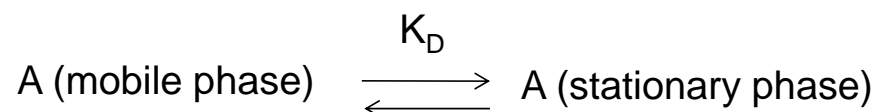
$$k' = (t_R - t_M) / t_M \quad \text{or}$$

$$k' = (V_R - V_M) / V_M$$

$$k' = \frac{\text{moles } A_{\text{stationary phase}}}{\text{moles } A_{\text{mobile phase}}}$$

k' is directly related to the strength of the interaction between a solute with the stationary and mobile phases.

A simple example relating k' to the interactions of a solute in a column is illustrated for partition chromatography:



where: K_D = equilibrium constant for the distribution of A between the mobile phase and stationary phase

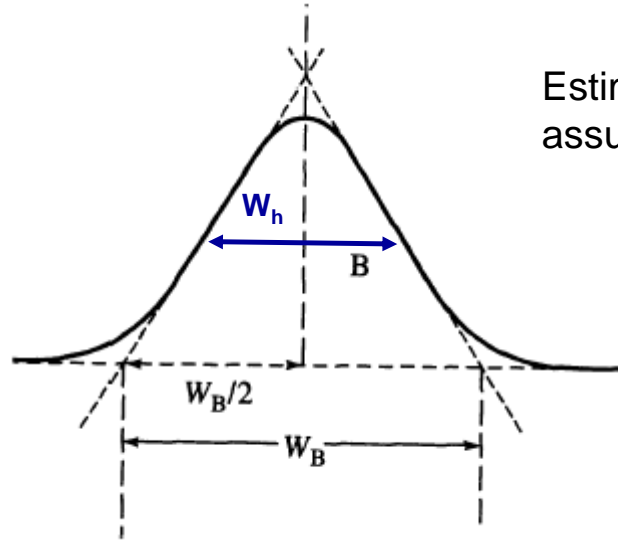
Assuming local equilibrium at the center of the chromatographic peak:

$$k' = \frac{[A]_{\text{stationary phase}} \text{Volume}_{\text{stationary phase}}}{[A]_{\text{mobile phase}} \text{Volume}_{\text{mobile phase}}}$$

$$k' = K_D \frac{\text{Volume}_{\text{stationary phase}}}{\text{Volume}_{\text{mobile phase}}}$$

As K_D increases, interaction of the solute with the stationary phase becomes more favorable and the solute's retention (k') increases

III. Efficiency: is related experimentally to a solute's peak width.



Estimate σ from peak widths, assuming Gaussian shaped peak:

$$W_b = 4\sigma$$

$$W_h = 2.354\sigma$$

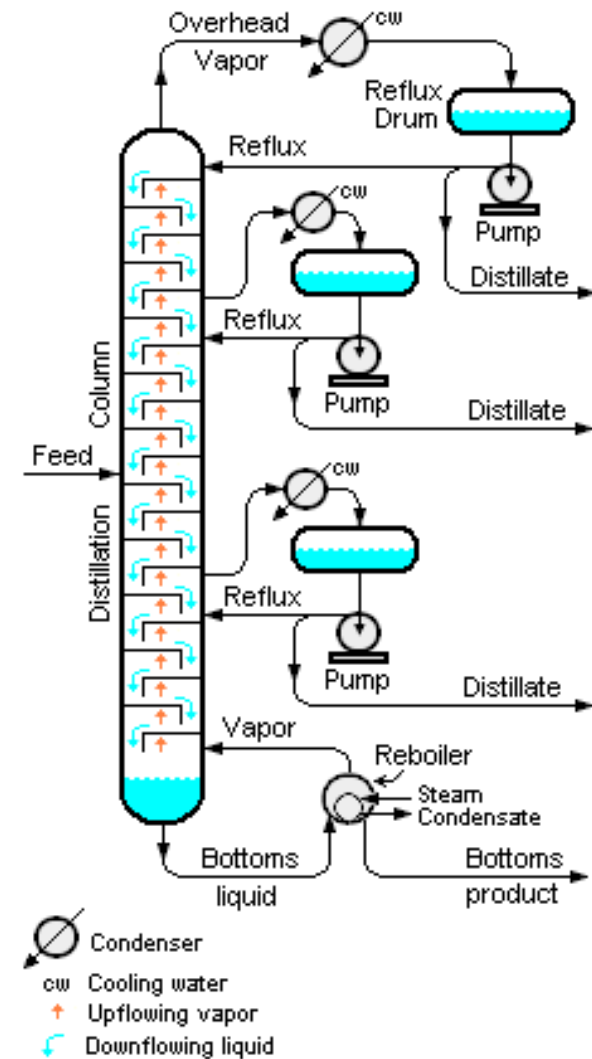
Number of theoretical plates (N)

$$N = (t_R / \sigma)^2$$

or for a Gaussian shaped peak

$$N = 16 (t_R / W_b)^2$$

$$N = 5.54 (t_R / W_h)^2$$



The larger the value of N is for a column, the better the column

- the better the ability to resolve solutes that have small differences in retention
- N is independent of solute retention
- N is dependent on the **length** of the column

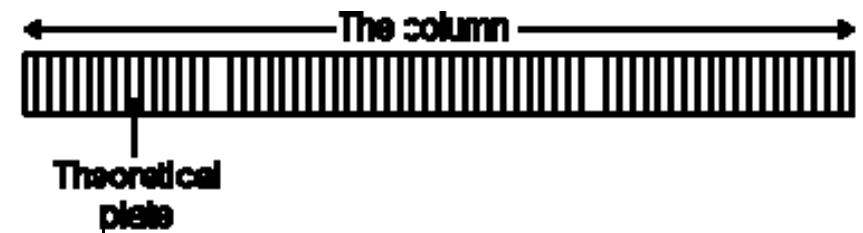


Plate height or height equivalent of a theoretical plate (H or HETP)

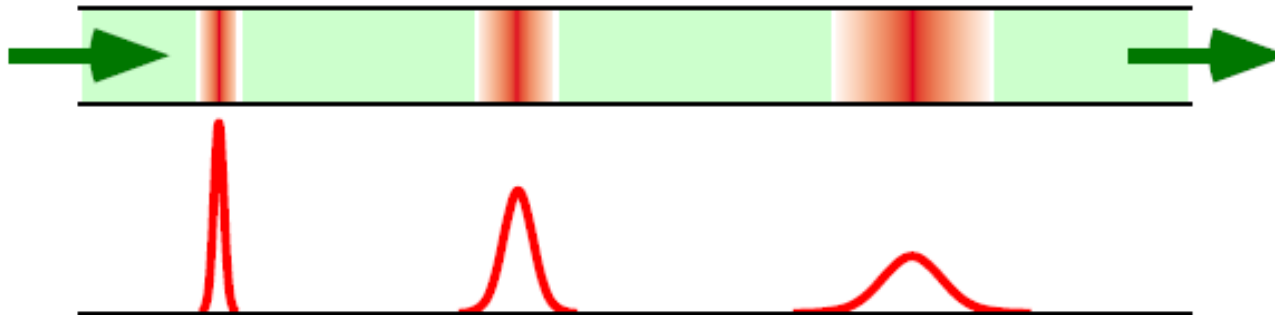
$$H = L / N$$

where: L = column length, N = number of theoretical plates for the column

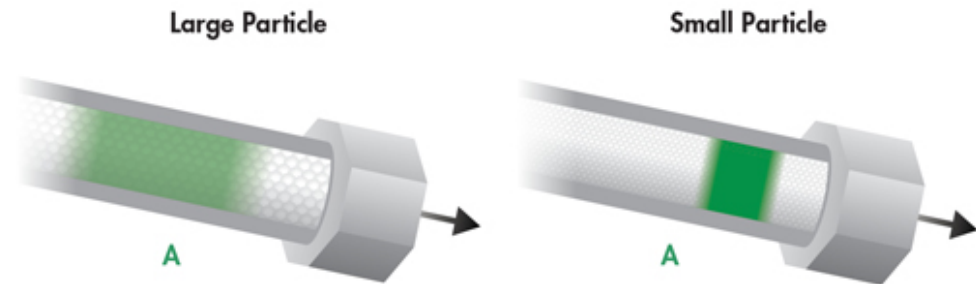
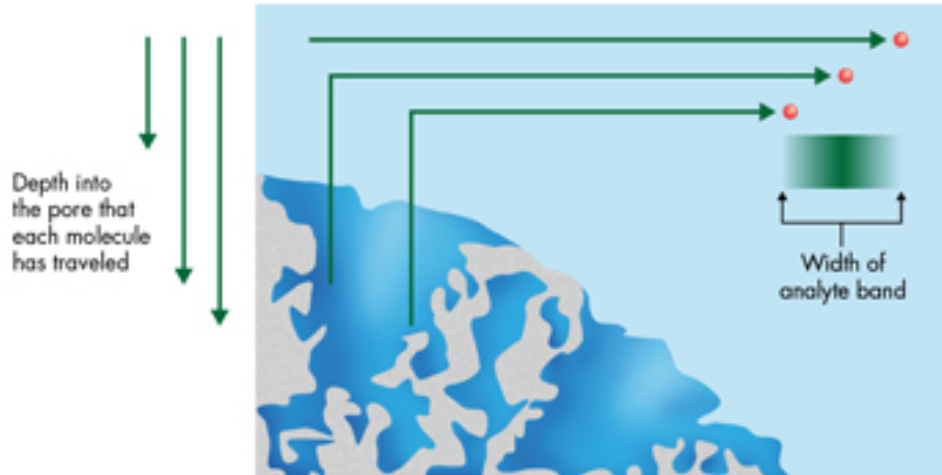
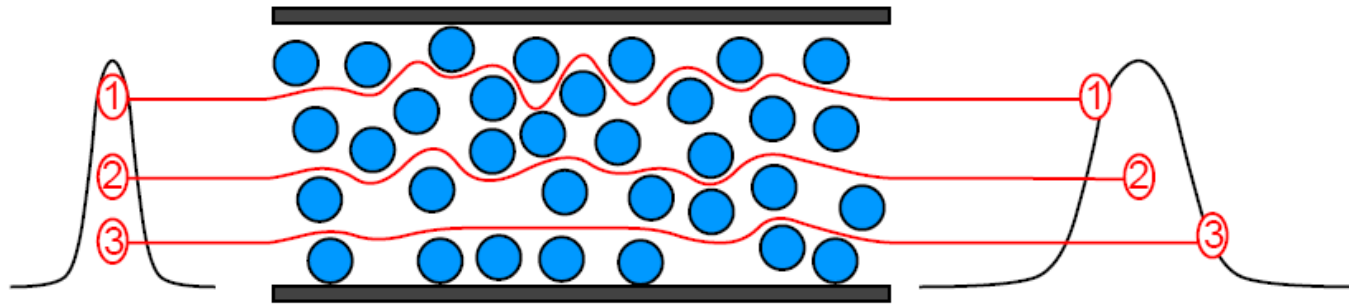
H can be also used to relate various chromatographic parameters (e.g., flow rate, particle size, etc.) to the kinetic processes that give rise to peak broadening

Why Do Bands Spread?

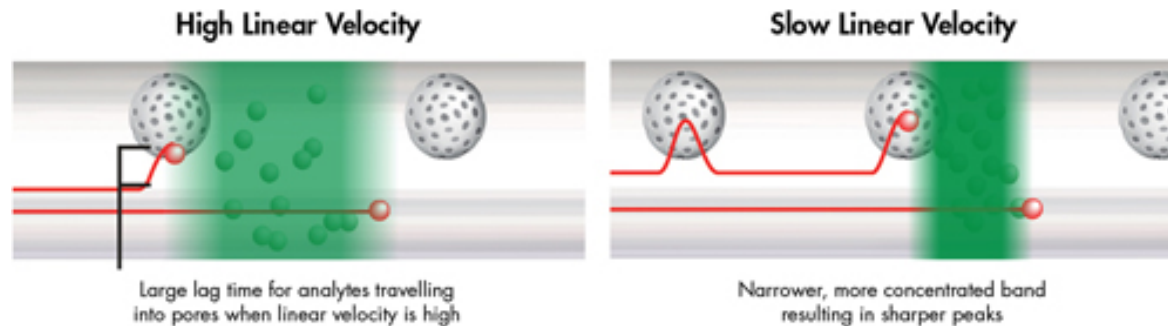
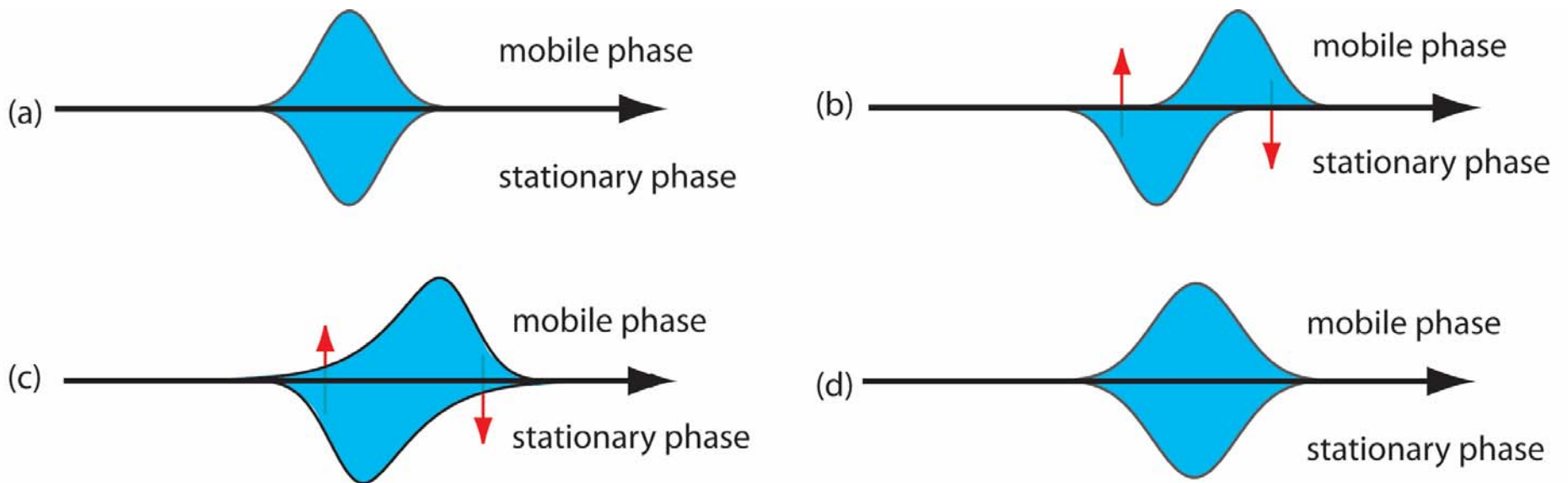
- a. Eddy diffusion (multiple Paths)
- b. Stationary phase mass transfer
- c. Longitudinal diffusion



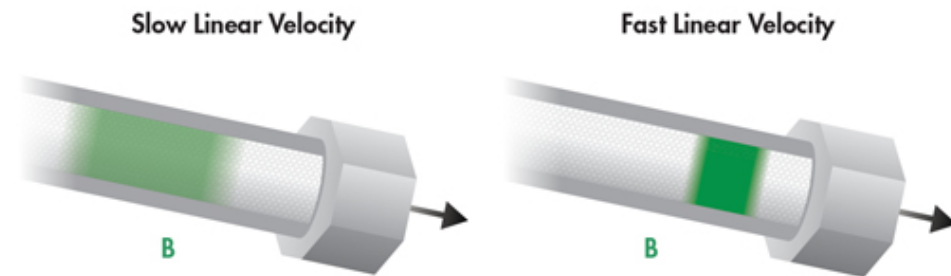
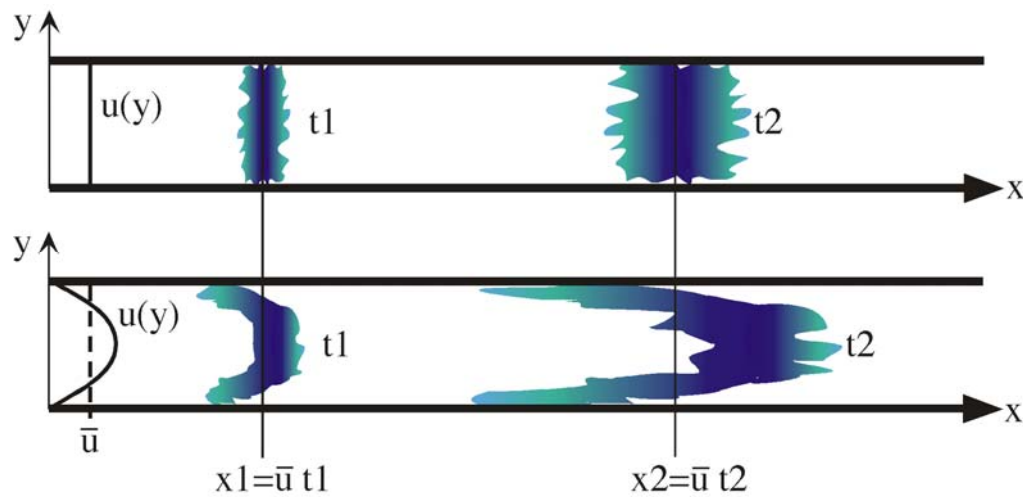
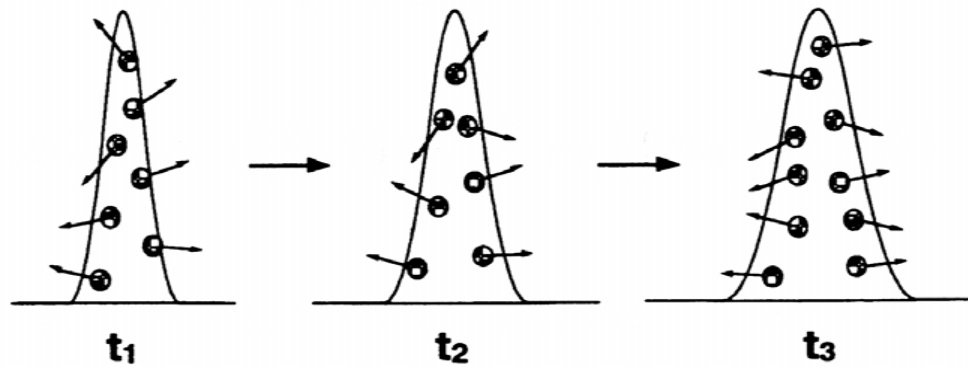
a) **Eddy diffusion** – multiple flow paths through a packed column.



b) Stationary phase mass transfer (equilibrium factor)

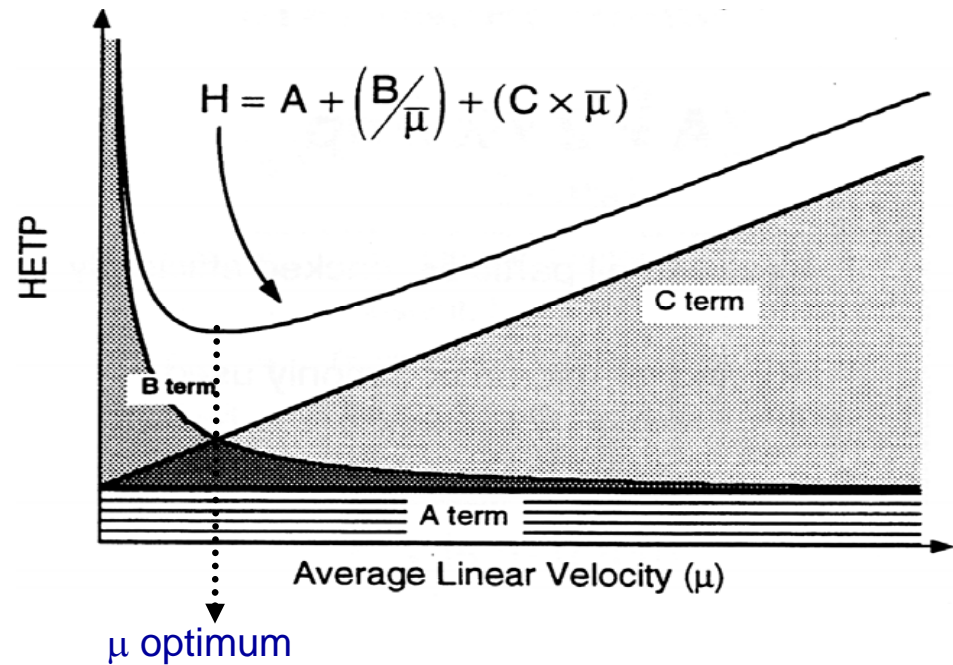


c) Longitudinal diffusion



Van Deemter equation:

$$H = A + \frac{B}{u} + Cu$$



Number of theoretical plates (N) = $5.54 (t_R / W_h)^2 = L / H$

Smaller $H \rightarrow$ better separation

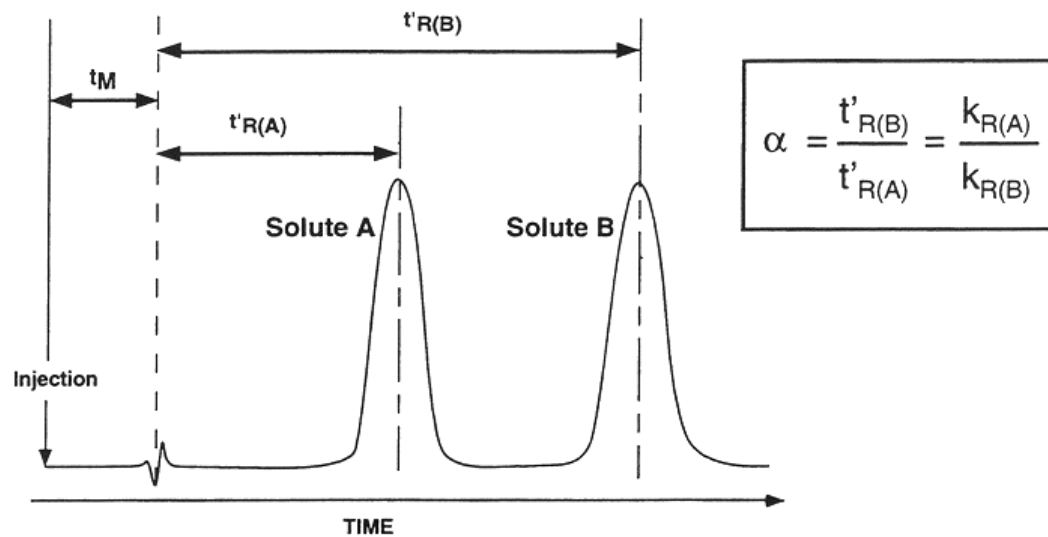
IV. Measures of Solute Separation:

Separation factor (α) = k'_2/k'_1 $k' = (t_R - t_M)/t_M$

where: k'_1 = the capacity factor of the first solute

k'_2 = the capacity factor of the second solute, with $k'_2 > k'_1$

A value of $\alpha > 1.1$ is usually indicative of a good separation

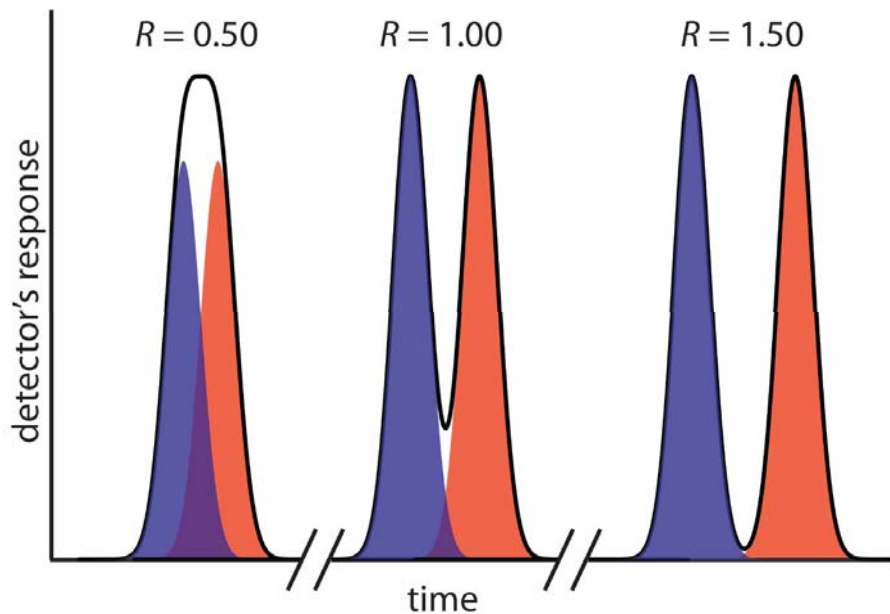


Does not consider the effect of column efficiency or peak widths, only retention.

Resolution (R_s) – resolution between two peaks is how well two peaks are separated:

$$R = \frac{t_{R2} - t_{R1}}{\frac{1}{2}(W_1 + W_2)} \quad \alpha = \frac{k_2}{k_1} = \frac{t_{R2} - t_0}{t_{R1} - t_0} \quad \Rightarrow \quad R = \frac{1}{4} \sqrt{N} \left(\frac{\alpha - 1}{\alpha} \right) \left(\frac{k}{1 + k} \right)$$

$$R \propto \sqrt{N} \propto \sqrt{L}, \quad (N = \frac{L}{H})$$

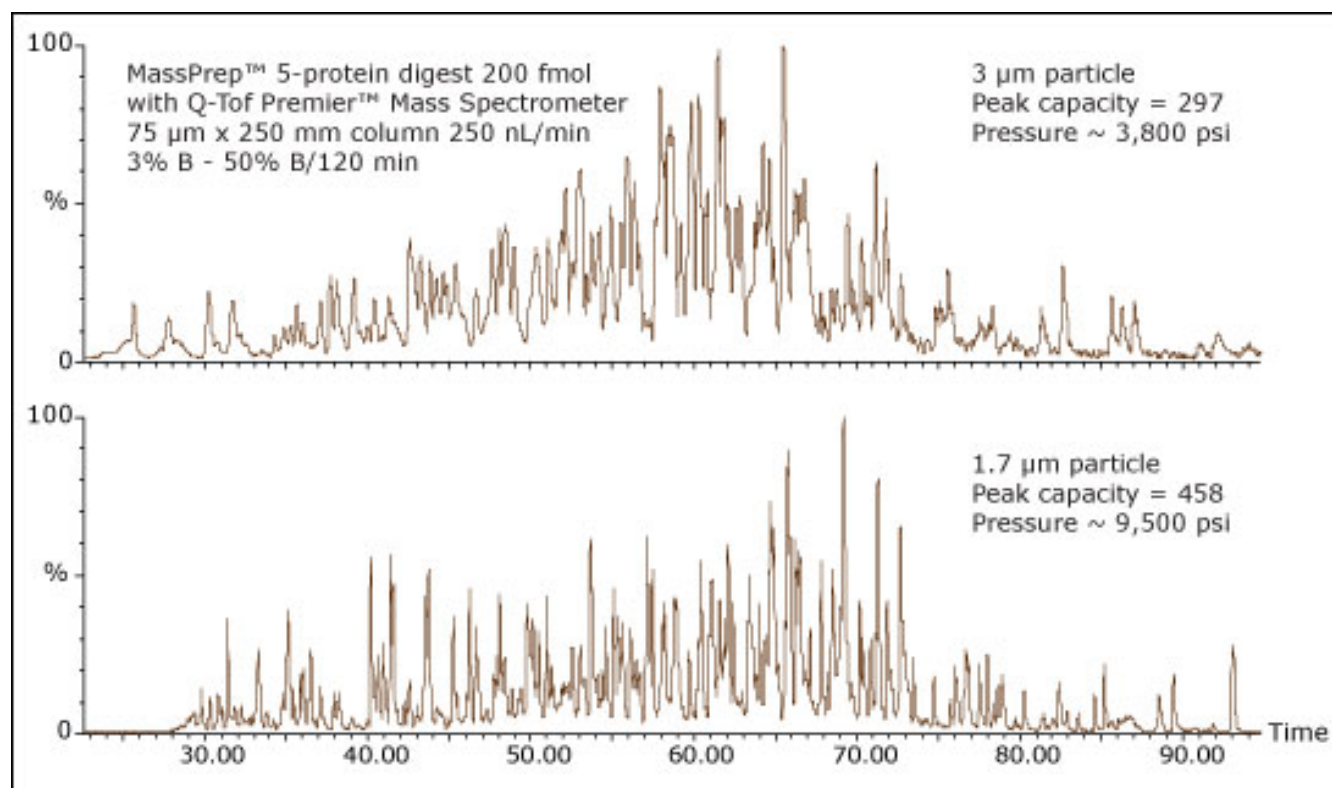


$R_s > 1.5$ represents *baseline resolution*, or complete separation of two neighboring solutes → ideal case.

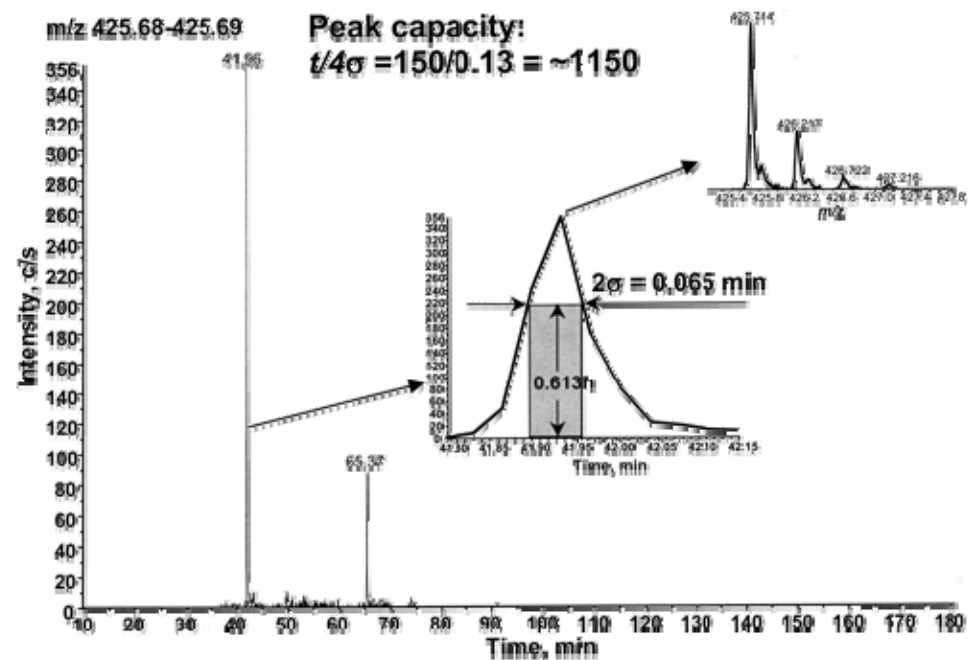
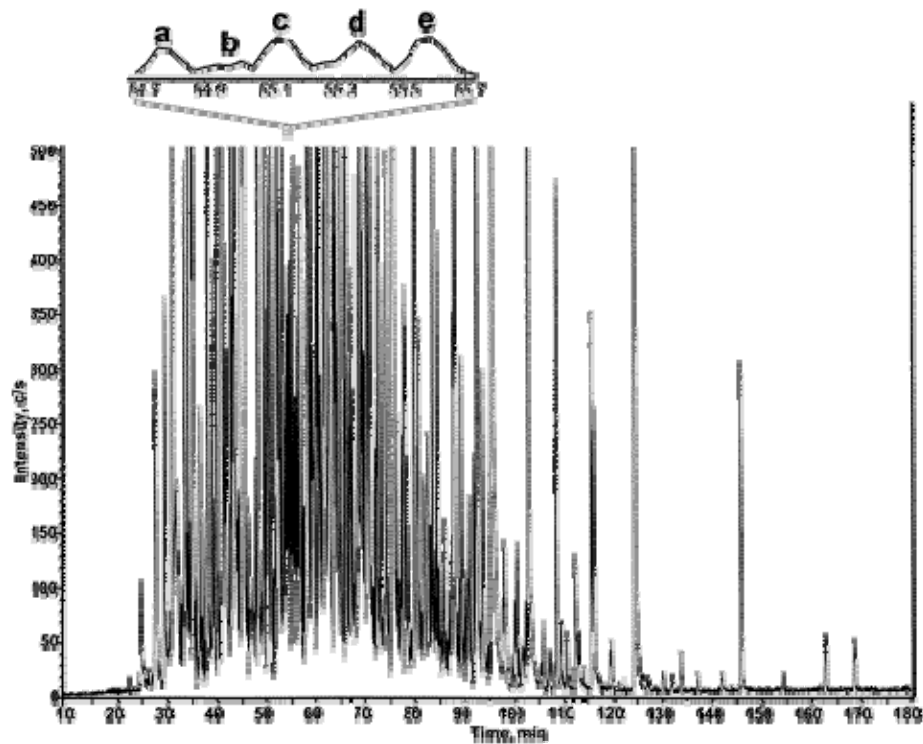
$R_s > 1.0$ considered adequate for most separations.

IV. Peak capacity

Defined as **number of peaks** that can be separated **within a retention window**; thus peak capacity can be simply the gradient run time divided by the average peak width



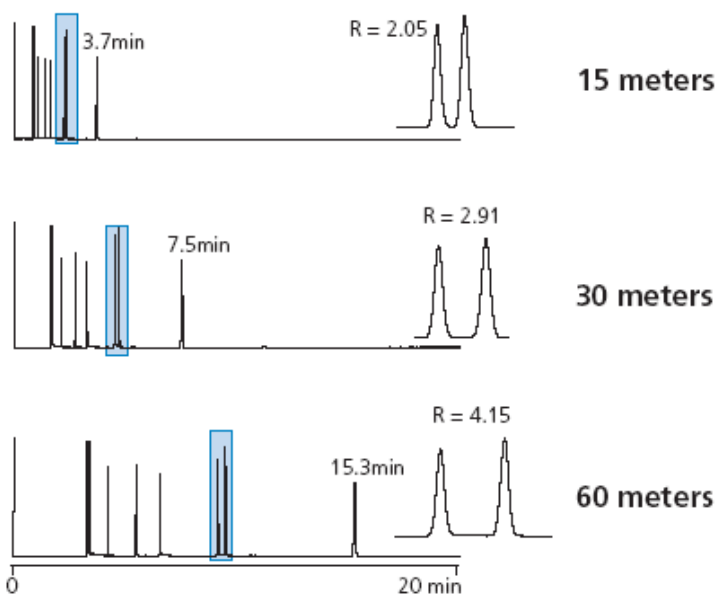
IV. Peak capacity



Summary: how to improve LC separation power ?

$$R \propto \sqrt{N} \propto \sqrt{L}, \quad (N = \frac{L}{H})$$

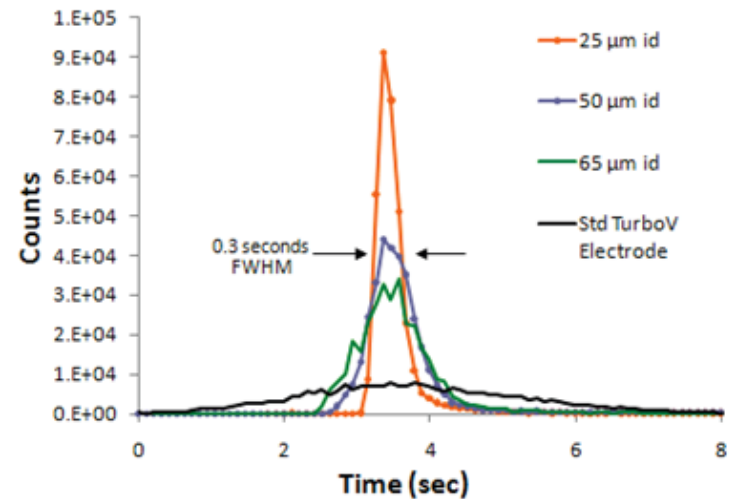
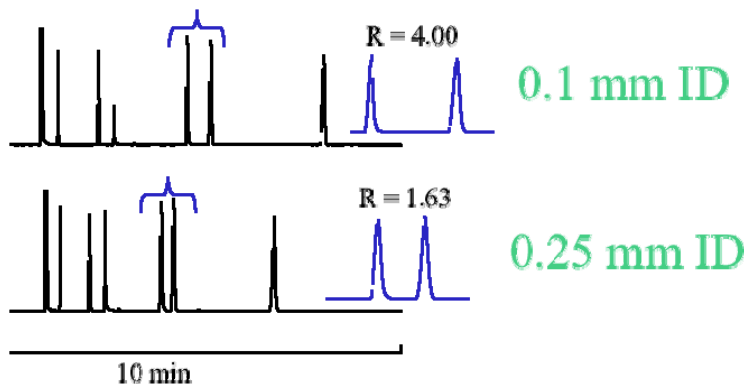
1. Increase column length

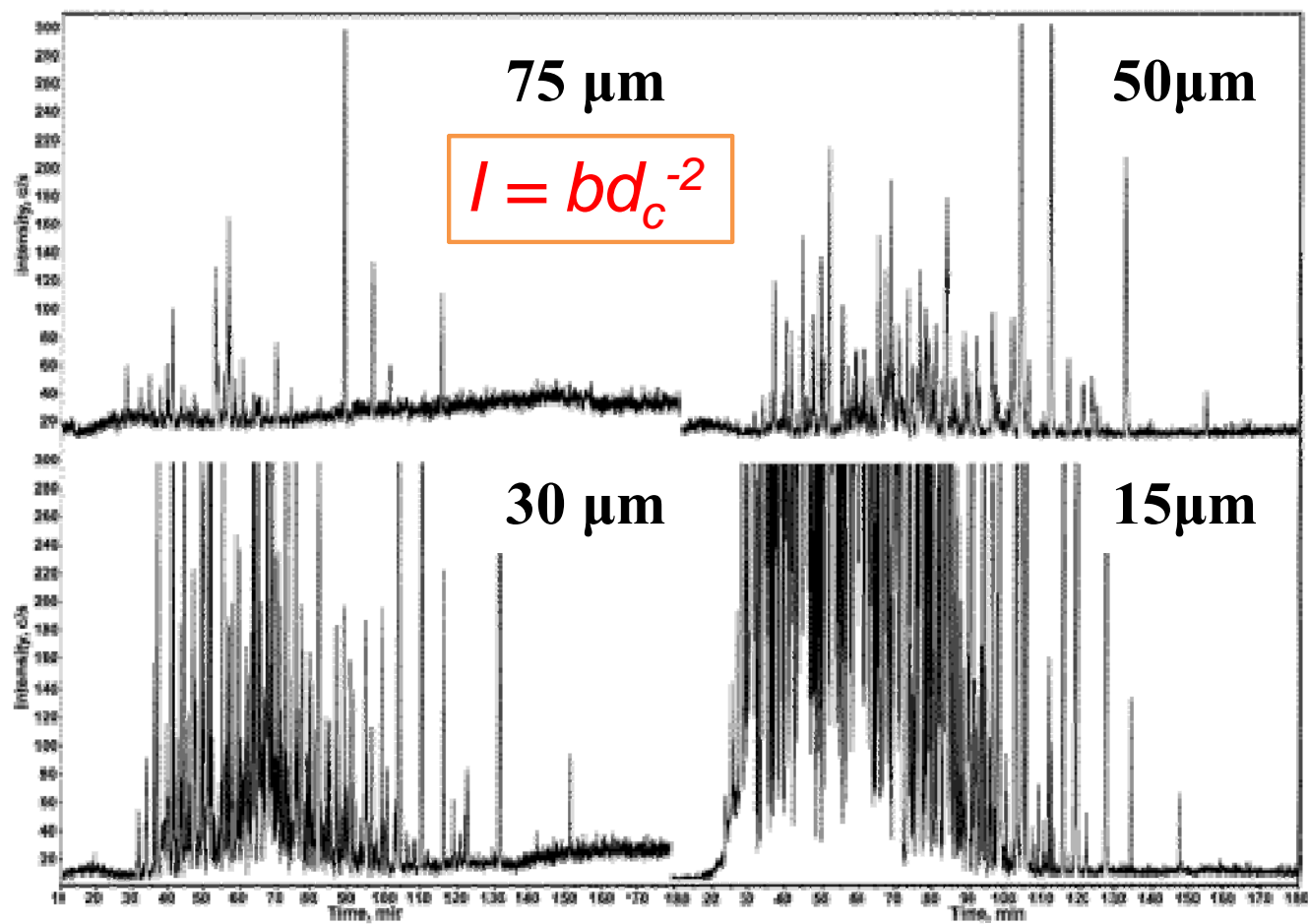


Summary: how to improve LC separation power ?

$$R \propto \sqrt{N} \propto \sqrt{L}, \quad (N = \frac{L}{H})$$

2. Decrease column diameter





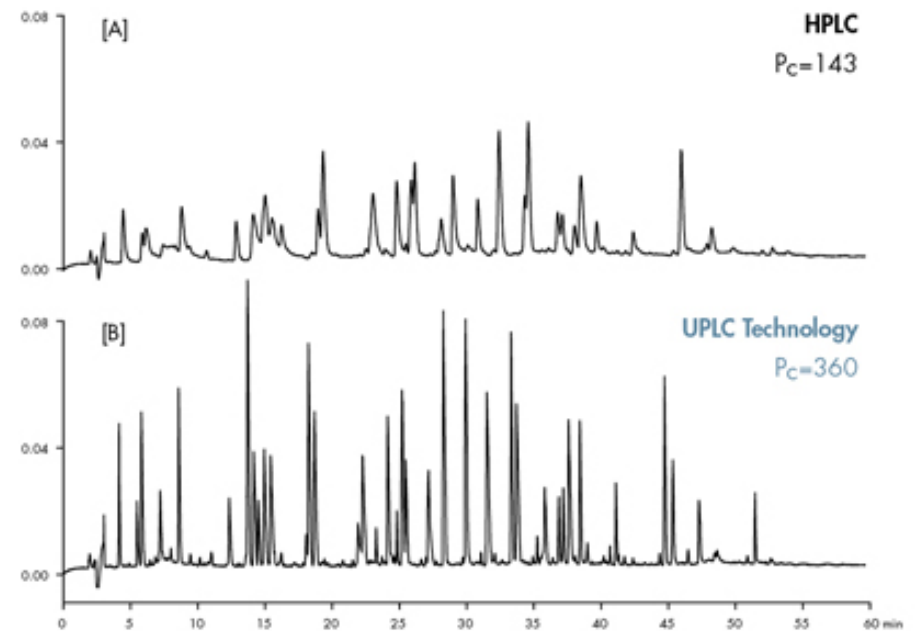
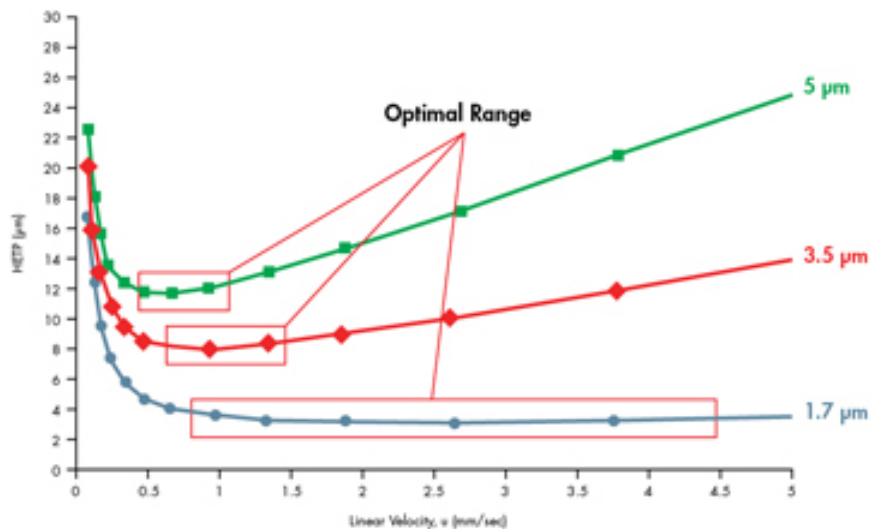
100 ng of a yeast soluble protein tryptic digest

Summary: how to improve LC separation power ?

$$R \propto \sqrt{N} \propto \sqrt{L}, \quad (N = \frac{L}{H})$$

$$H = A + \frac{B}{u} + Cu$$

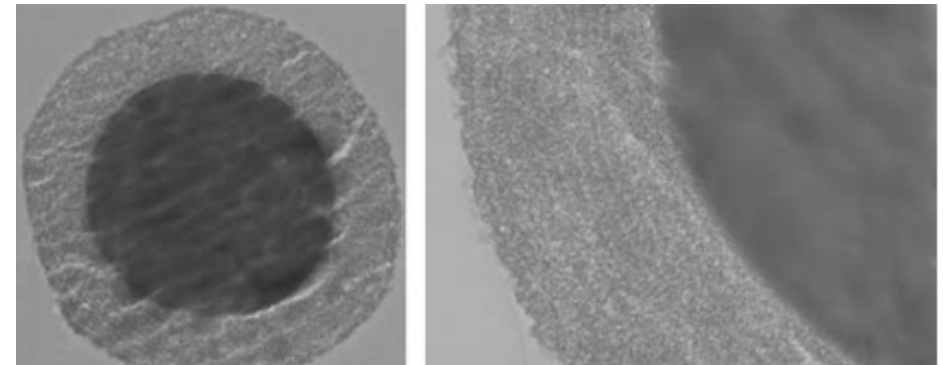
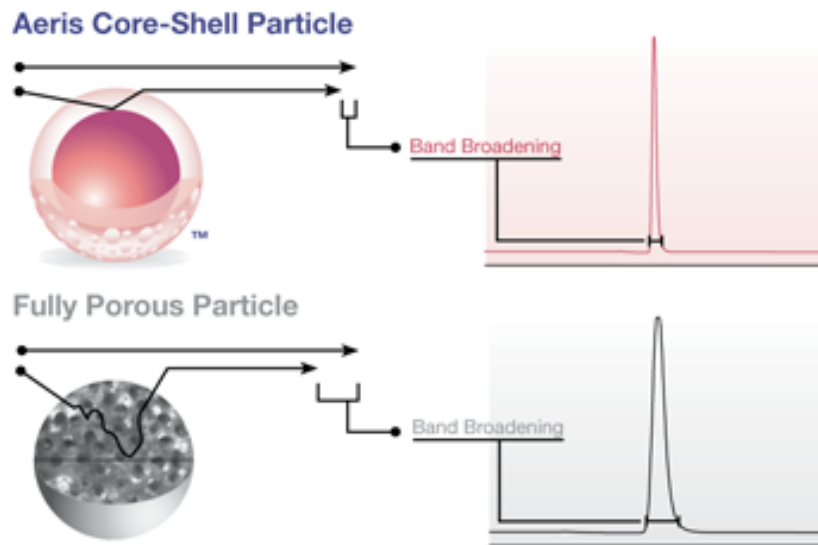
3. Smaller particle



Summary: how to improve LC separation power ?

$$R \propto \sqrt{N} \propto \sqrt{L}, \quad (N = \frac{L}{H}) \quad H = A + \frac{B}{u} + Cu$$

4. Core-shell particle



Beyond HPLC (high performance LC): Ultra performance LC (UPLC)

High pressure → allow to use smaller particle, long column, narrower column
→ **high performance**

HPLC \leq 400 bar, 6000 psi

UPLC (or UHPLC) \geq 690 bar, 10000 psi

Recently, UHPLC w/ 1500 bar (22000 psi) of capacity has been released

History of chromatography

Chromato + graphy = color + writing



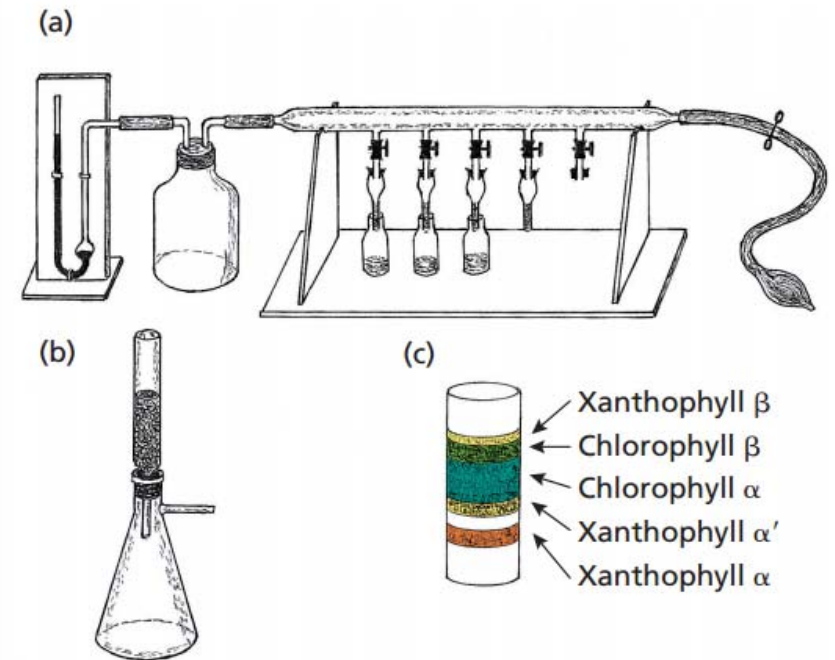
Mikhail Tsvet

Adsorption chromatography

Stationary phase = solid (CaCO_3)

Mobile phase = liquid (CS_2)

Figure 2: Illustrations from Tswett's 1906 paper.^{6,10,11}
(a) Apparatus for the simultaneous use of as many as five columns. The lower part of the small funnel-like glass pieces (2–3 mm i.d. and 20–30 mm length) served as the packed column. (b) Apparatus for larger samples (1–3 cm i.d., packing length: 5–9 cm). (c) Chromatographic separation of plant pigments as drawn by Tswett. Stationary phase: calcium carbonate; eluent: carbon disulphide.



M. Tswett, *Ber. Dtsch. Botan. Ges.*, 24, 384–393 (1906)

History of chromatography : partition chromatography

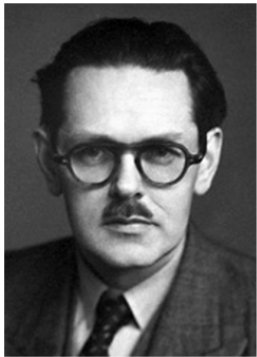


The Nobel Prize in Chemistry 1952

Archer J.P. Martin, Richard L.M. Synge

Share this: 3

The Nobel Prize in Chemistry 1952



Archer John Porter Martin

Prize share: 1/2



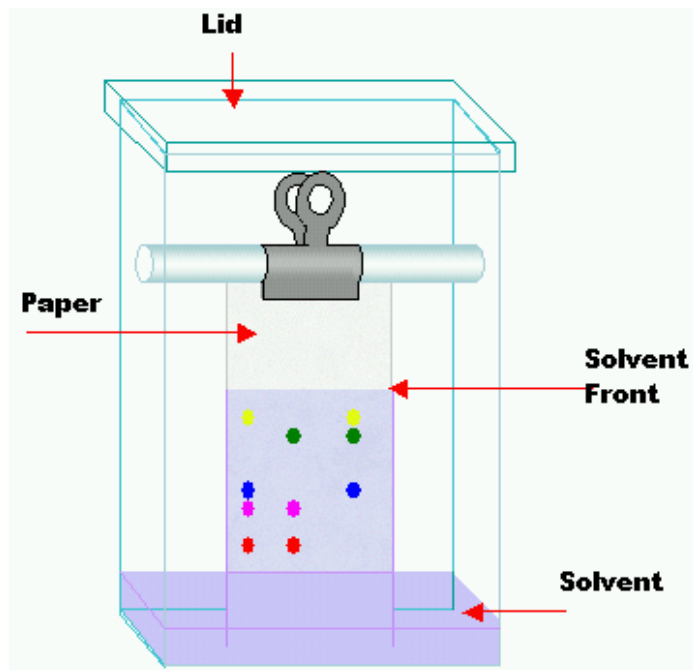
Richard Laurence Millington Synge

Prize share: 1/2

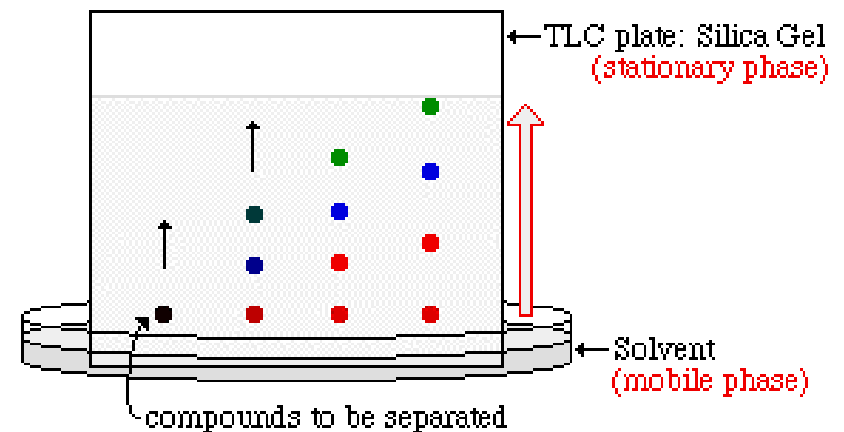
The Nobel Prize in Chemistry 1952 was awarded jointly to Archer John Porter Martin and Richard Laurence Millington Synge "*for their invention of partition chromatography*"

History of chromatography : **partition chromatography**

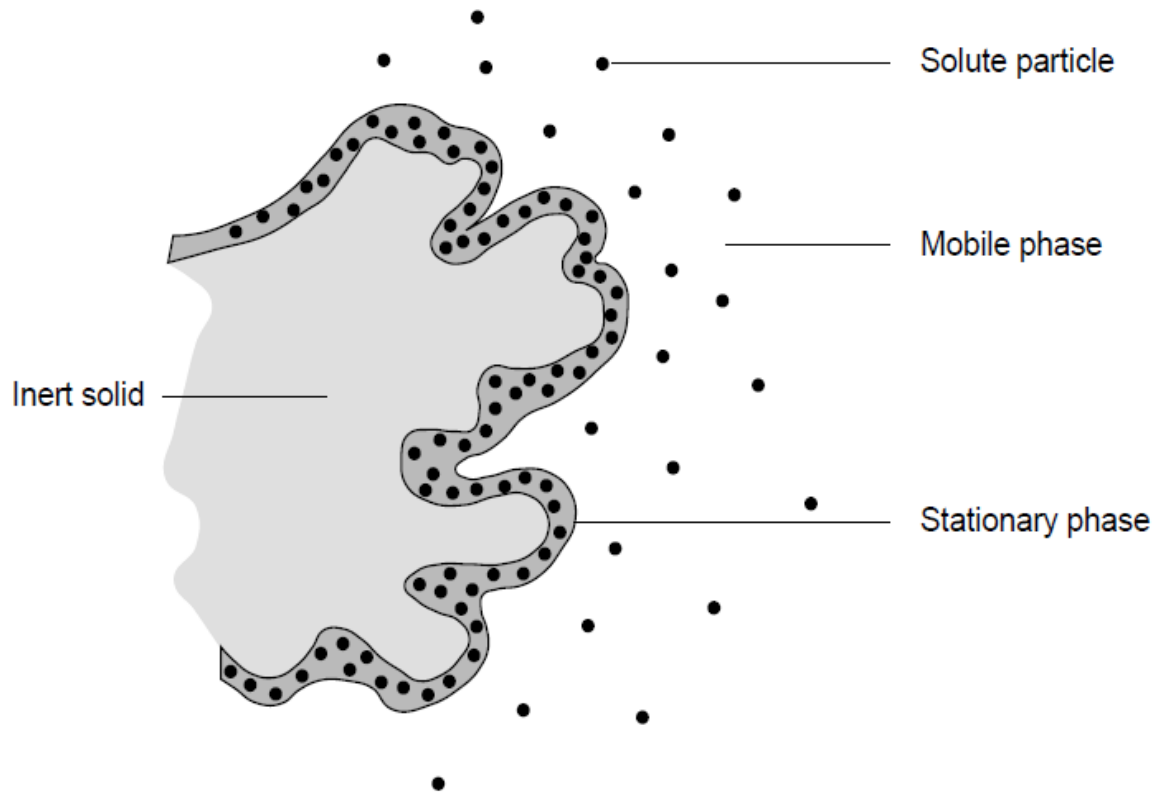
1. Paper chromatography



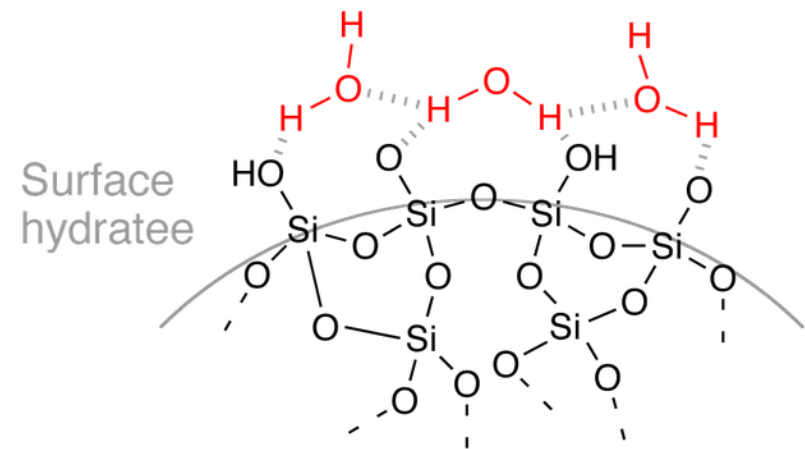
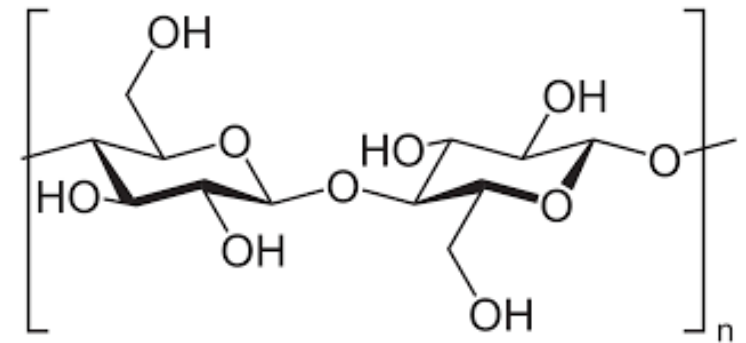
2. Thin layer chromatography



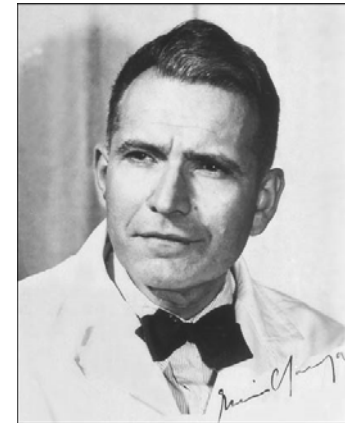
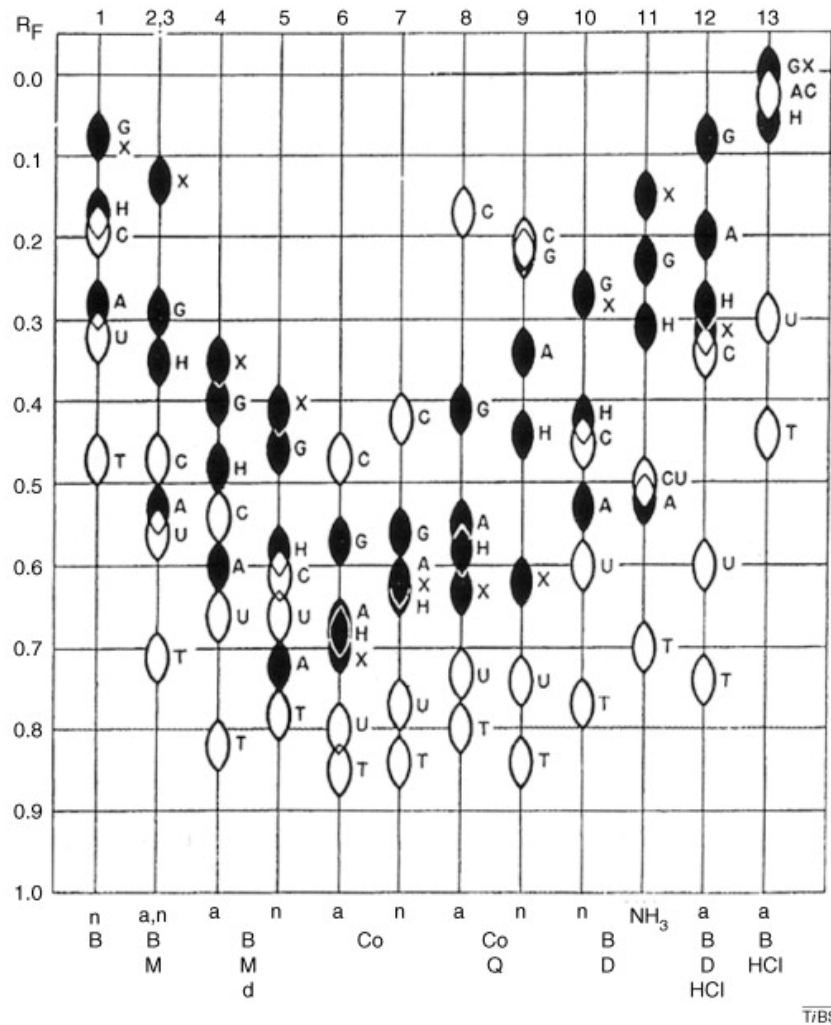
Partition chromatography



Each solute partitions itself between the stationary phase and the mobile phase



Schematic representation of the position of the purines and pyrimidines



Erwin Chargaff

Table 1. Molar proportions of purines and pyrimidines in DNA from different species and organisms^a

Species	A/G	T/C	Pu/Py
Ox <i>Bos taurus</i>	1.29	1.43	1.1
Man <i>Homo sapiens</i>	1.56	1.75	1.0
Wheatgerm <i>Triticum vulgare</i>	1.22	1.18	1.0
Yeast <i>Saccharomyces cerevisiae</i>	1.72	1.9	1.0
Avian tubercle bacillus (<i>Mycobacterium avium</i>)	0.4	0.4	1.1
Haemophilus influenzae, Type C	1.74	1.54	1.0
<i>Escherichia coli</i> , K-12	1.05	0.95	1.0
<i>Serratia Marcescens</i>	0.7	0.7	0.9
Hydrogen organism <i>Bacillus Schatz</i>	0.7	0.6	1.0

Chargaff's rule

King's College, London. One of us (J. D. W.) has been aided by a fellowship from the National Foundation for Infantile Paralysis.

J. D. WATSON
F. H. C. CRICK

Medical Research Council Unit for the
Study of the Molecular Structure of
Biological Systems,
Cavendish Laboratory, Cambridge.
April 2.

- ¹ Pauling, L., and Corey, R. B., *Nature*, **171**, 346 (1953); *Proc. U.S. Nat. Acad. Sci.*, **39**, 84 (1953).
- ² Furberg, S., *Acta Chem. Scand.*, **6**, 634 (1952).
- ³ Chargaff, E., for references see Zamenhof, S., Brawerman, G., and Chargaff, E., *Biochim. et Biophys. Acta*, **9**, 402 (1952).
- ⁴ Wyatt, G. R., *J. Gen. Physiol.*, **36**, 201 (1952).
- ⁵ Astbury, W. T., *Symp. Soc. Exp. Biol.* 1, Nucleic Acid, 66 (Camb. Univ. Press, 1947).
- ⁶ Wilkins, M. H. F., and Randall, J. T., *Biochim. et Biophys. Acta*, **10**, 192 (1953).

Insulin sequencing

The Nobel Prize in Chemistry 1958



Frederick Sanger
Prize share: 1/1

The Nobel Prize in Chemistry 1958 was awarded to Frederick Sanger *"for his work on the structure of proteins, especially that of insulin"*.

The Nobel Prize in Chemistry 1980



Paul Berg
Prize share: 1/2



Walter Gilbert
Prize share: 1/4



Frederick Sanger
Prize share: 1/4

The Nobel Prize in Chemistry 1980 was divided, one half awarded to Paul Berg *"for his fundamental studies of the biochemistry of nucleic acids, with particular regard to recombinant-DNA"*, the other half jointly to Walter Gilbert and Frederick Sanger *"for their contributions concerning the determination of base sequences in nucleic acids"*.

Insulin sequencing

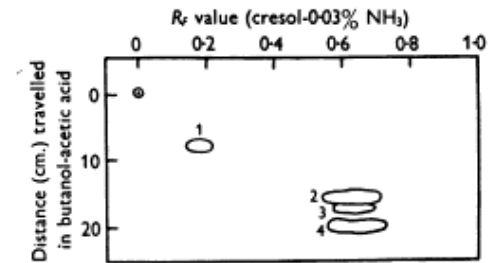
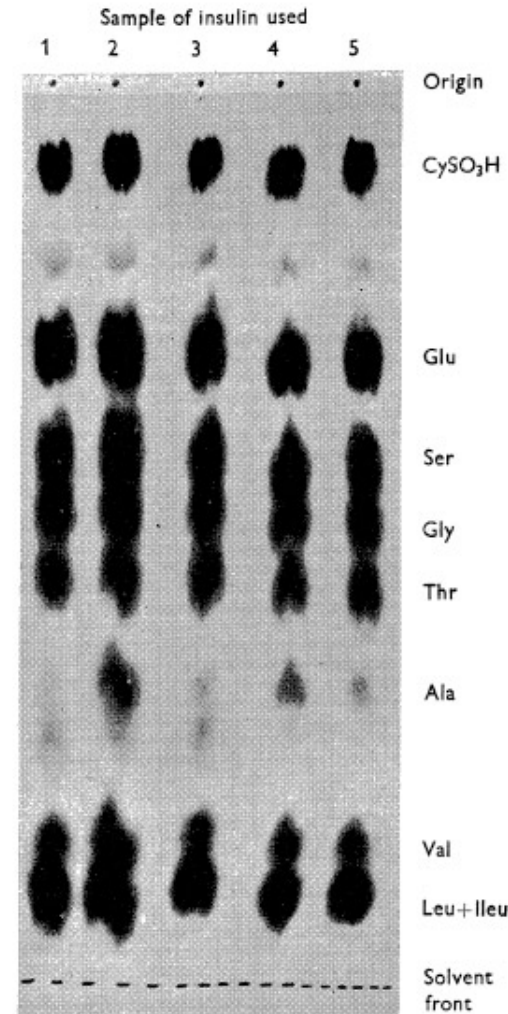
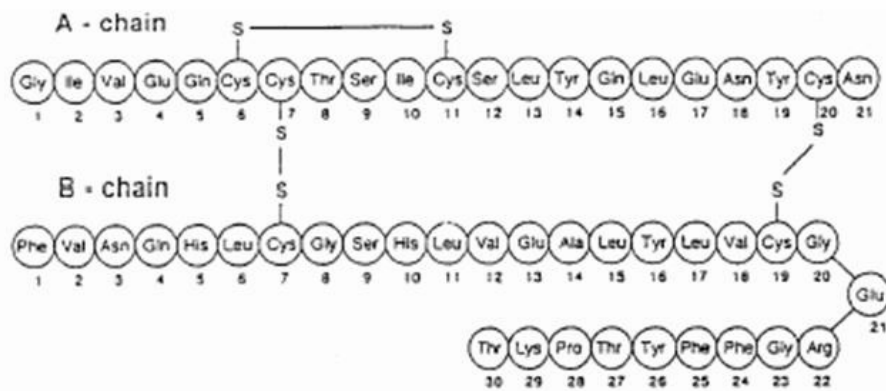


Fig. 3. Chromatogram of the neutral fraction of the chymotryptic hydrolysate (*Bc*) (see Table 2).

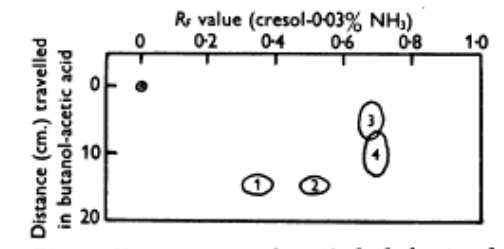


Fig. 5. Chromatogram of peptic hydrolysate of fraction *Bc2* (see Table 2).

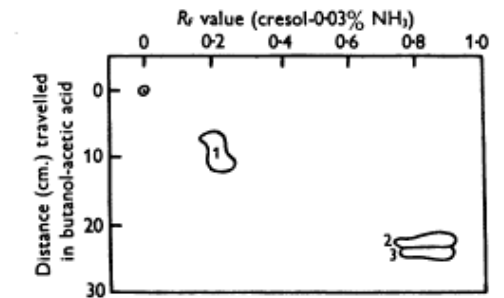
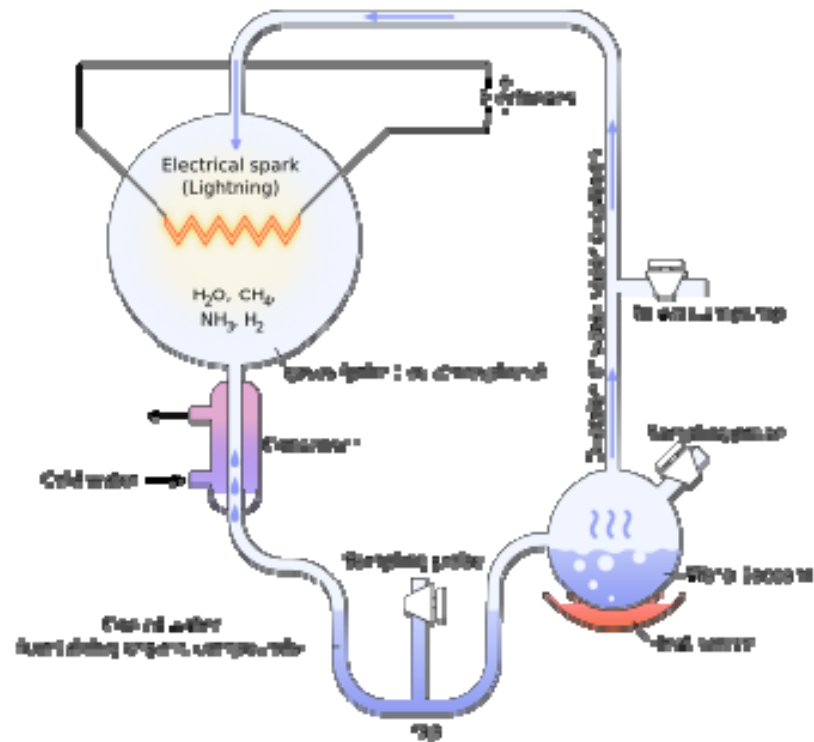
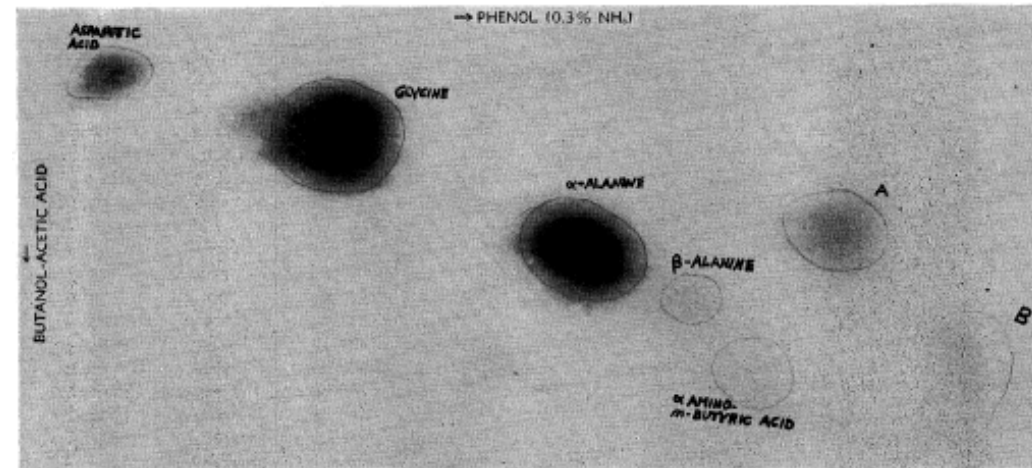


Fig. 6. Chromatogram of the tryptic hydrolysate (*Bt*) (see Table 2).

Miller–Urey experiment



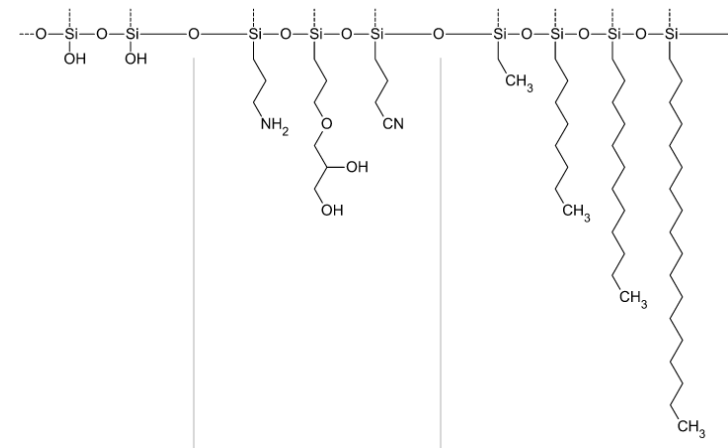
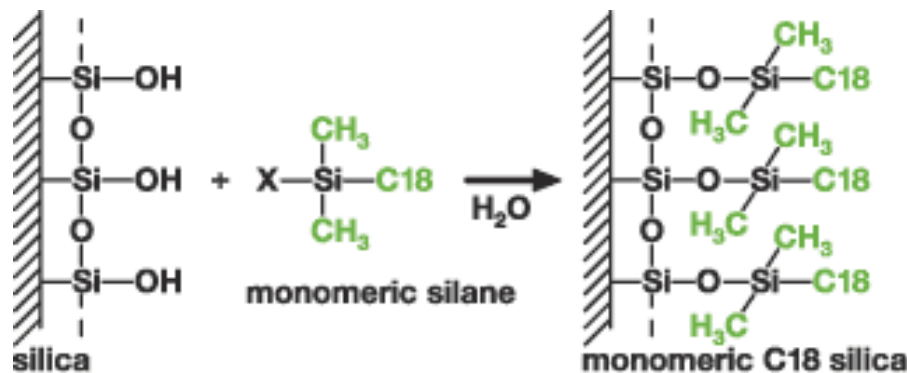
Stanley L Miller



Science 117 (3046): 528–9, 1953

History of chromatography : reversed phase chromatography

- Until ~ 1970s, a **hydrophilic** solid support (silica or alumina) was used as a stationary phase → called “**normal phase** chromatography”
- A technique using alkyl chains covalently bonded to the solid support created a **hydrophobic** stationary phase → **reversed phase** chromatography



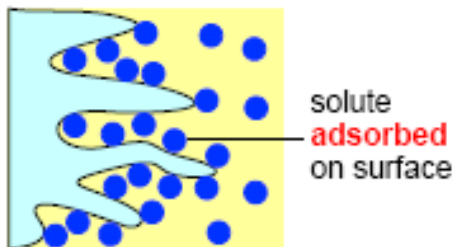
Reversed phase chromatography : **impact !**

- Mobile solvent: organic → aqueous !! (\$\$\$ → \$)
- Amphiphilic molecules can be easily separated: **PEPTIDE**
- **Electrospray-friendly (volatile organic solvent)**

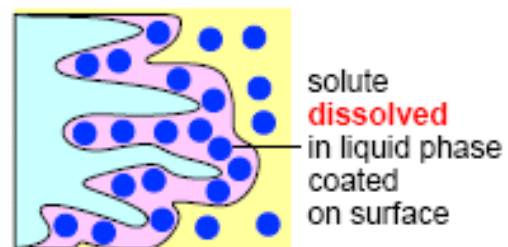
Other types of Liquid Chromatography:

- ② Adsorption chromatography
- ② Partition chromatography
- ② Ion-exchange chromatography

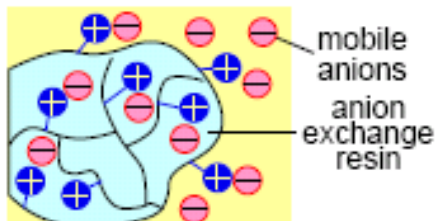
- ② Affinity chromatography
- ② Size-exclusion chromatography



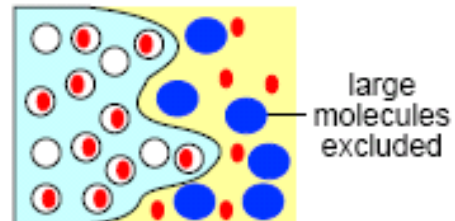
Adsorption Chromatography



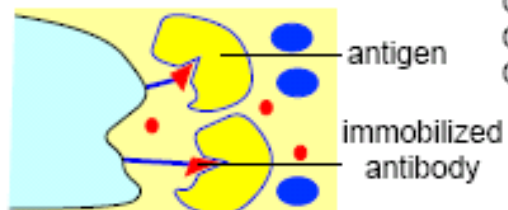
Partition Chromatography



Ion-Exchange Chromatography



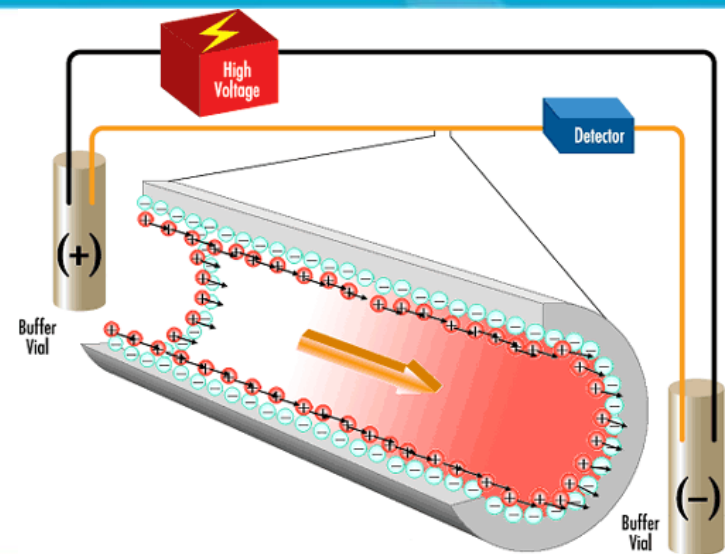
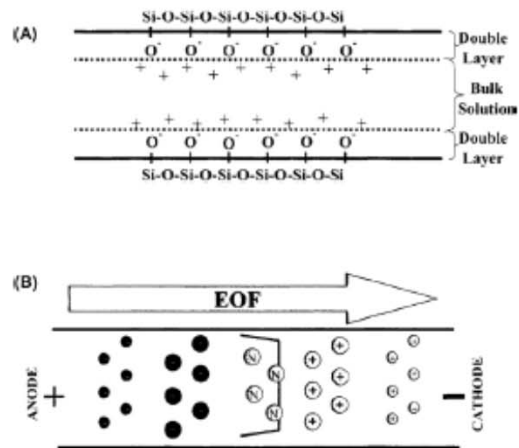
Molecular Exclusion Chromatography
Gel Permeation Chromatography
Gel-Filtration Chromatography
Gel Chromatography



Affinity Chromatography

***Separation tools for
electrospray mass spectrometry
- Capillary Electrophoresis***

ElectroOsmotic Flow (EOF)

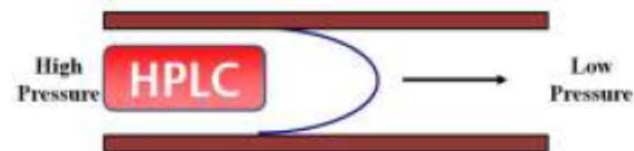


High Efficiency
(높은 분리효율)



Electroosmotic velocity profile

Flat flow



Hydrodynamic velocity profile

Laminar flow

Open Tubular Capillary: How high voltage enhances separation

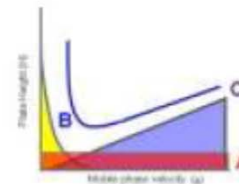
Minimize sample pretreatment

van Deemter equation:

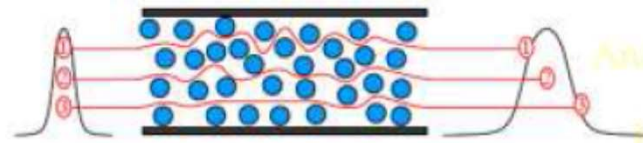
$$H = A + \frac{B}{u} + Cu$$

$$H = A \cdot d_p + \frac{B}{\mu} + C \cdot d_s^2 \cdot \mu$$

Eddy Diffusion Multi-path Effect Longitudinal Diffusion Mass Transfer (Kinetics)



A

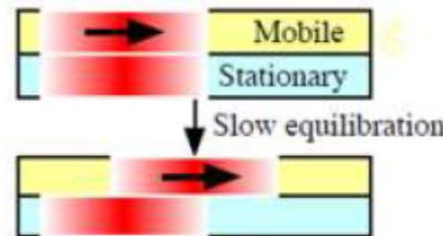


Multiple Path
다통로

B

Longitudinal
Diffusion
확산

C



Mass Transfer
질량이동

Open Tubular Column

van Deemter equation:

$$H = \cancel{A} + \frac{B}{u} + \cancel{Cu}$$

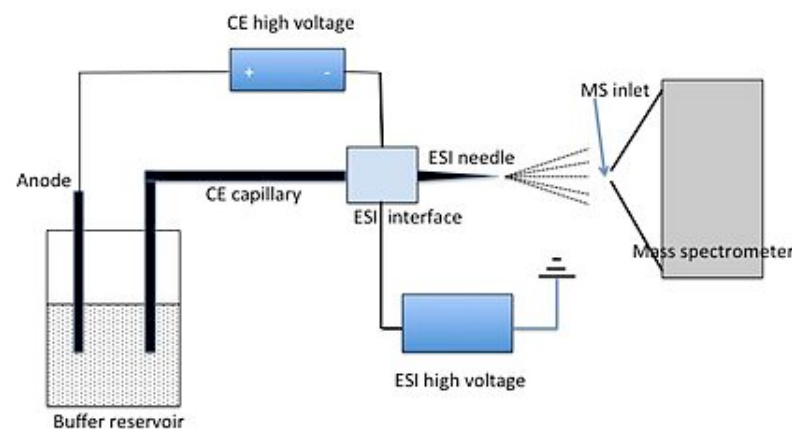
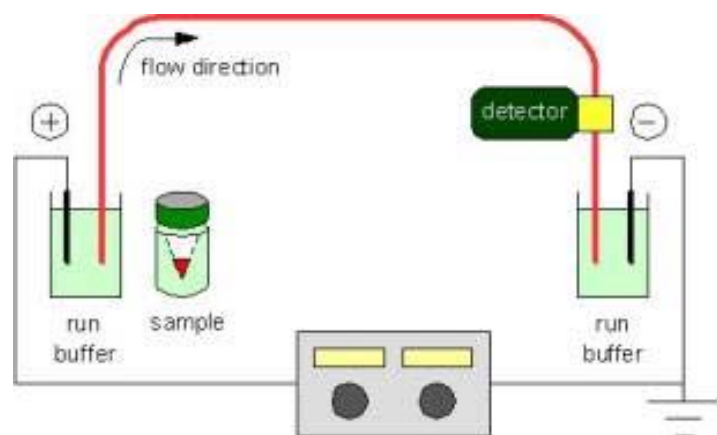
+

Flat
Flow

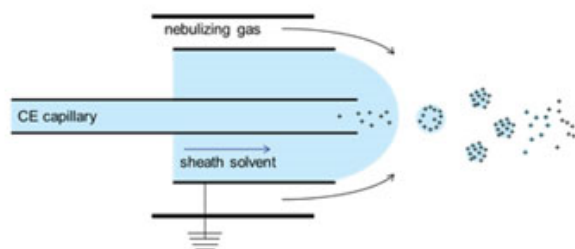
Sharp Peaks

SCIEV

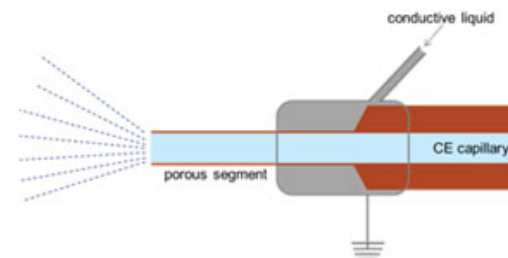
Capillary electrophoresis mass spectrometry (CE-MS)



Sheath flow interface



Sheathless interface



Sheath liquid-less CE-MS

