



DNA separation

- Introduction to separation science
- Conventional gel electrophoresis
- Conventional pulsed-field gel electrophoresis
- Capillary electrophoresis
- Advanced DNA separation techniques
(On-chip electrophoresis and pulsed-field electrophoresis, DNA prism, etc.)

References:

- JC Giddings, *"Unified Separation Science"* (Wiley-Interscience, New York, 1991)
- *Electrophoresis* (Journal)

The big column for separation...

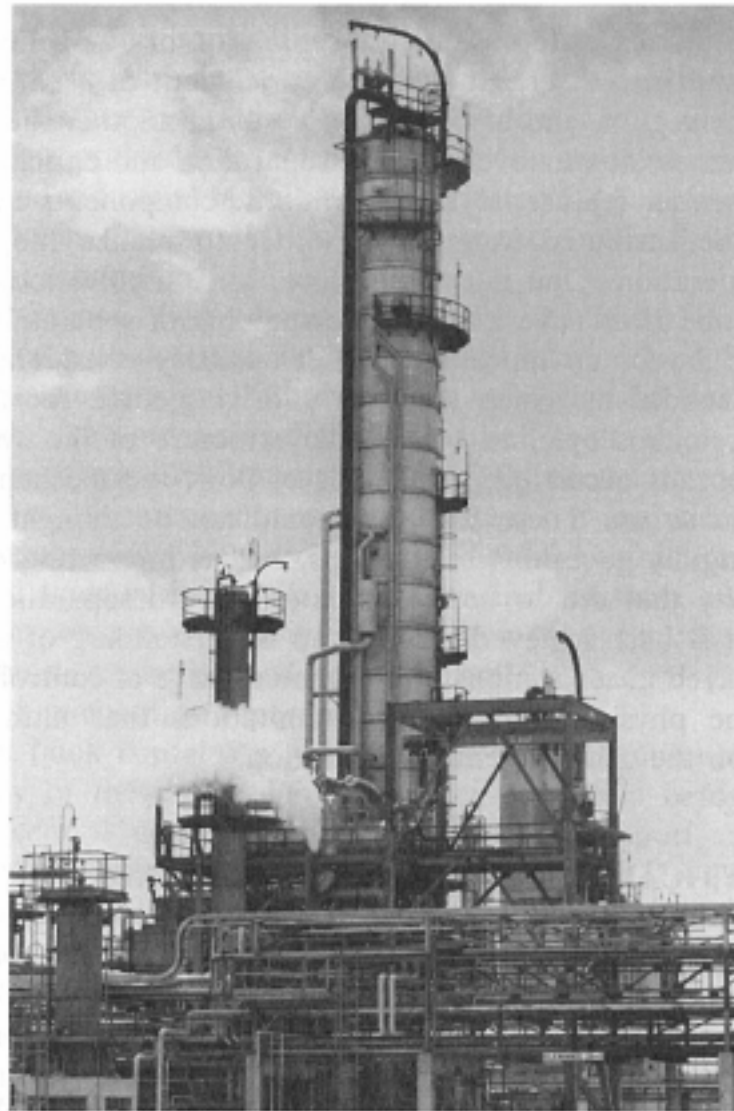


Figure 1.1. Crude fractionation tower at the Phillips Oil Refinery in Woods Cross, Utah. In this crude fractionator, 25,000 barrels per day of incoming crude oil is divided into six fractions by distillation. (Photo by Alexis Kelner.)



Figure 1.2. Preparative scale liquid chromatography unit with 6 ft (1.8 m) long column (in center) having an inner diameter of 6 in (15 cm). Up to 2 L of sample solution can be injected and processed in a single 90-min run. (Courtesy Thomas J. Filipi and Whatman Chemical Separation Division.)

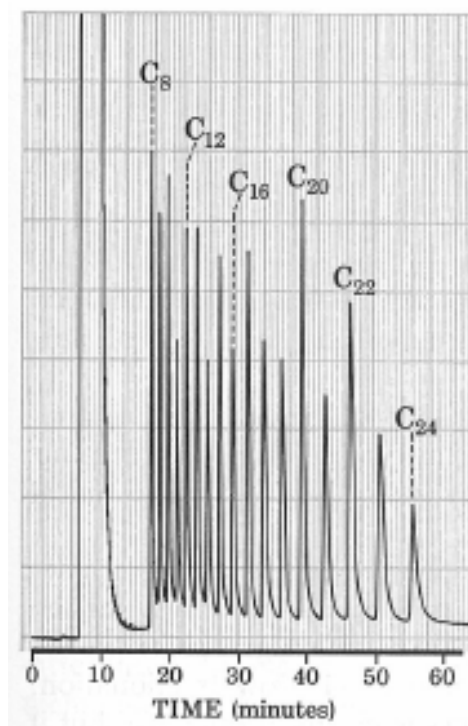
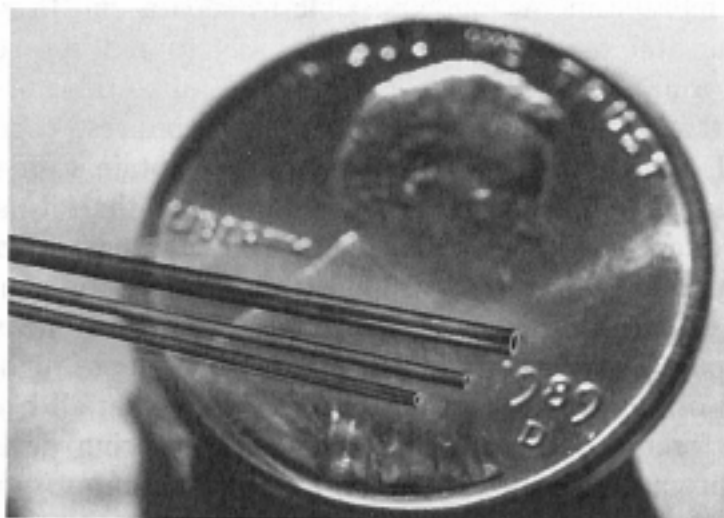
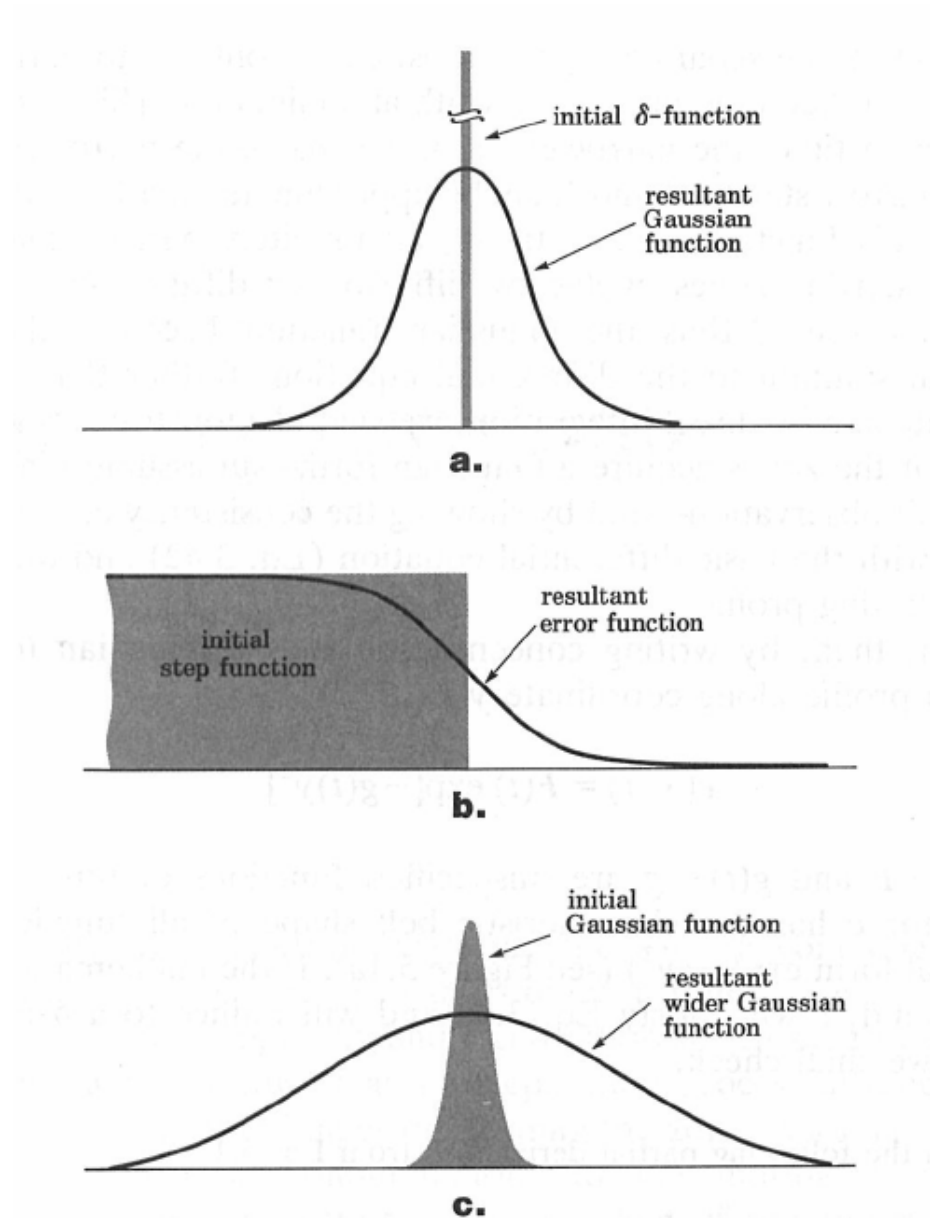


Figure 1.4. High-resolution separation of free fatty acids of indicated carbon numbers by supercritical fluid chromatography using a compressed CO_2 mobile phase at 170°C . Column is 50 cm long fused silica capillary tube of $250\ \mu\text{m}$ internal diameter. (Courtesy of Frank J. Yang.)

Concentration (band) broadening by diffusion



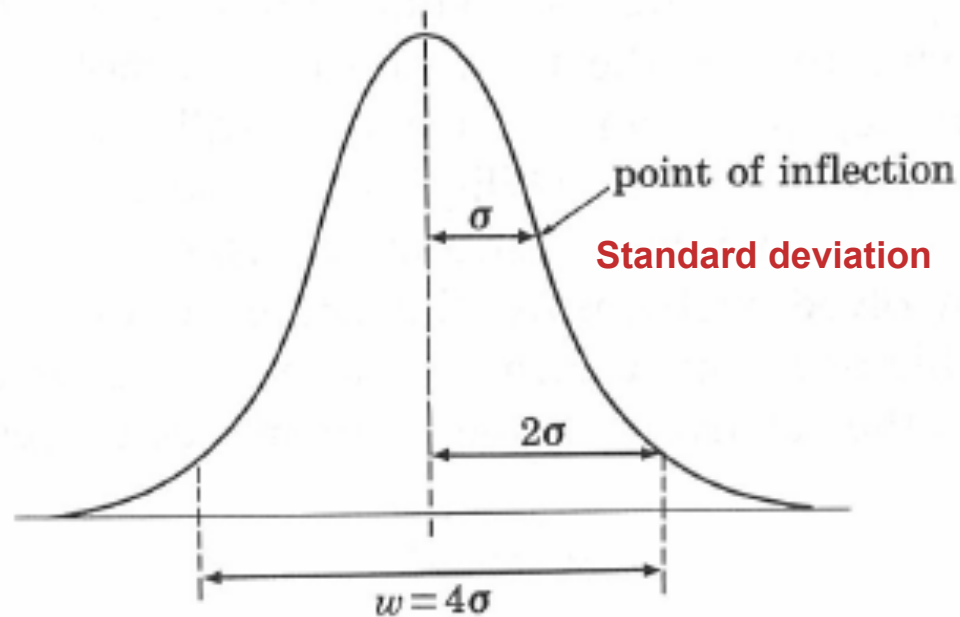
Normalized Gaussian Profile

$$c = \frac{1}{(4\pi Dt)^{1/2}} \exp\left(\frac{-y^2}{4Dt}\right)$$

$$c = \frac{n}{(4\pi Dt)^{1/2}} \exp\left(-\frac{y^2}{4Dt}\right)$$

$$\sigma^2 = 2Dt$$

$$\sigma = (2Dt)^{1/2}$$



Effective zone width



$$X = v t$$

$$\sigma^2 = (2D/v) X$$

Plate height: H 

$$\sigma^2 = H X$$

$$H = \sigma^2 / X$$

Theoretical plates: $N = X / H$

Resolution

$$R_s = \frac{\Delta X}{4\sigma}$$

$$\sigma = (HX)^{1/2}$$

$$R_s = \frac{\Delta \nu}{\nu} \left(\frac{X}{16H} \right)^{1/2} = \frac{\Delta \nu}{\nu} \left(\frac{N}{16} \right)^{1/2}$$

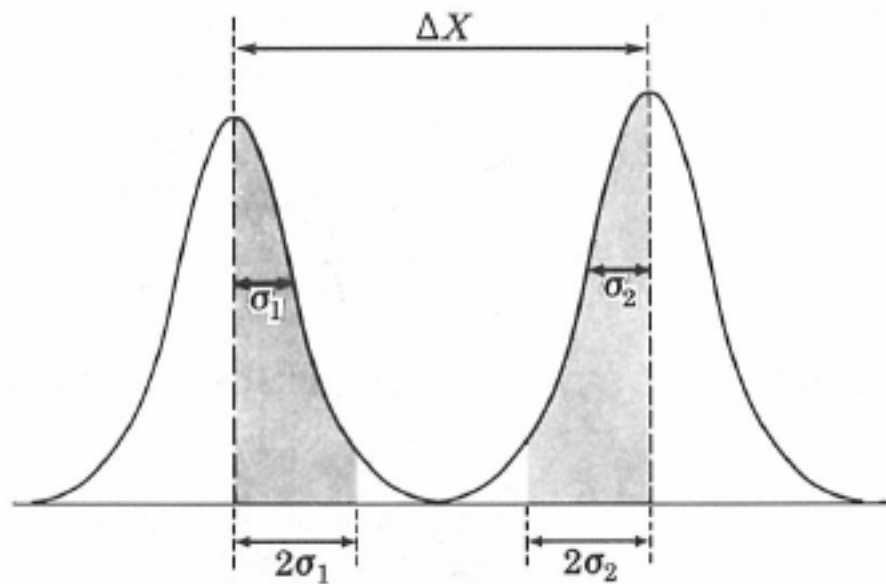


Figure 5.7. The resolution of two peaks, defined as $R_s = \Delta X / (2\sigma_1 + 2\sigma_2)$, measures peak separation ΔX relative to the zone dispersion ($2\sigma_1$ and $2\sigma_2$) leading to overlap and contamination.

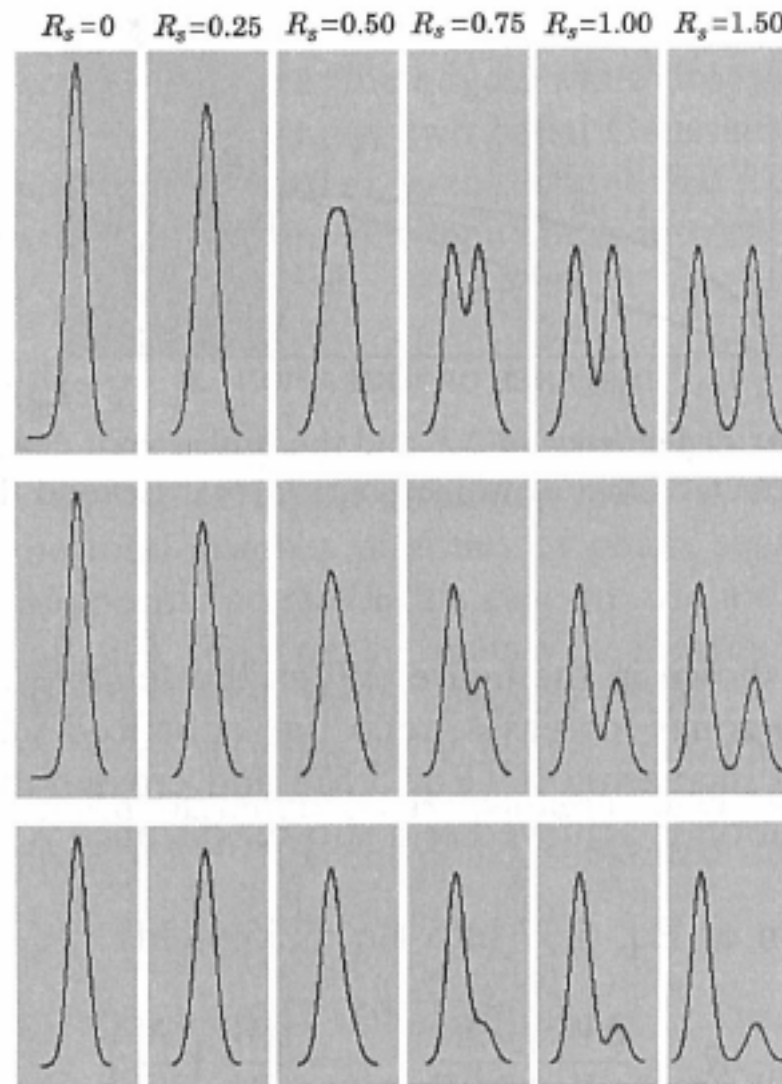


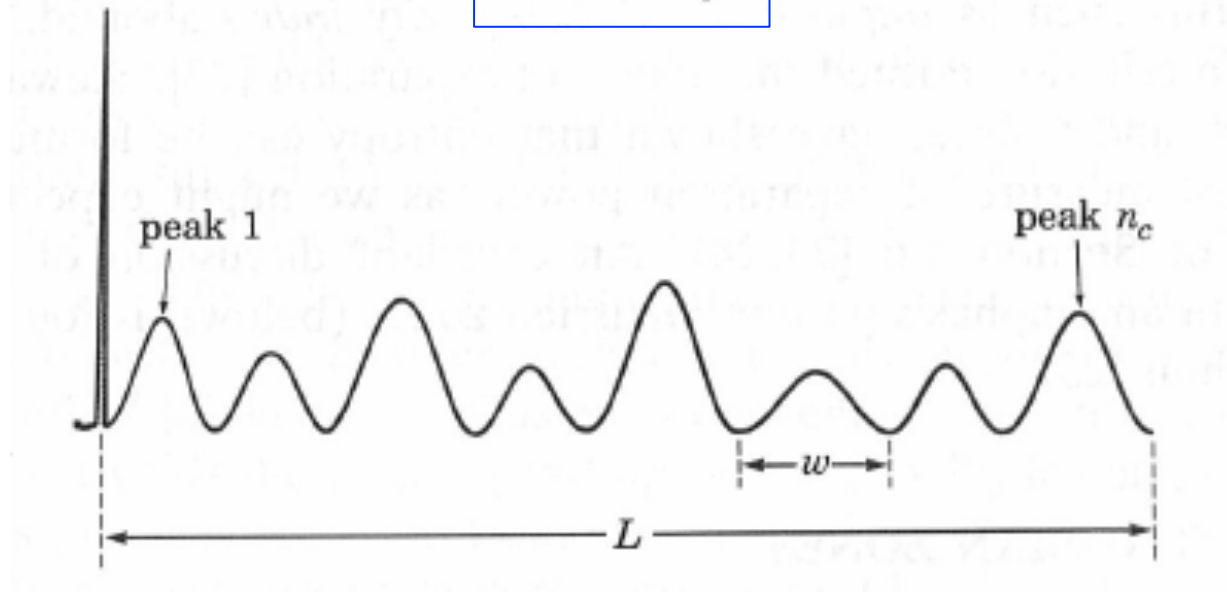
Figure 5.8. Left to right sequence gives the profile of a pair of Gaussian zones of equal σ at increasing levels of resolution, as shown. Top row shows this sequence for two zones of equal peak height; middle and bottom rows show the sequence for 2:1 and 5:1 peak height ratios (profiles courtesy of Joe M. Davis).



Peak capacity: $n_c = L/w = L/4\sigma$ (at $R_s = 1$)

the max number of separated peaks that can be fit (with adjacent peaks at some specified R_s value) into the path length.

$$n_c = \frac{L}{4\sigma R_s}$$



Theoretical plates: $N = L / H$

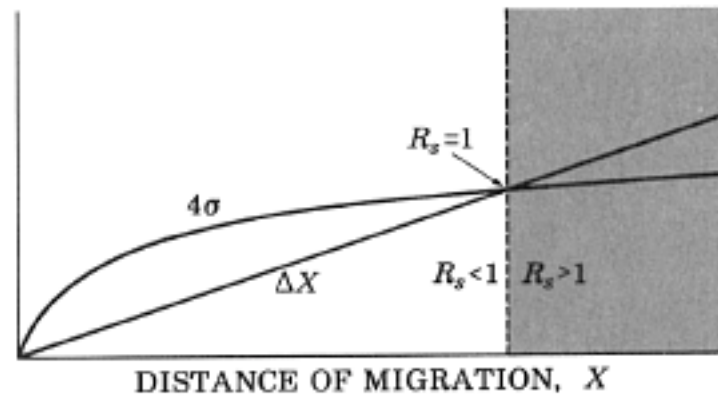


Figure 5.9. The linear dependence of ΔX and the square root dependence of 4σ on X lead to their inevitable crossing, at which point $R_s = 1$. Beyond the crossing (shaded area), $R_s > 1$.

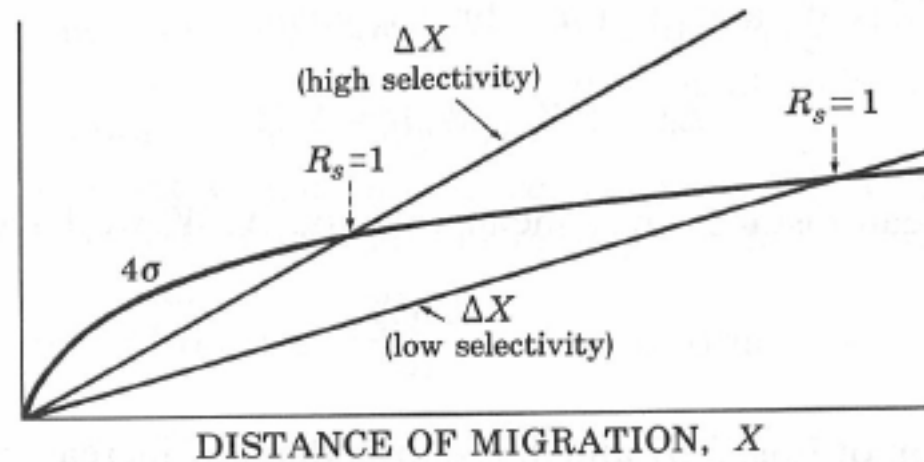




TABLE 8.1 Types of “Fields” that Cause Relative Displacement and thus (potentially at least) Some Degree of Relative Separation^a

| Field or Gradient | Static (<i>Sc</i>) Separation Methods or Phenomena | Other (non- <i>Sc</i>) Separation Methods |
|-------------------------------|---------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------|
| 1. Electrical | Electrophoresis, isoelectric focusing, isotachophoresis | Electrodialysis, electrodeposition, electrostatic precipitation, countercurrent electrophoresis, electrical FFF ^b |
| 2. Sedimentation | Rate-zonal sedimentation, isopycnic sedimentation | Equilibrium sedimentation, centrifugal elutriation, sedimentation FFF ^b |
| 3. Temperature gradient | Thermophoresis | Thermogravitational methods, thermal FFF ^b |
| 4. Electrical (nonuniform) | Dielectrophoresis | |
| 5. Magnetic (nonuniform) | Magnetophoresis | Mass spectroscopy ^c , magnetic separations, magnetic FFF ^a |
| 6. em radiation | Photophoresis | |
| 7. Concentration gradient | Diffusophoresis | |

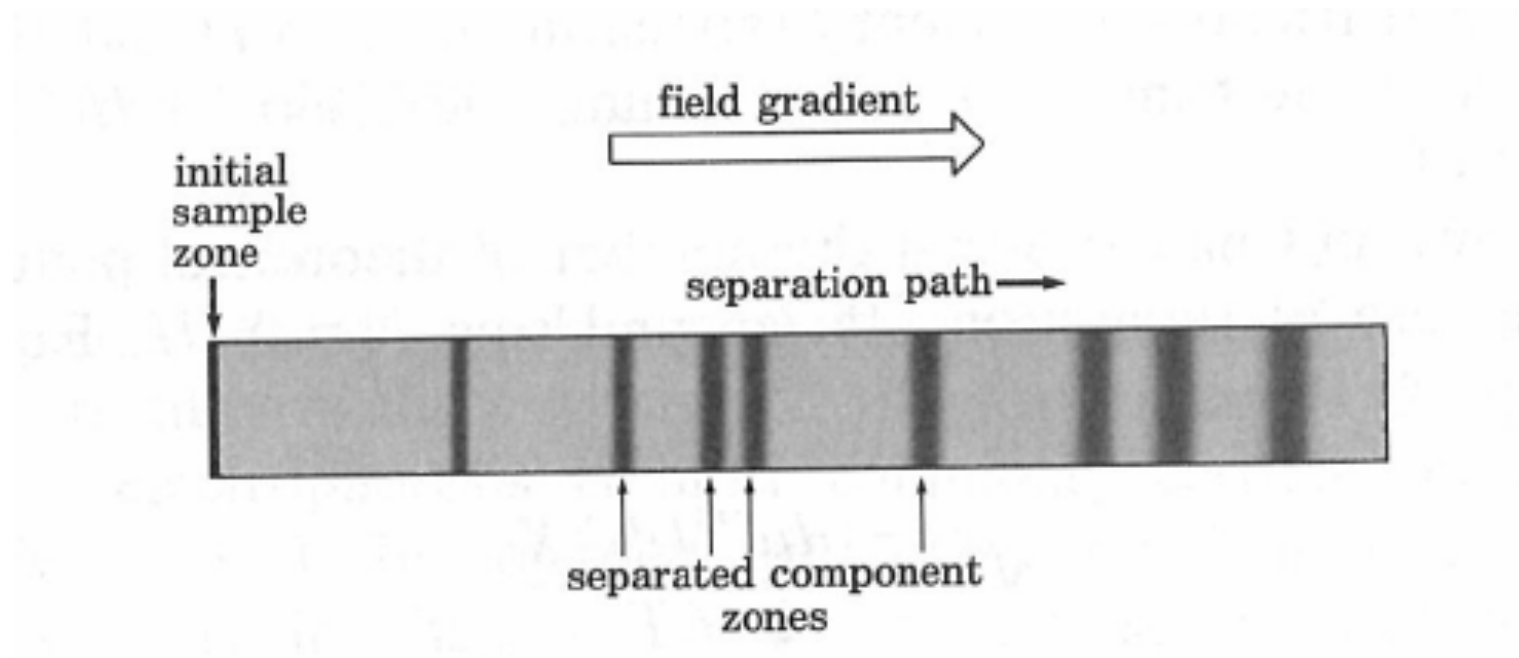
^aPrincipal separation techniques, if they exist, are listed for each field.

^bFFF = field-flow fractionation

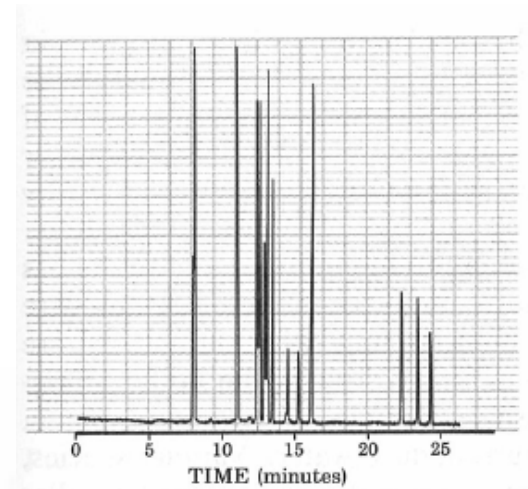
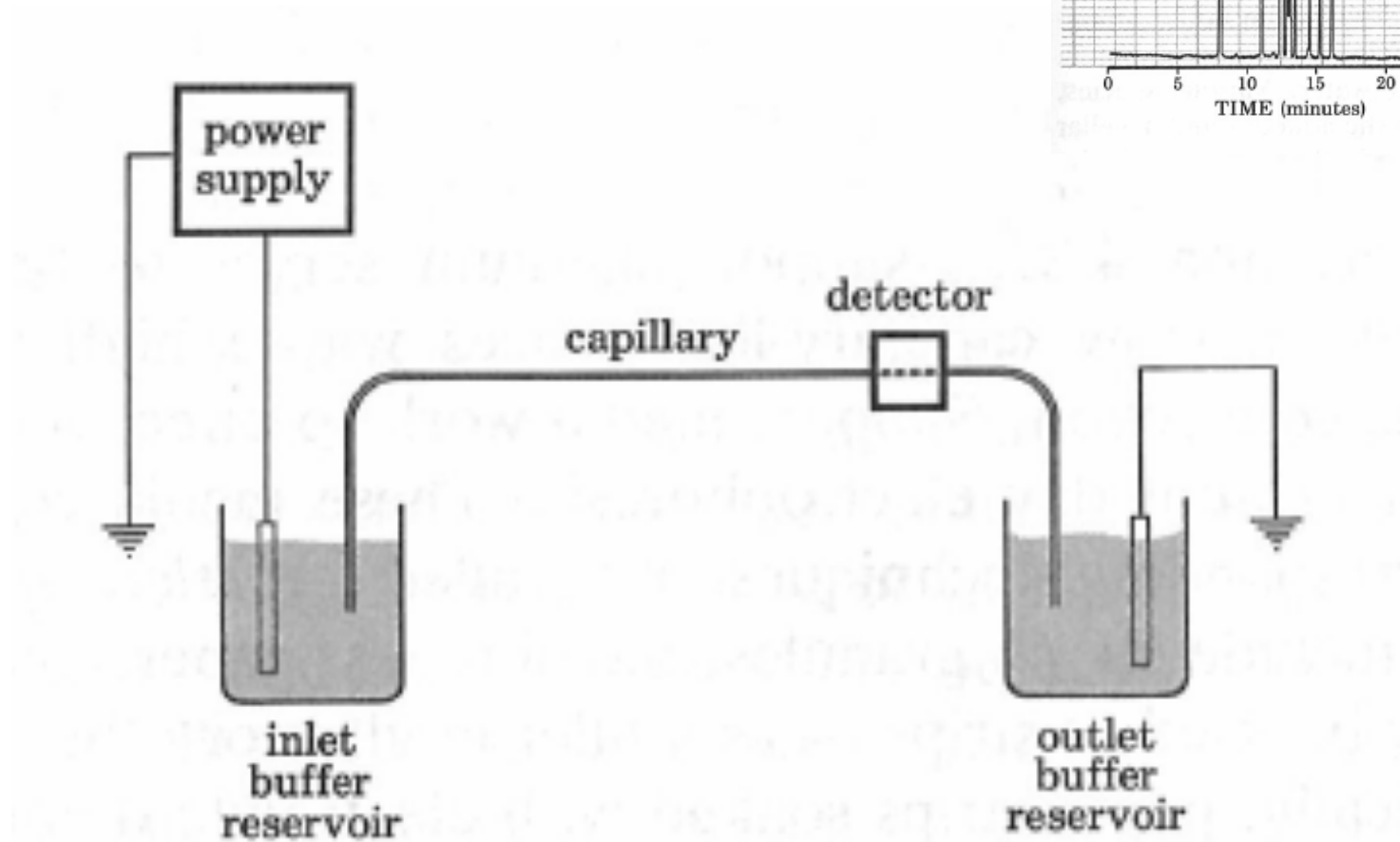
^cInertial transport term important



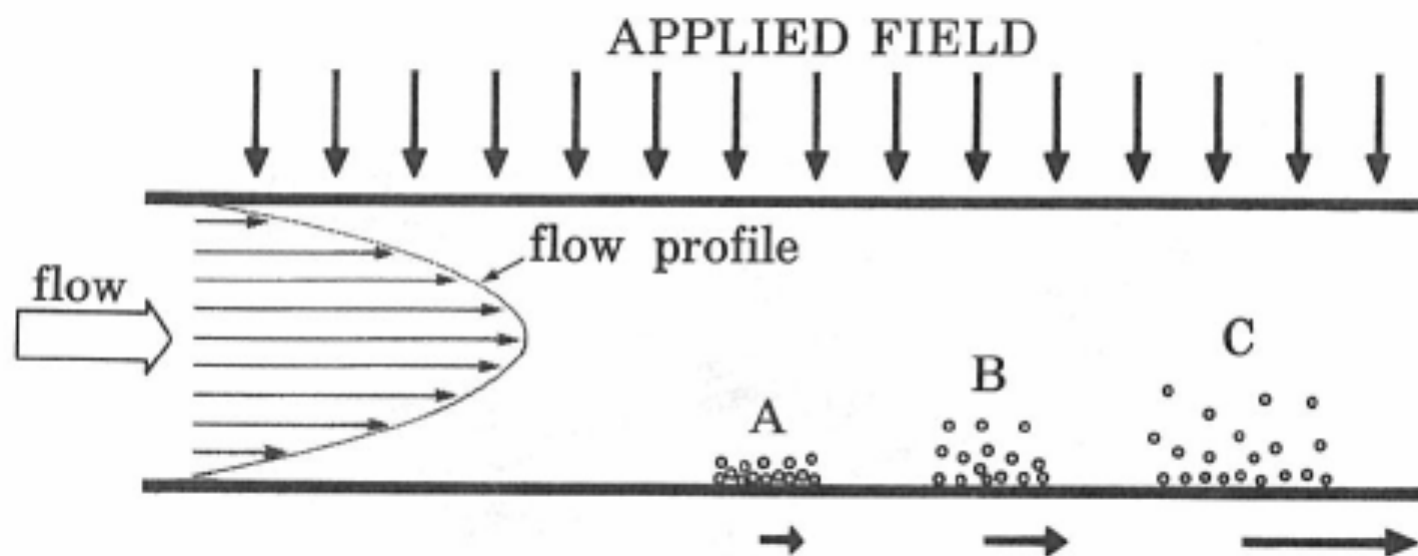
Electrophoresis

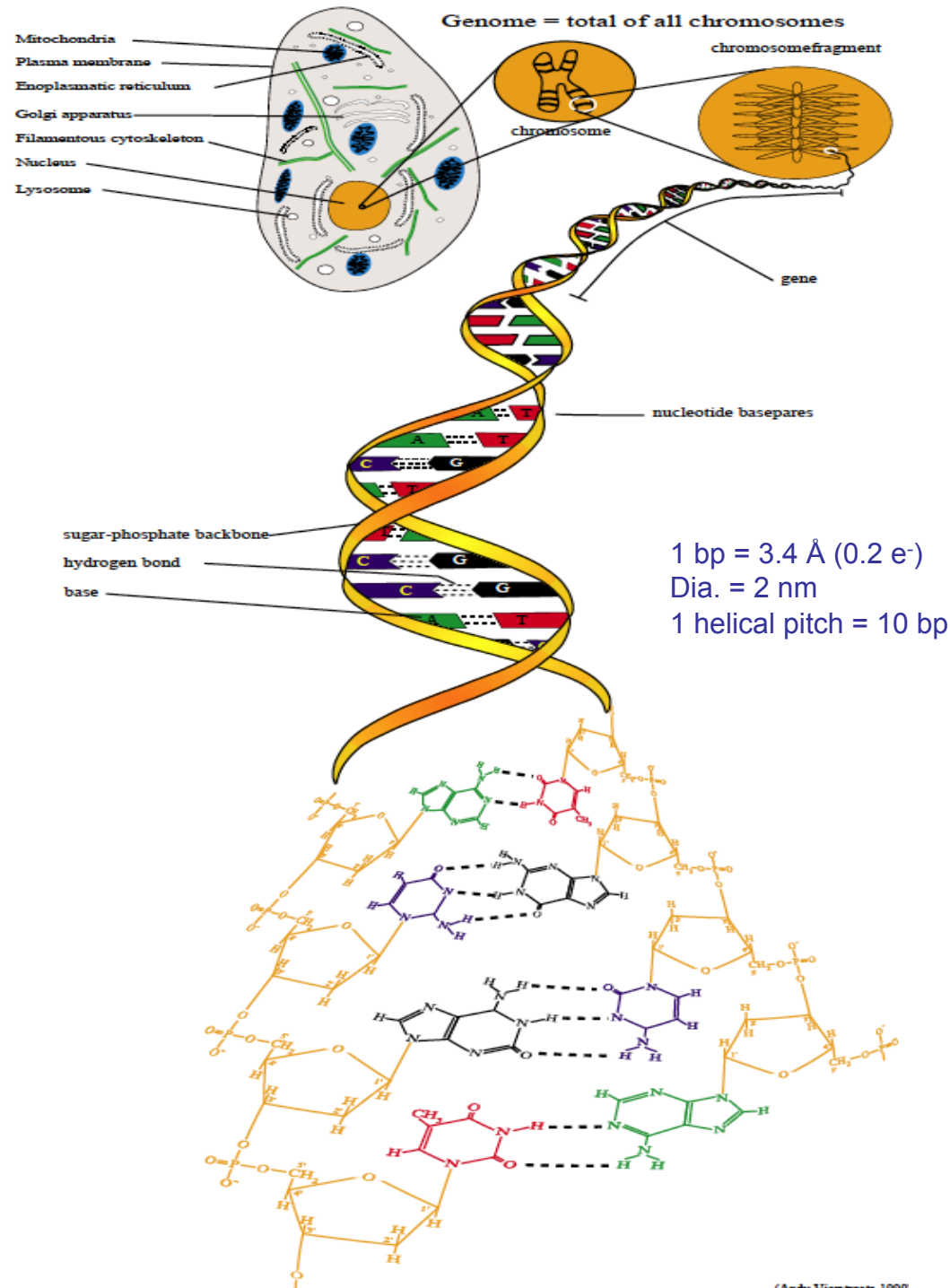


Capillary electrophoresis

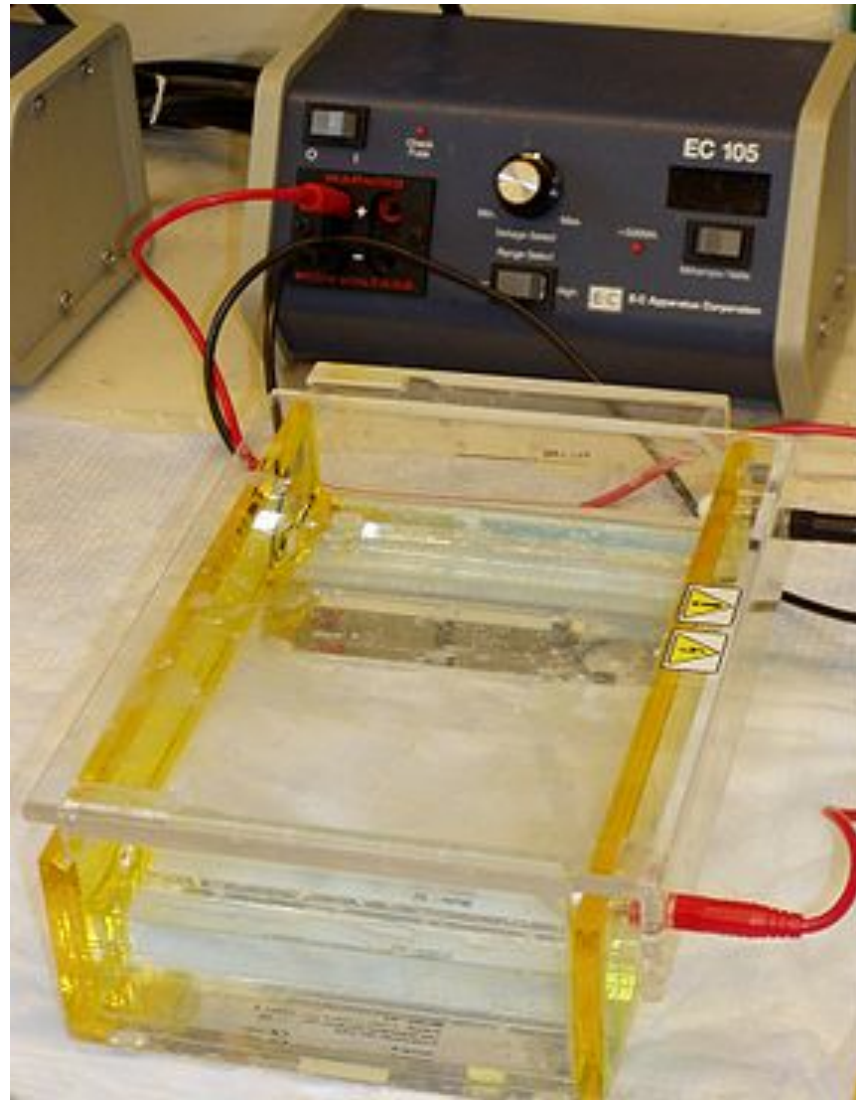


Field flow fractionation (FFF)

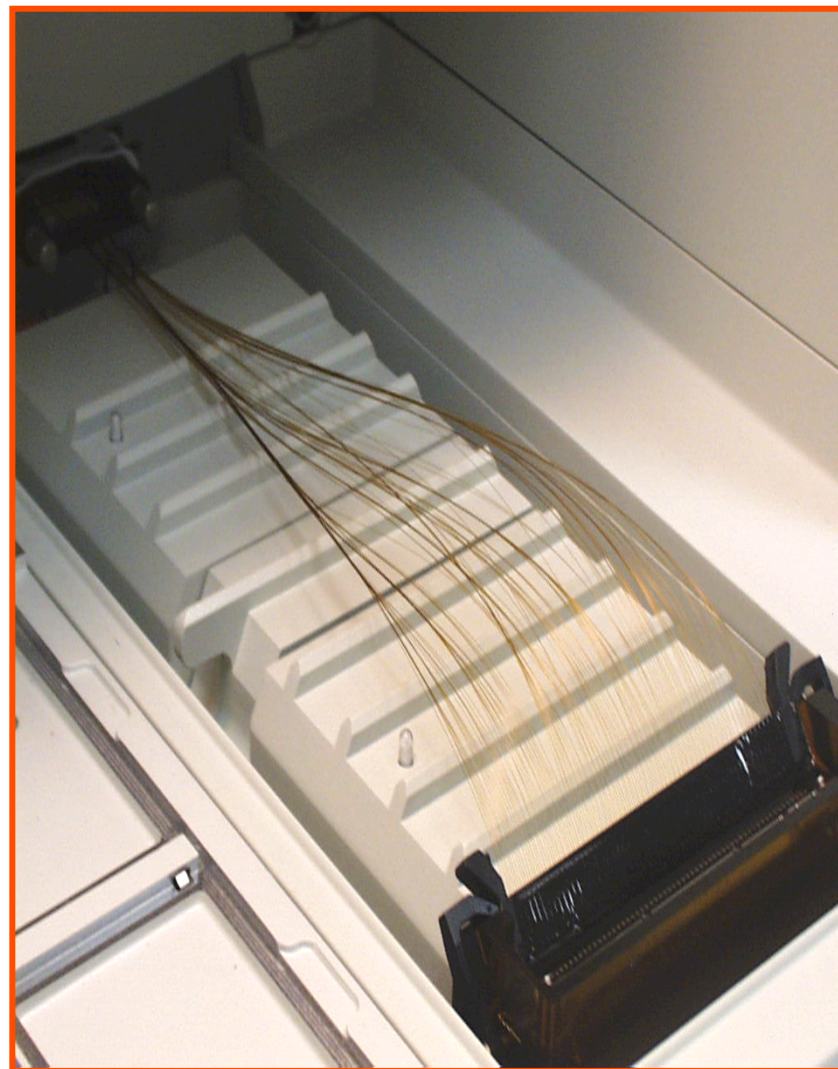
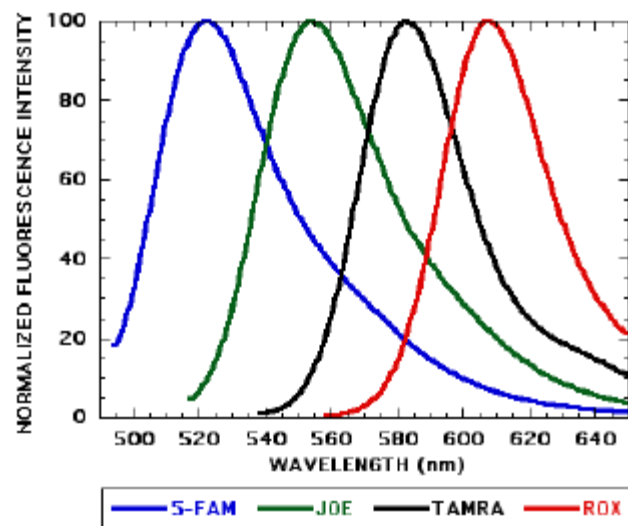
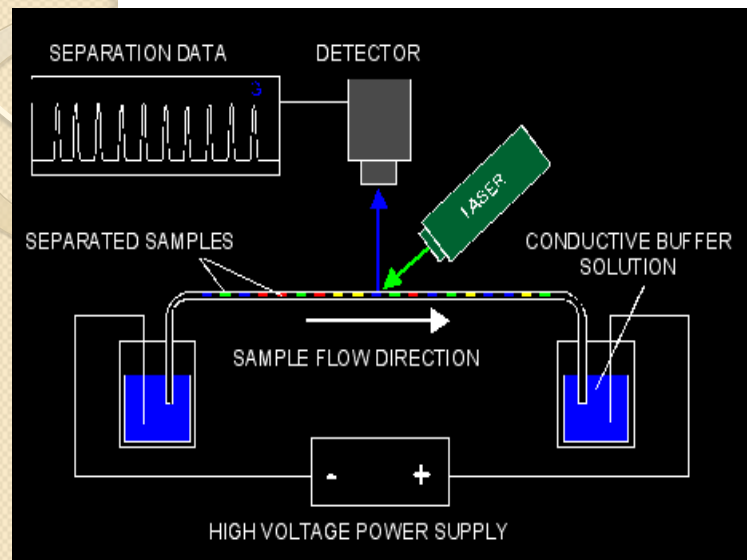




Gel electrophoresis apparatus

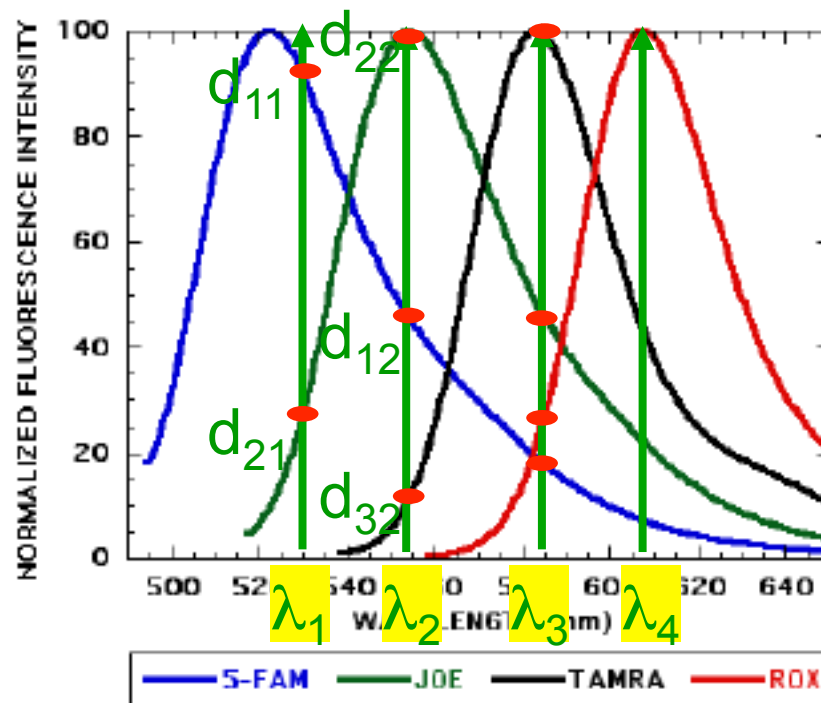


Capillary (Array) Electrophoresis



CE DNA Readout → Least-square fitting

What is needed: optical spectrum of 4 channels vs. time



Calibrated Coefficient:

$d_{11}, d_{21}, d_{31}, d_{41}, \dots$

$$I(\lambda_1) = d_{11}c_1 + d_{21}c_2 + d_{31}c_3 + d_{41}c_4$$

$$I(\lambda_2) = d_{12}c_1 + d_{22}c_2 + d_{32}c_3 + d_{42}c_4$$

$$I(\lambda_3) = d_{13}c_1 + d_{23}c_2 + d_{33}c_3 + d_{43}c_4$$

$$I(\lambda_4) = d_{14}c_1 + d_{24}c_2 + d_{34}c_3 + d_{44}c_4$$

$$I(\lambda_5) = d_{15}c_1 + d_{25}c_2 + d_{35}c_3 + d_{45}c_4$$

c_1, c_2, c_3, c_4 are the dye (DNA) concentration



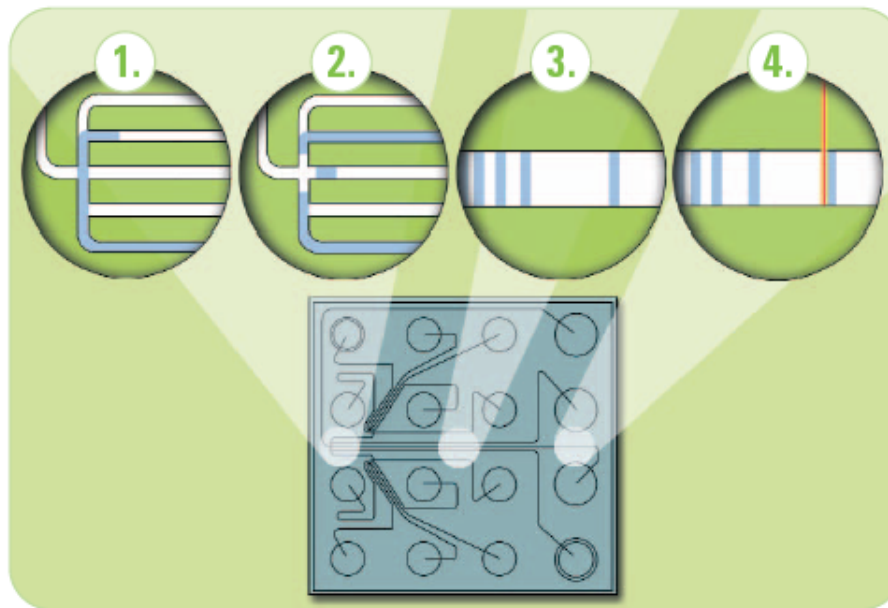
Agilent 2100 Bioanalyzer

Applied microfluidic lab-on-a-chip technology

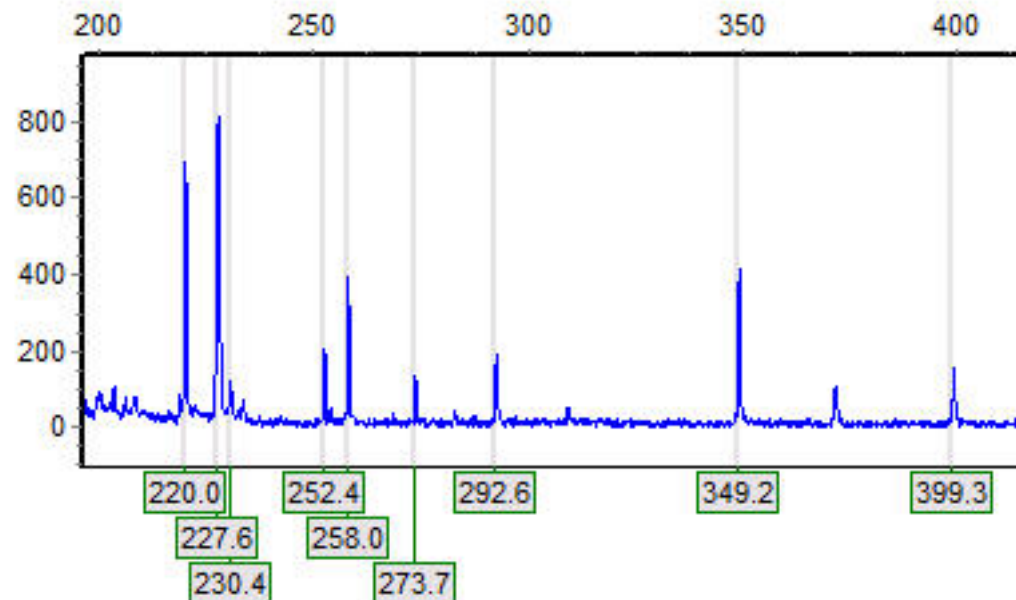




Principle of lab-on-a-chip operation

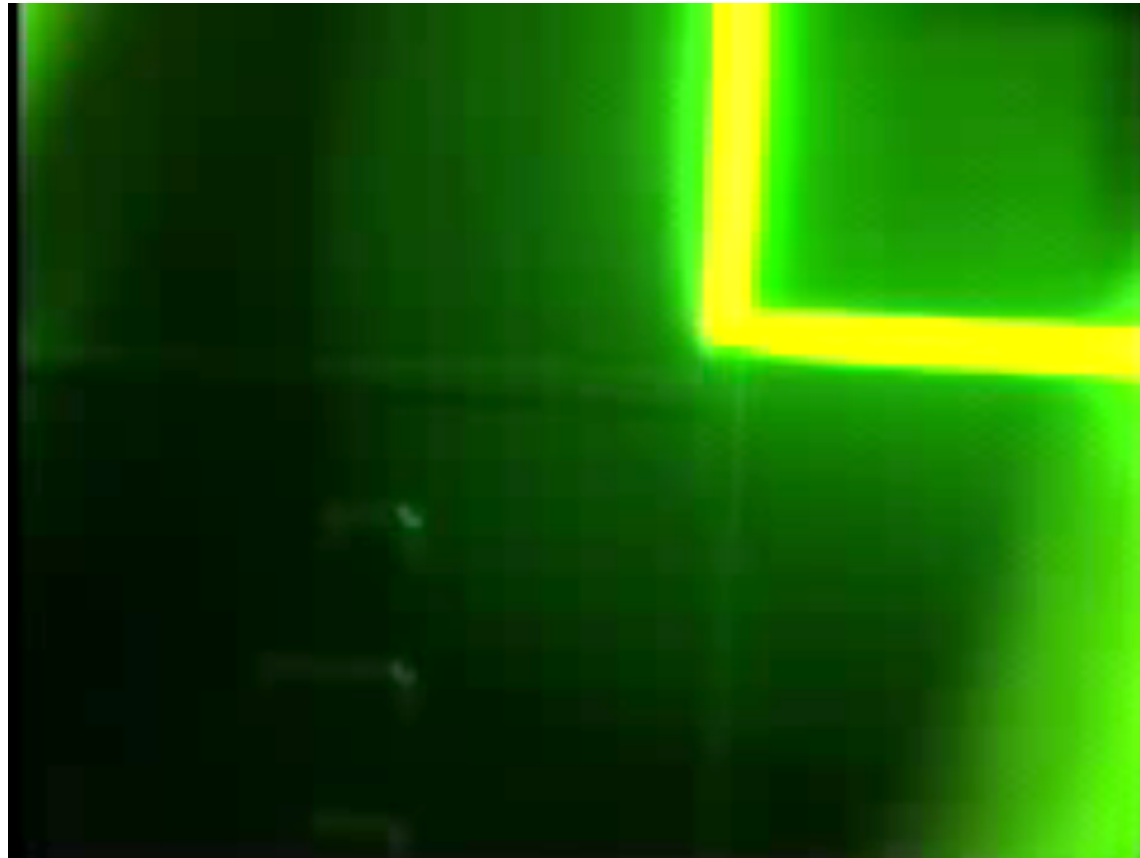


1. The sample moves through the microchannels from the sample well.
2. The sample is injected into the separation channel.
3. Sample components are electrophoretically separated.
4. Components are detected by their fluorescence and translated into gel-like images (bands) and electropherograms (peaks).





Micro-CE injection





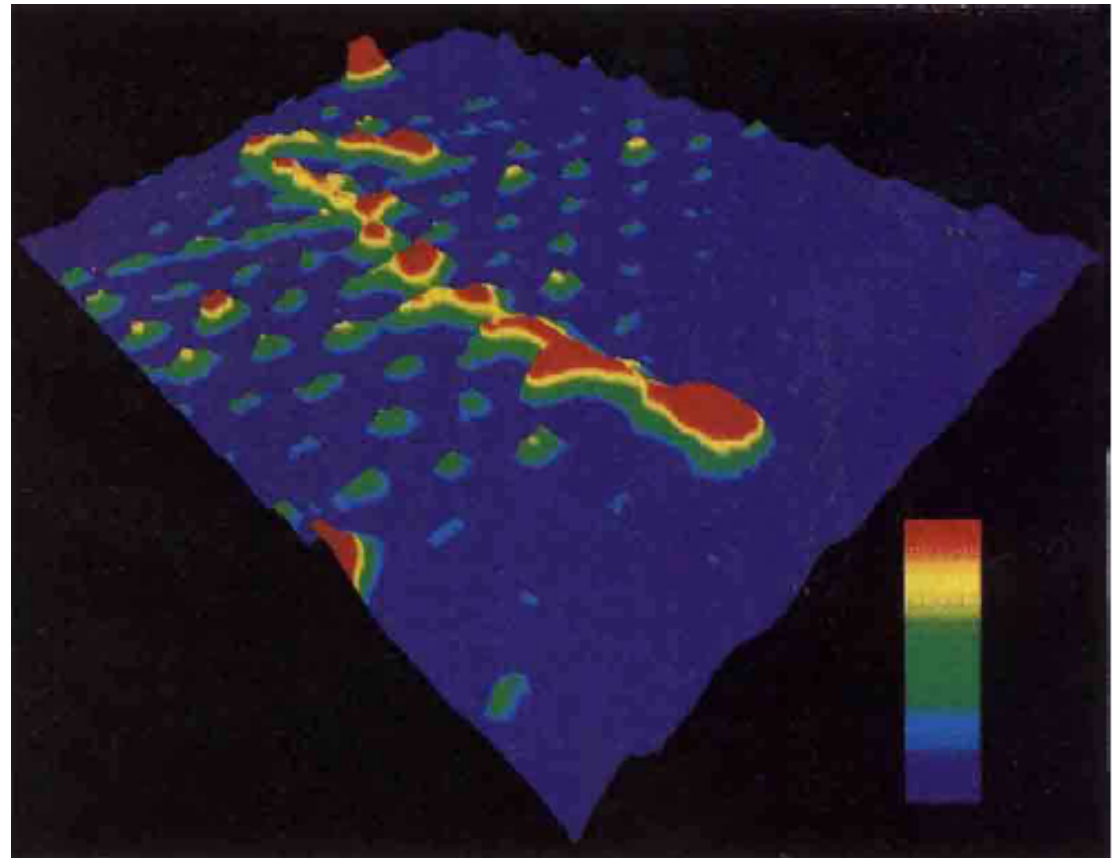
LETTERS TO NATURE

Nature 358, 600 (1992)

DNA electrophoresis in microlithographic arrays

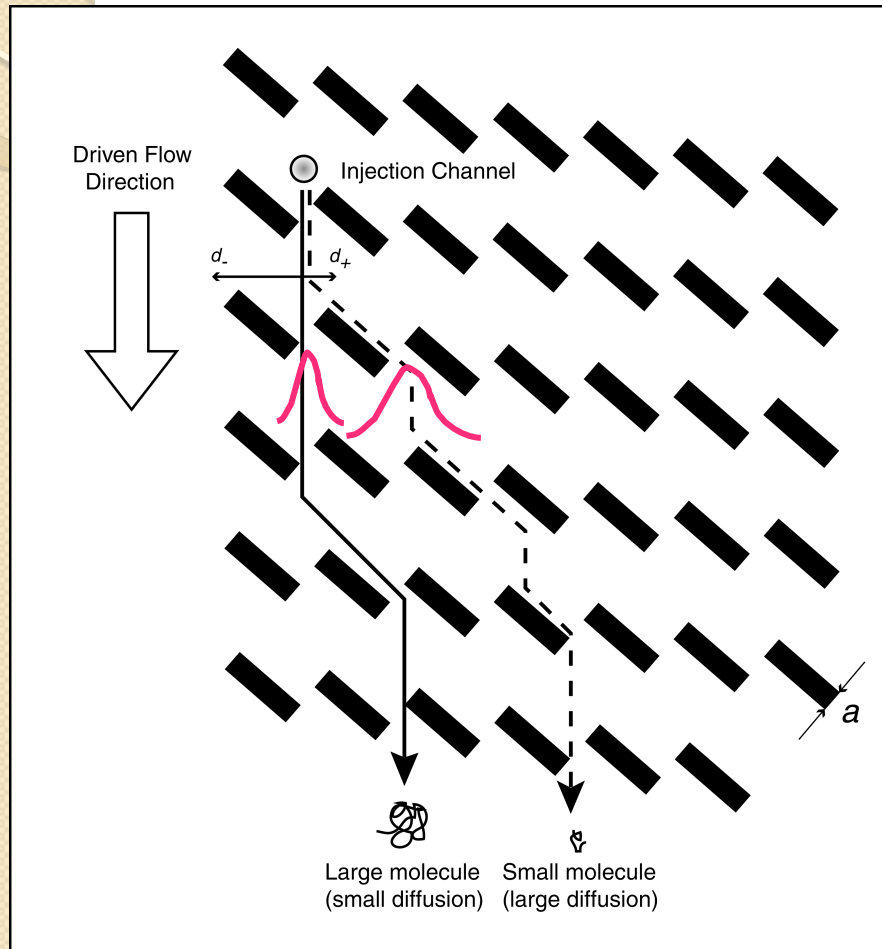
W. D. Volkmuth & R. H. Austin

Department of Physics, Princeton University, Princeton,
New Jersey 08544, USA





Separation based on Rectified Brownian Motion

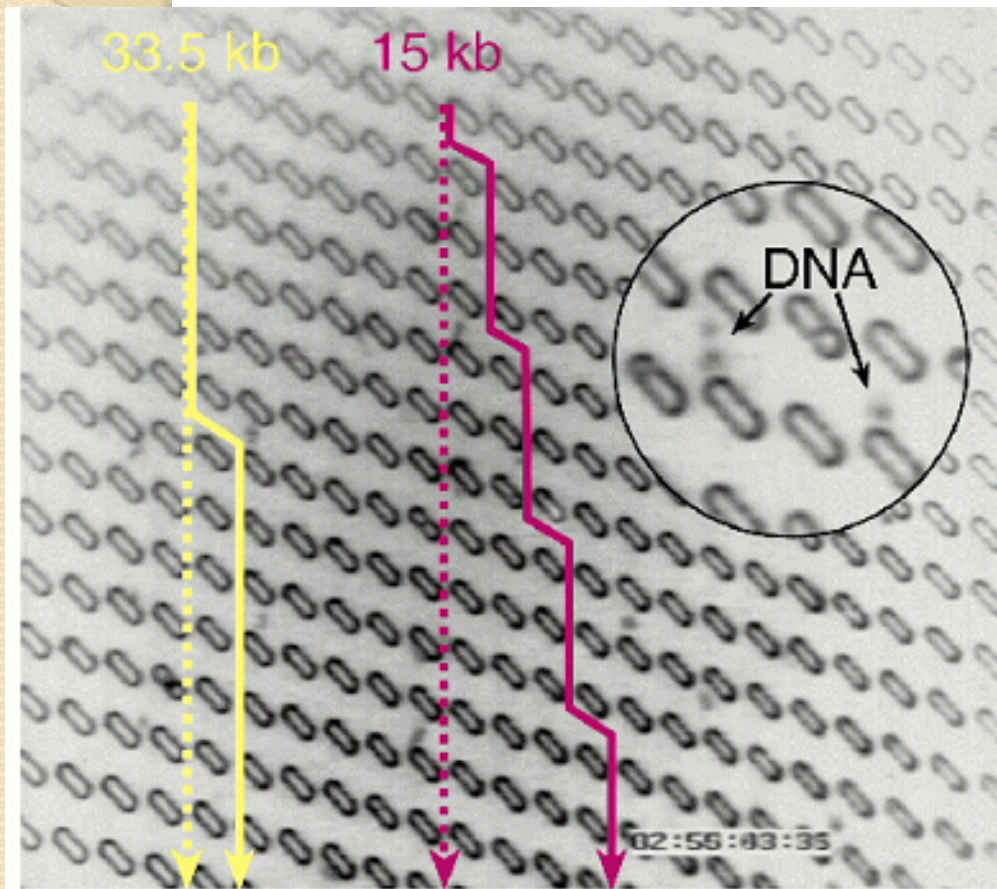


A simple idea: let the altered probabilities effectively move molecules to one side
— Rectified Brownian motion

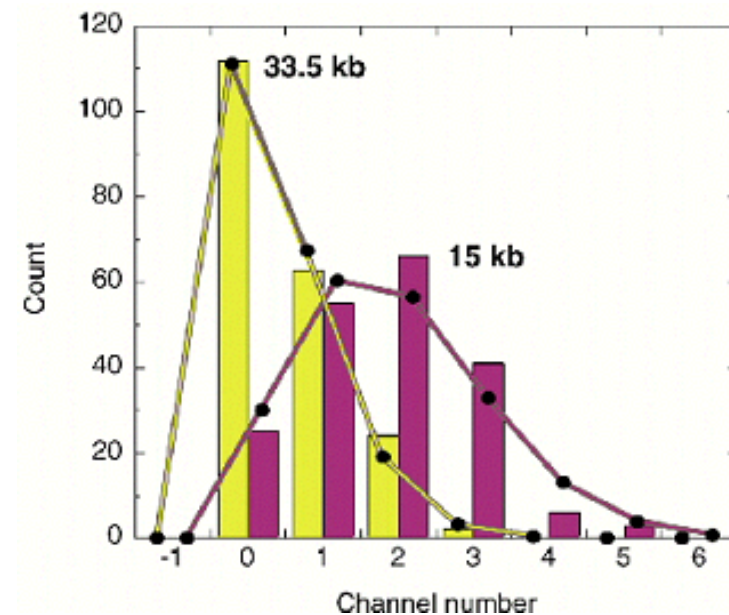
Duke and Austin, *PRL* (1998)
Ertas, *PRL* (1998)

Continuous molecular sorting by rectified Brownian motion

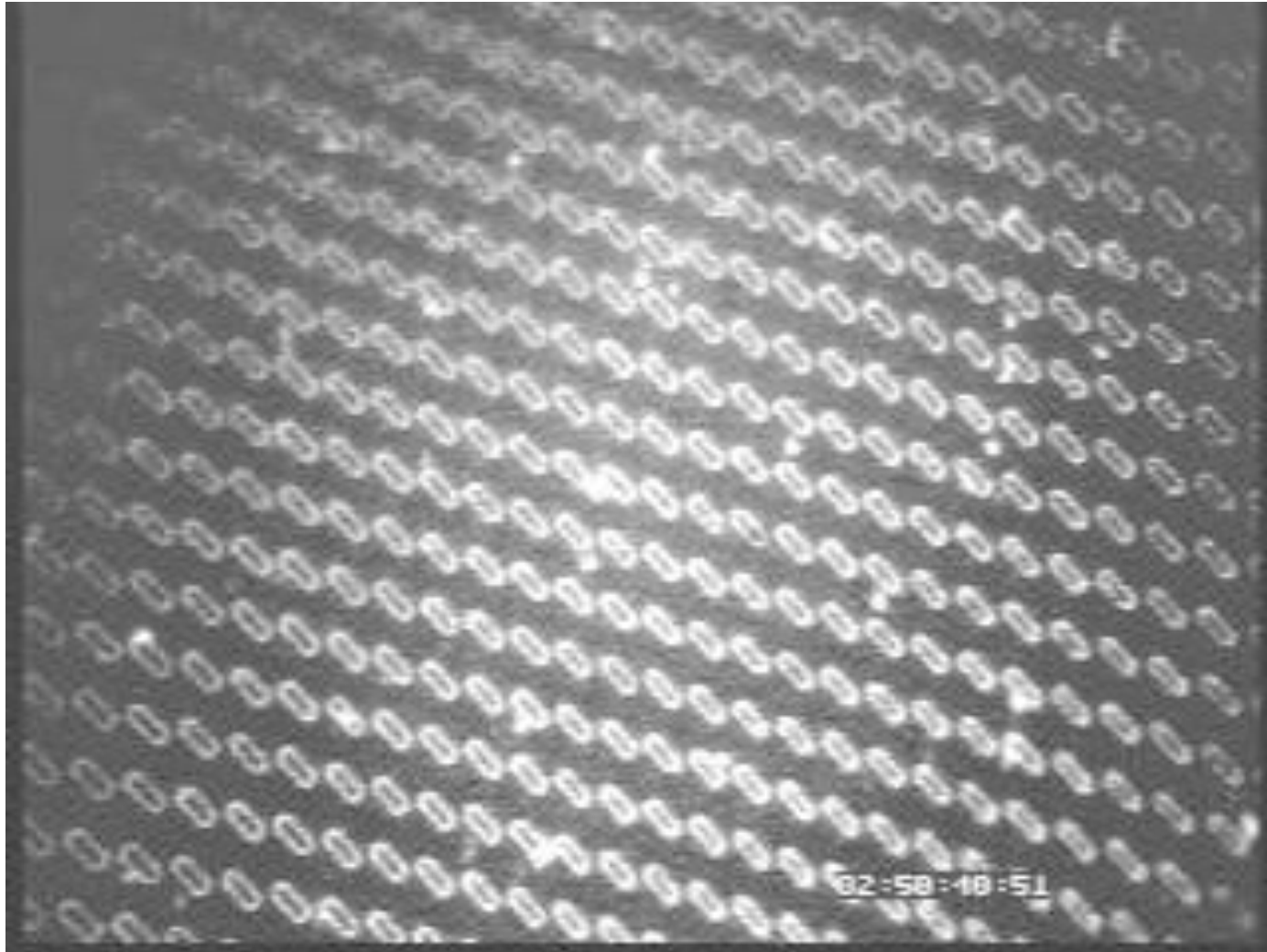
Continuous Lateral Separation



Chou et al, *PNAS* (1999)



Continuous DNA sorting by rectified Brownian motion



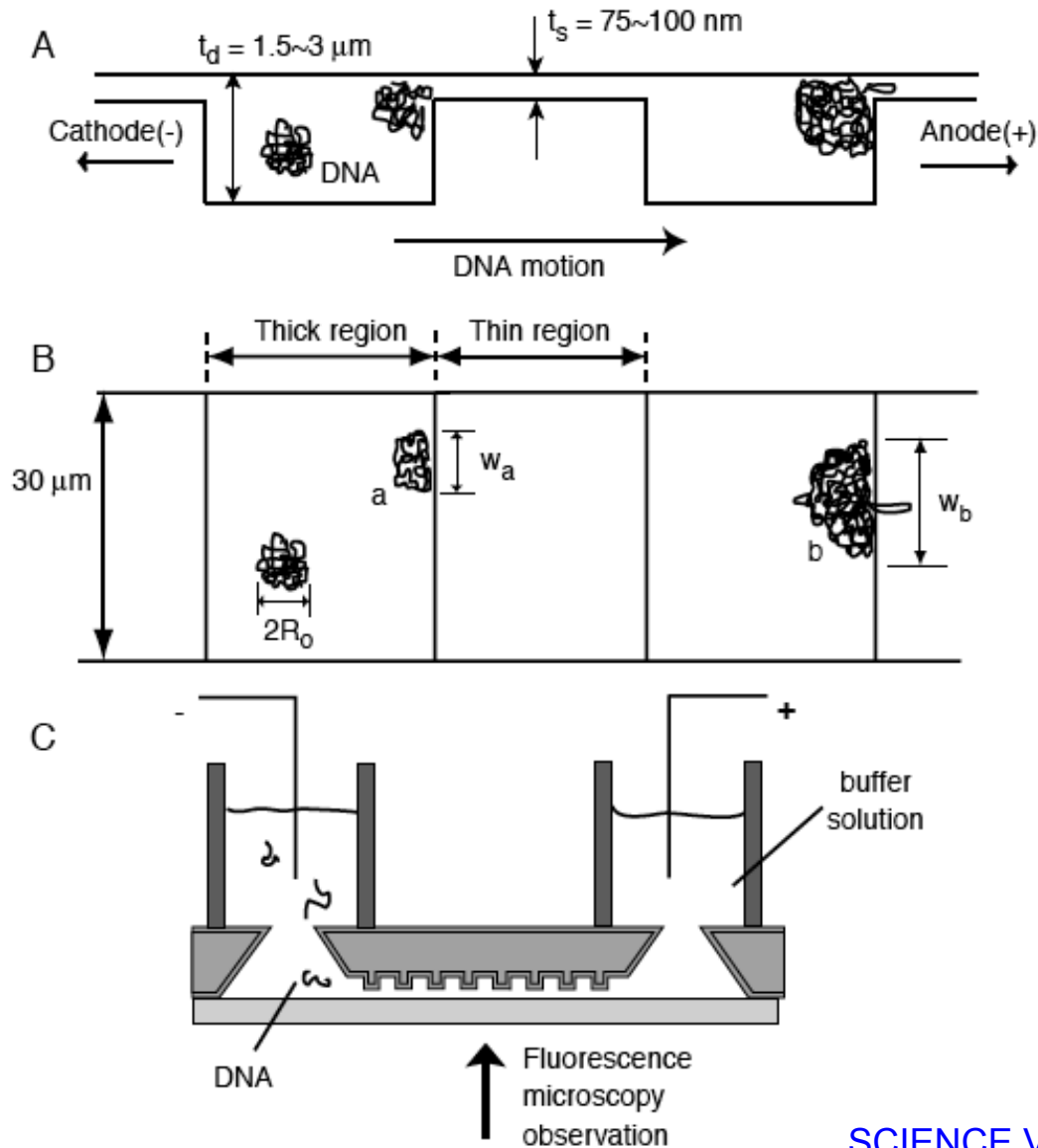


Separation of Long DNA Molecules in a Microfabricated Entropic Trap Array

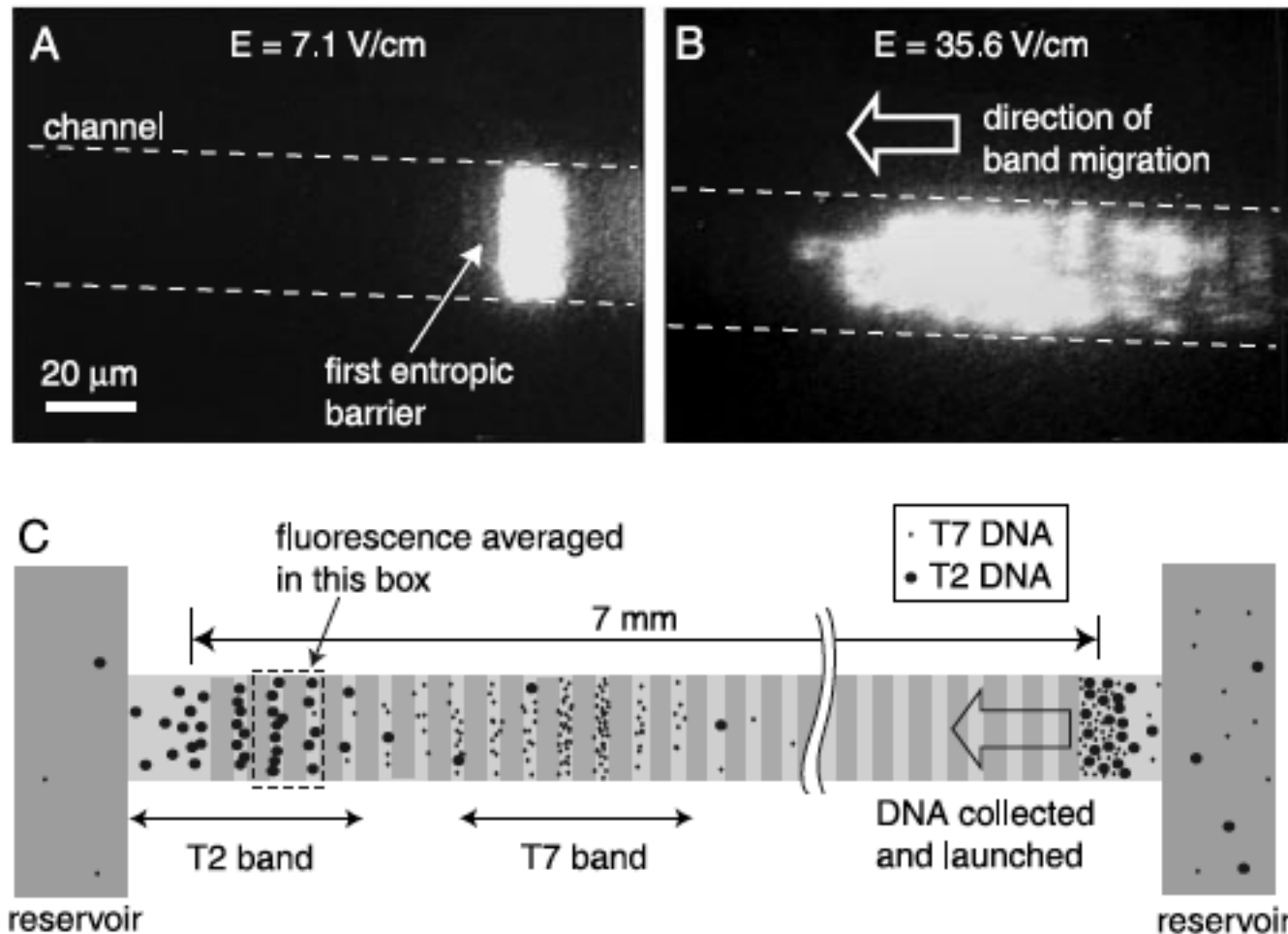
J. Han and H. G. Craighead*

A nanofluidic channel device, consisting of many entropic traps, was designed and fabricated for the separation of long DNA molecules. The channel comprises narrow constrictions and wider regions that cause size-dependent trapping of DNA at the onset of a constriction. This process creates electrophoretic mobility differences, thus enabling efficient separation without the use of a gel matrix or pulsed electric fields. Samples of long DNA molecules (5000 to $\sim 160,000$ base pairs) were efficiently separated into bands in 15-millimeter-long channels. Multiple-channel devices operating in parallel were demonstrated. The efficiency, compactness, and ease of fabrication of the device suggest the possibility of more practical integrated DNA analysis systems.

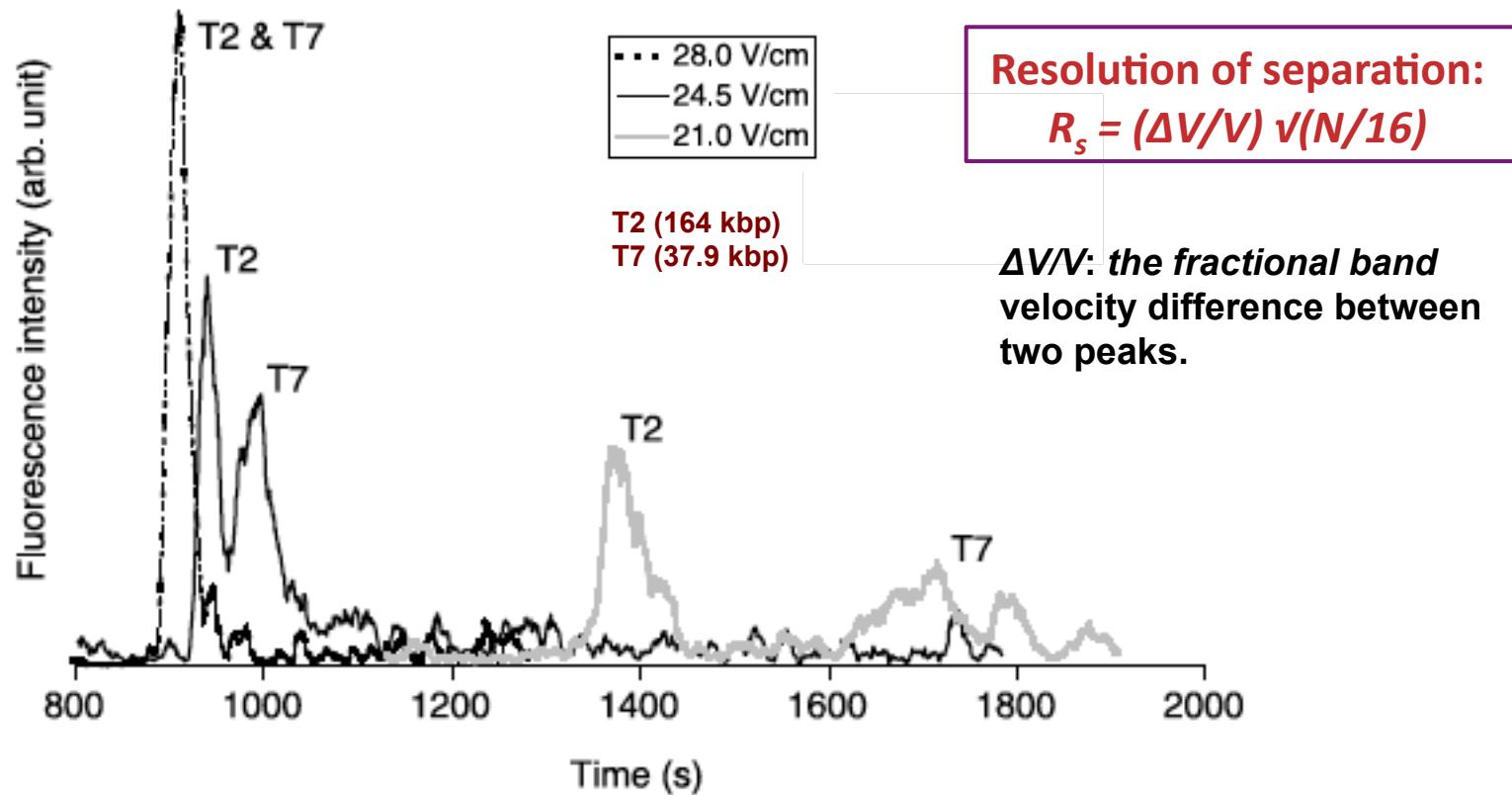
Nanofluidic DNA separation device with entropic trapping array



Band launching and analysis of bands



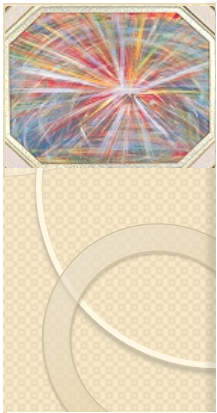
Electric field dependence of the separation of T2-T7 DNA mixture



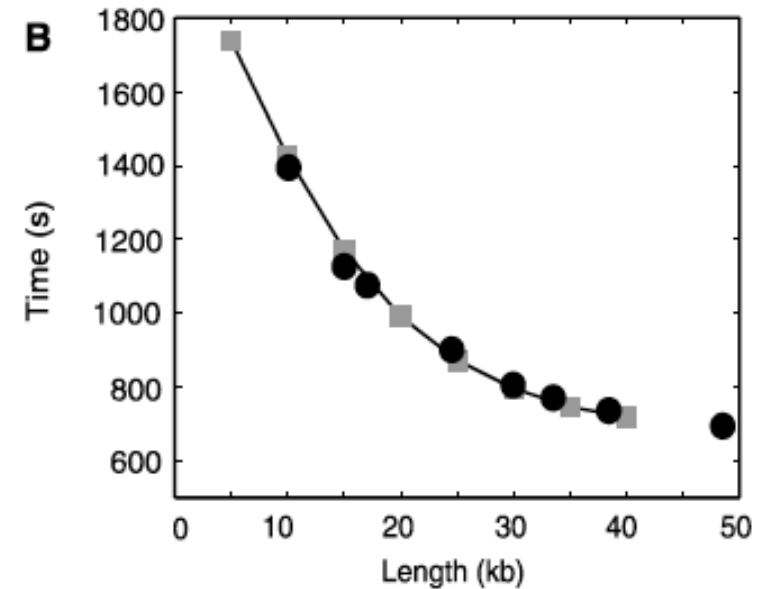
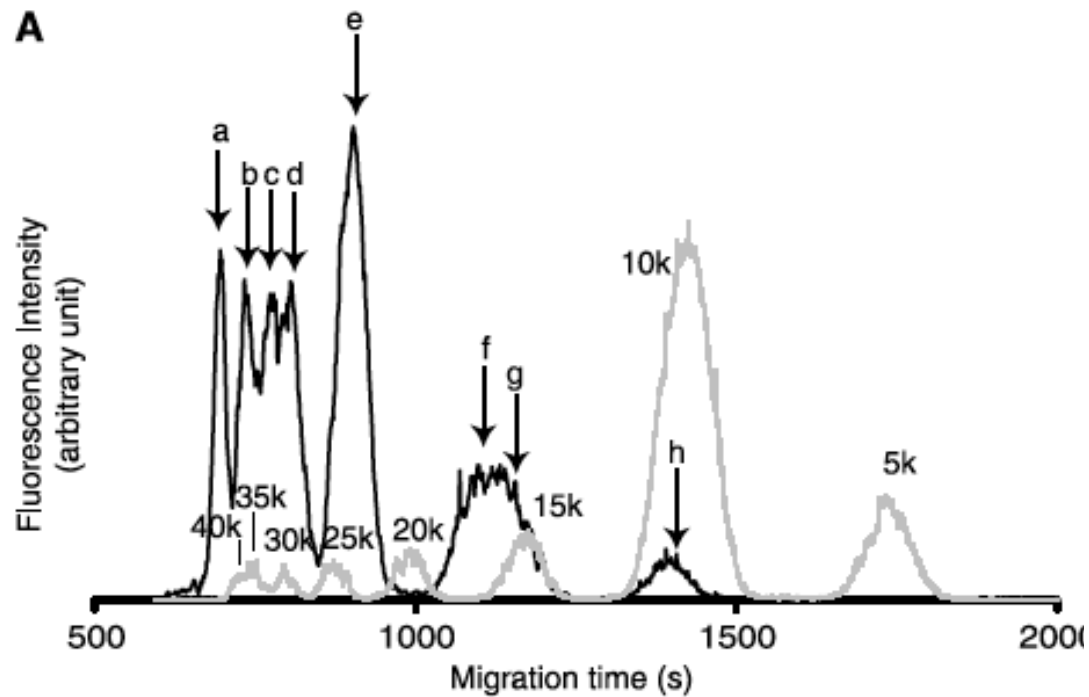
At 21.0 V/cm (gray line), the **theoretical plate number (N)** was 4900 for the T2 peak and 970 for the T7 peak, and **the resolution** was 1.95.

At 24.5 V/cm (black line), $N = 8500$ for the T2 peak and 3400 for the T7 peak, and the resolution was 0.89.

At 28.0 V/cm, no separation was achieved (broken line).



Simultaneous separation of the Mono Cut Mix sample (black line) and 5-kbp ladder sample (gray line) by the entropic trap array, run at 80 V/cm.



Large molecules move faster!!

On-chip pulsed-field electrophoresis

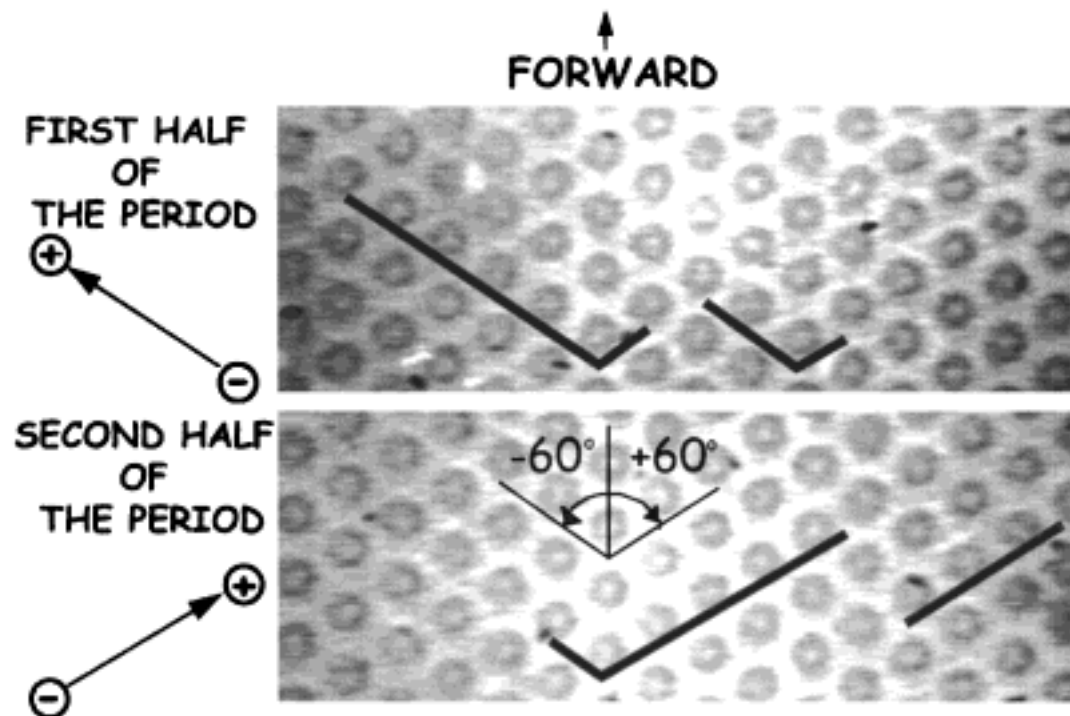
Separation of 100-Kilobase DNA Molecules in 10 Seconds

Anal. Chem. 2001, 73, 6053-6056

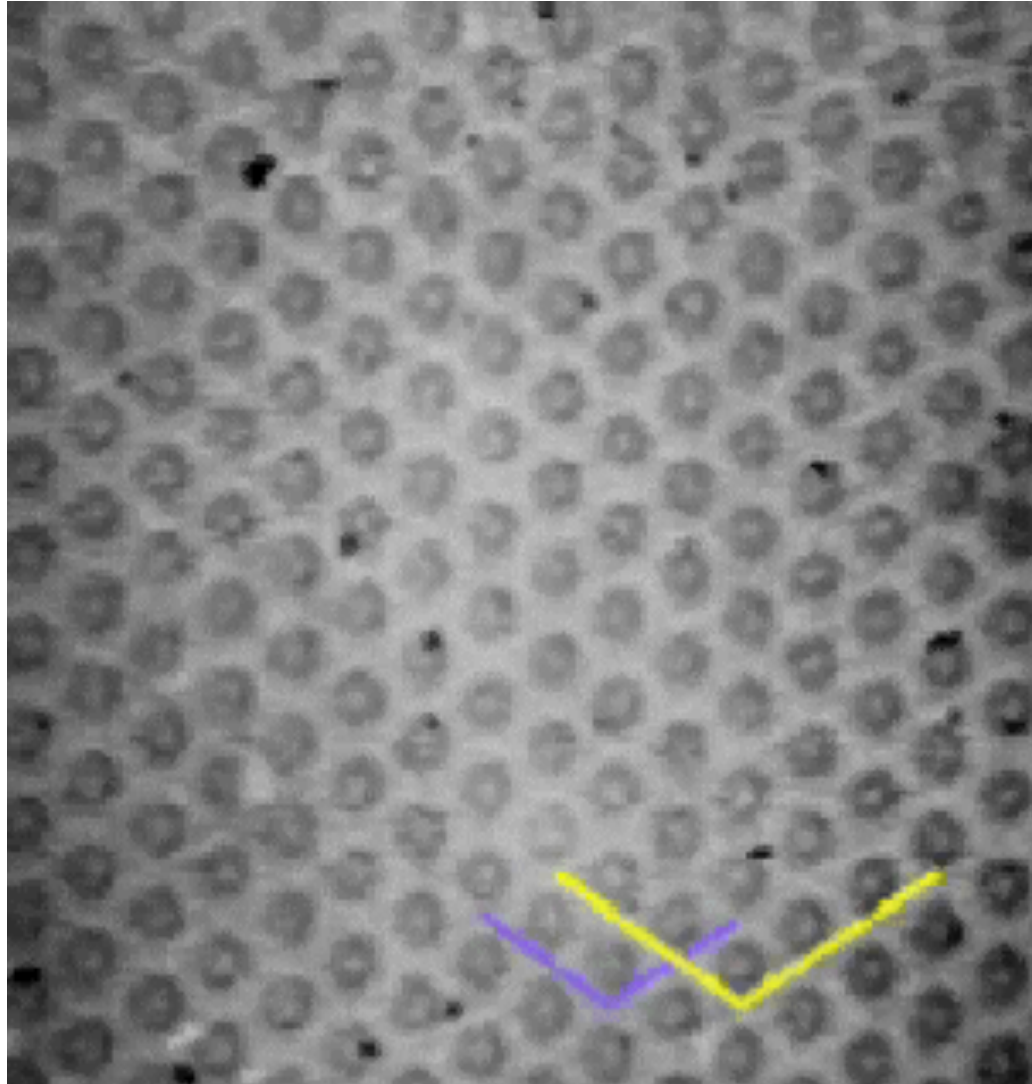
Olgica Bakajin,^{†,‡} Thomas A. J. Duke,[‡] Jonas Tegenfeldt,[†] Chia-Fu Chou,[†] Shirley S. Chan,[†] Robert H. Austin,^{*,†} and Edward C. Cox[§]

Physics Department, Princeton University, Princeton, New Jersey 08544, Cavendish Laboratory, Madingley Road, Cambridge CB3 0HE, U.K., and Department of Molecular Biology, Princeton University, Princeton, New Jersey 08544

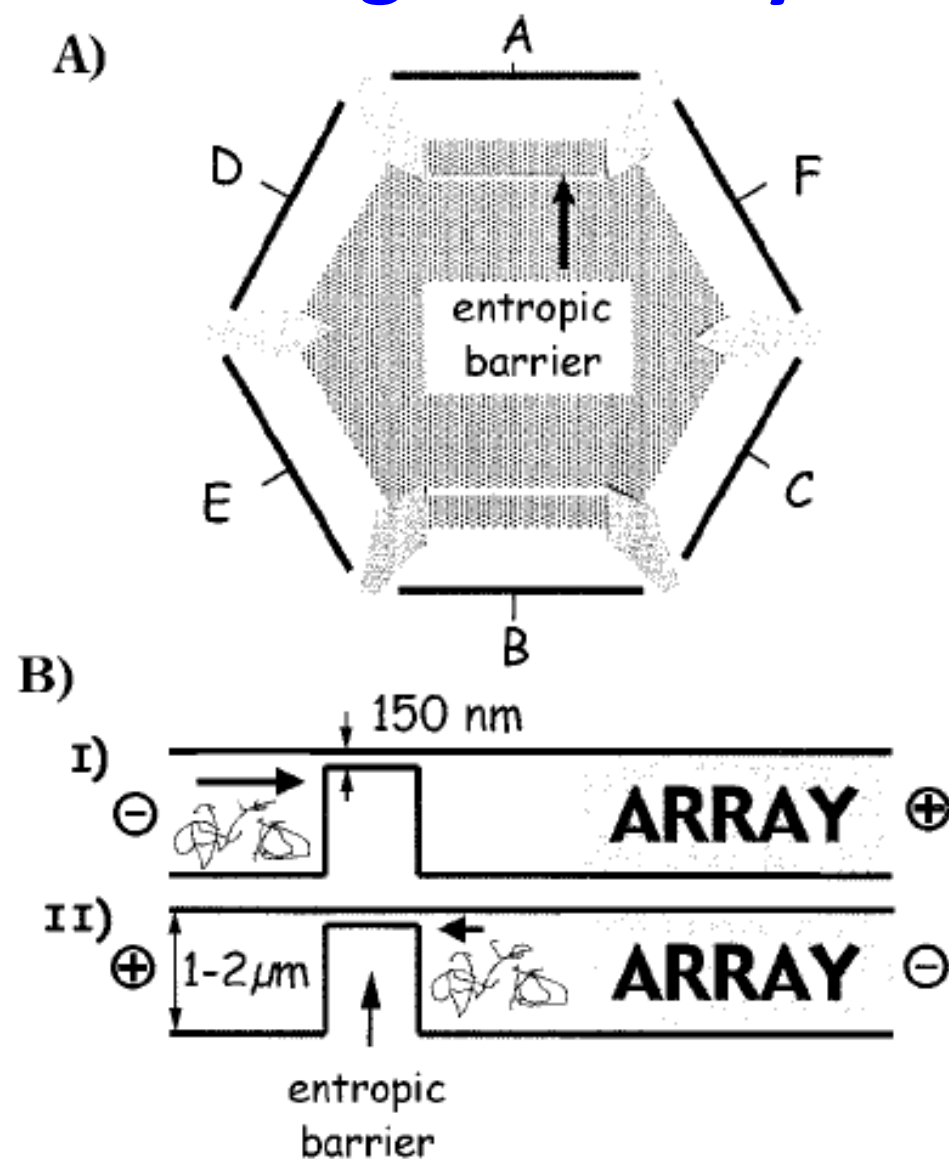
Microscopic view of the pulsed hexagonal array



On-chip pulsed-field electrophoresis (animation)



Device configuration of a pulsed hexagonal array





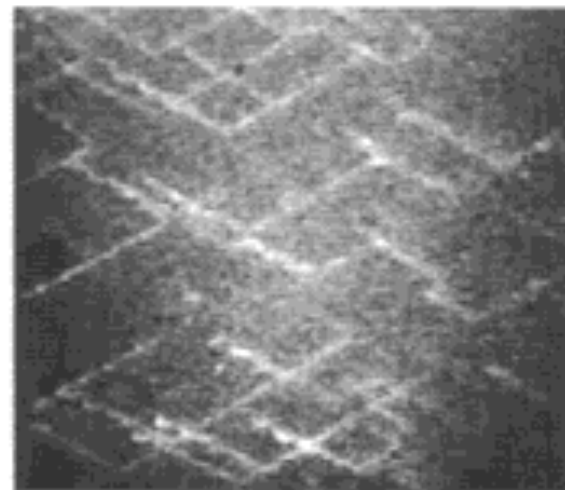
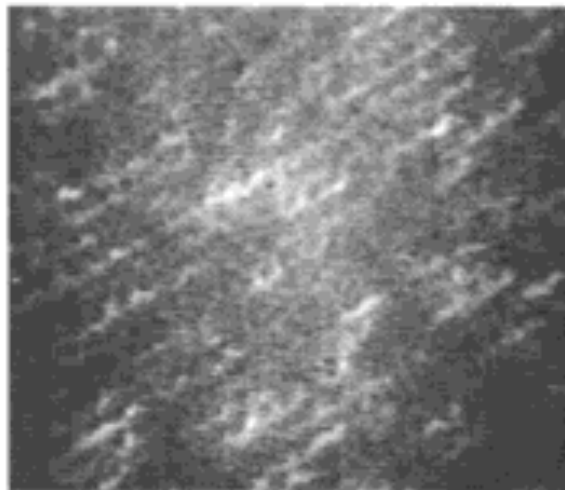
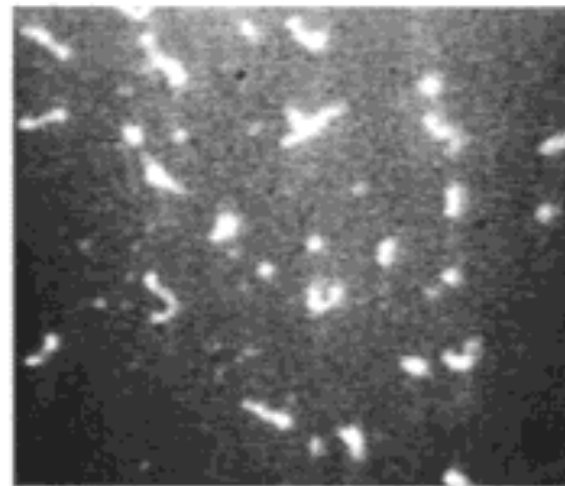
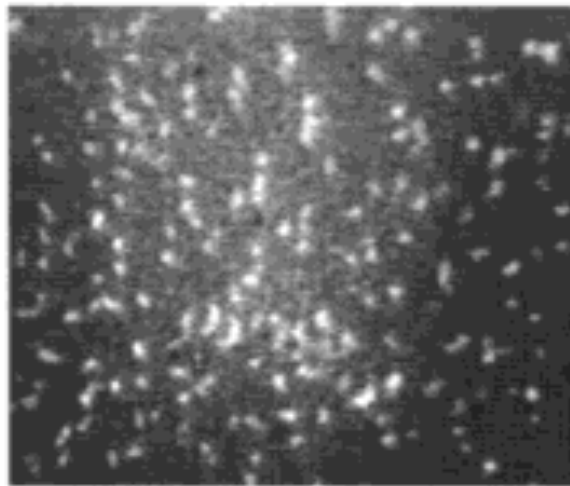
T4 and λ SEPARATION IN HEXAGONAL ARRAYS

12-24-16-43

Single-molecule images

λ : 48.5 kbp

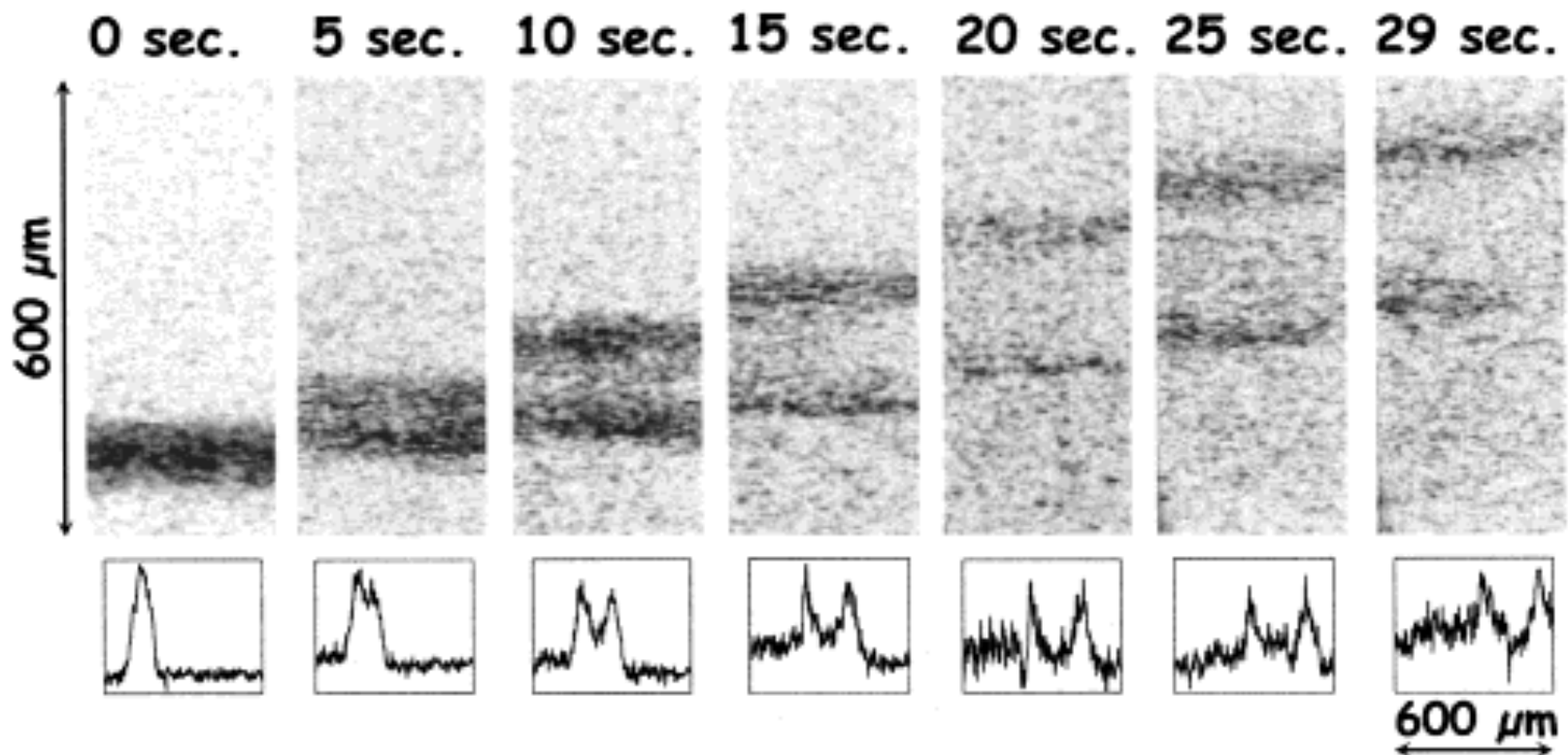
T4 : 168.9 kbp



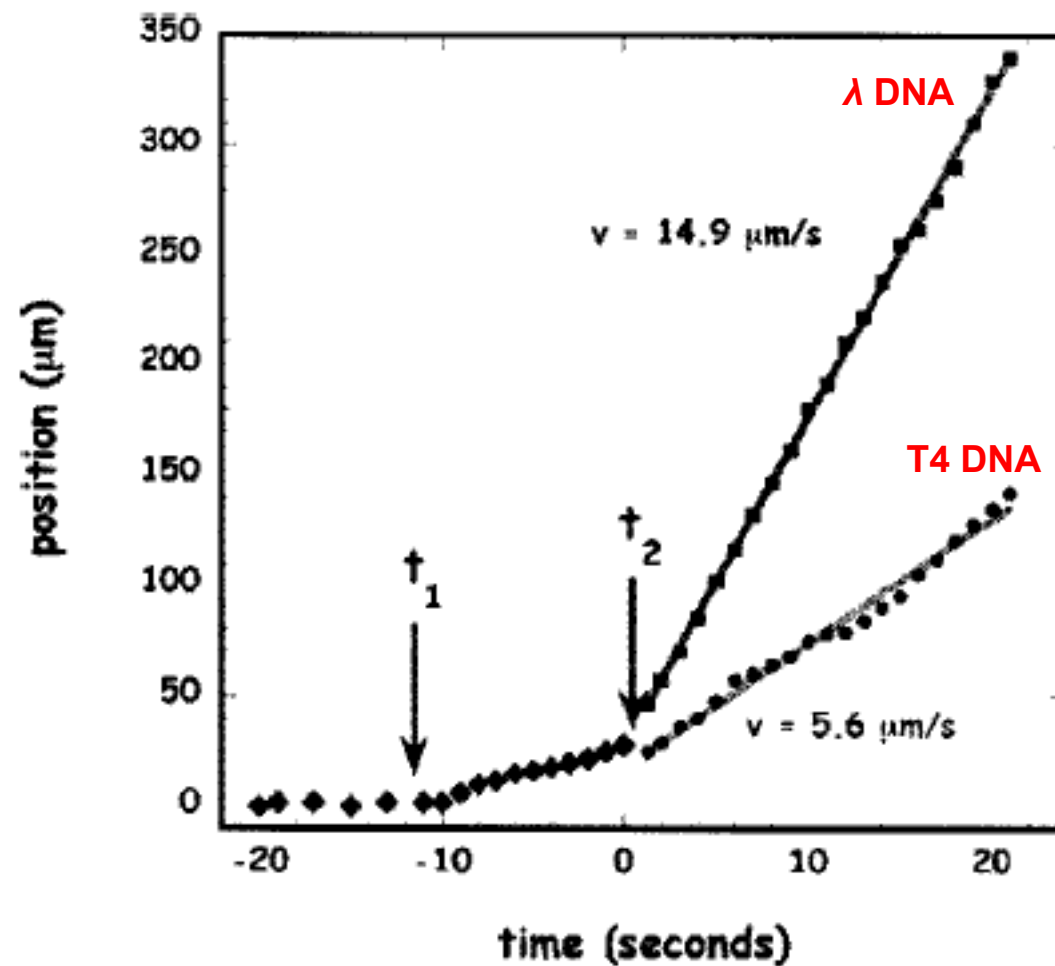
30 μ m

λ and T4 DNA separation

λ and T4 DNA pulsed at 244 V and with period $T = 1$ s after release from the entropic trap



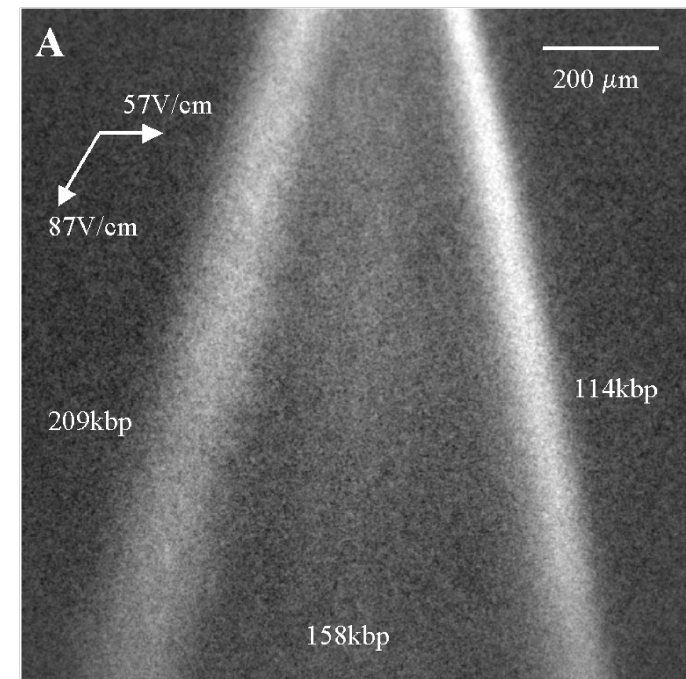
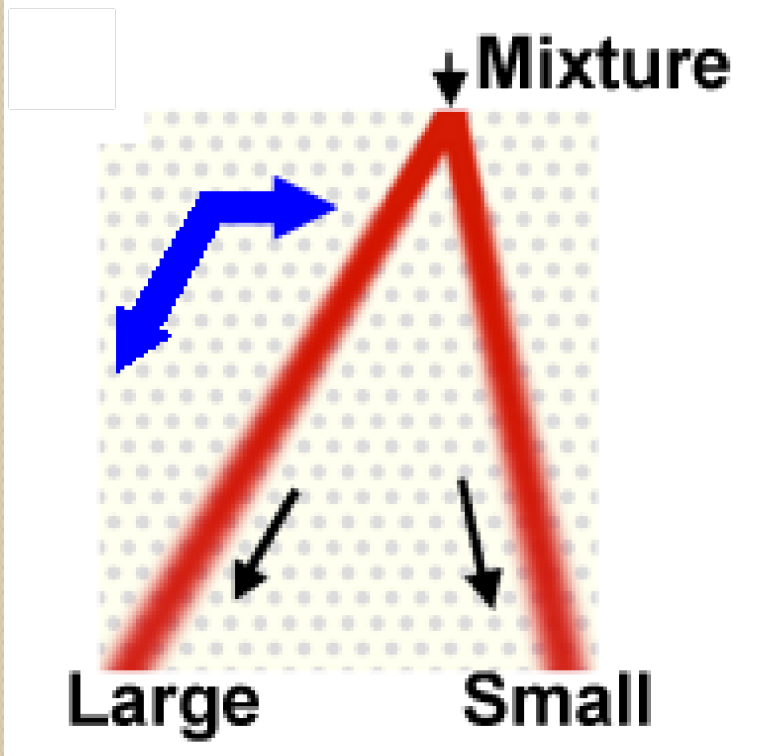
DNA position as a function of time



DNA Prism

Asymmetric pulsed fields:

The direction of the net flow motion of the DNA is angle-dependent.



Continuous separation of BAC/PAC mixture of 114, 158, and 209 kbp fragments (Huang *et al.*, Nat. Biotech. 2001)