

Capillary Electrophoresis in DNA Analysis

Intro to CE and ABI 310

DNA Academy Workshop
 Albany, NY
 June 13-14, 2005
 Dr. John M. Butler
 Dr. Bruce R. McCord



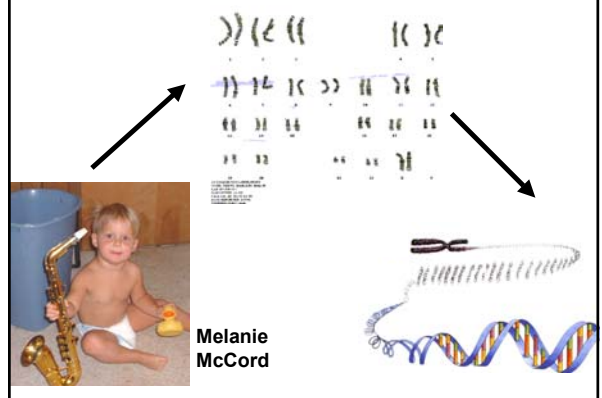
Outline for Workshop

- Introductions
- STR Analysis
- Introduction to CE and ABI 310
- Validation and Interlaboratory Studies
- Real-time qPCR and miniSTRs
- Stats and Higher Throughput Approaches
- Y-Chromosome Analysis
- Troubleshooting the ABI 310
- Review and Test

Historical Perspective

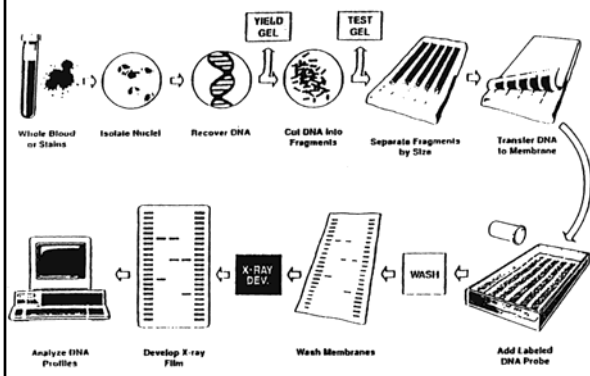
- 1928 - Griffith demonstrates DNA carries genetic information
- 1953 - Structure of DNA determined by Watson and Crick
- 1959 - Hjerten describes electrophoresis in tubes
- 1970 - Restriction enzymes begin to be used to cut DNA
- 1981 - Jorgenson performs CZE
- 1984 - Alec Jeffries utilizes restriction fragment digestion of DNA
- 1985 - Kary Mullis describes PCR

There are 23 pairs of Chromosomes



Melanie McCord

DNA Analysis by RFLP



From the Inventor of PCR

"I think I might have been stupid, in some respects, if it weren't for my psychedelic experiences."

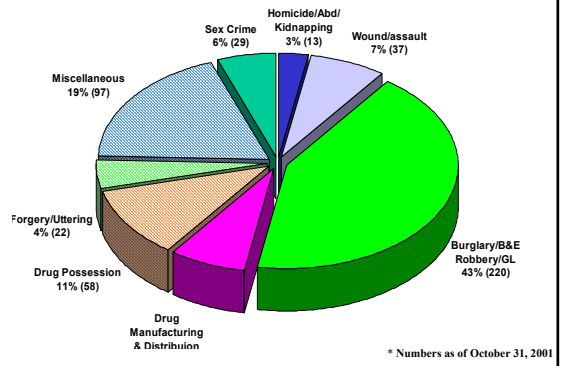
"Out of a natural laziness, I always start with the easiest possible protocol and work from there. Better yet, I suggest that someone start from there, and I come back in a month to see how things worked out."

- Kary Mullis, Ph.D., Nobel Prize Laureate, Chemistry, 1993

Issues Being Faced in the Mid-1990s

1. RFLP techniques were time consuming and insensitive
2. PCR is rapid and highly sensitive
3. The old results don't translate
4. Soon all sexual offenders (and other felons) would be required to submit a sample for testing
5. CODIS -Combined DNA Indexing System
6. How will all these samples be run ?

Using CODIS Virginia has 788 had cold 788 hits with 180,000 samples analyzed – Prior conviction:



GENE JEWELS

Scientist Hopes to Fashion a Fortune From Celebrity DNA

The Application

Speed and detection capabilities of DNA analyses have improved since the development of the PCR

Increase in number of complex assays necessitates automated testing procedures

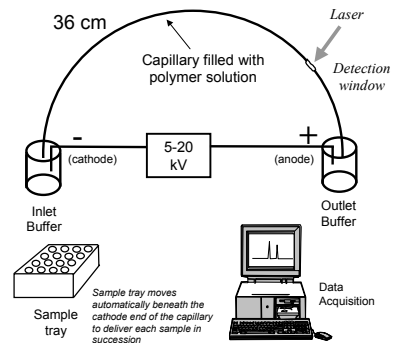
Automated systems are needed to increased sample throughput

Automated systems must be robust and must demonstrate long term stability

Why Use CE for DNA Analysis?

1. Injection, separation, and detection are automated.
2. Rapid separations are possible
3. Excellent sensitivity and resolution
4. The time at which any band elutes is precisely determined
5. Peak information is automatically stored for easy retrieval

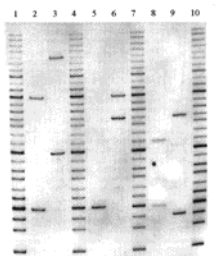
Components of a Capillary Electrophoresis System



Butler, J.M. (2001) Forensic DNA Typing, Figure 9.3, ©Academic Press

This DNA stuff was powerful and needed to be automated

BUT HOW TO MOVE
FROM GELS TO
CAPILLARIES?

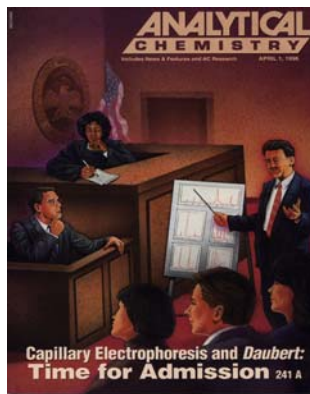


In the Early 90's the real question was how to transition from a gel to a capillary

- X-linked acrylamide gel filled capillaries tried first
 - Reusable?
 - Bubble formation
 - Thermal degradation
- Alternative was to not use a gel at all
 - Entangled polymer solutions
 - Refillable
 - Resolution?

Early Work with CE and STRs

- Barry Karger's group (1988-1990)
 - Utilized gel-filled capillaries to separate ssDNA
 - Introduced sieving polymers in the form of linear polyacrylamide to separate restriction digests
- Commercial CE systems (1992) with lasers appear (Beckman P/ACE)
- Initial STR results at FBI and AFDIL (1992-)
- First STR typing with multi-color CE (and multi-capillary) using dye-labeled primers
- ABI 310 is introduced in July 1995 as the first commercially available multi-color CE



1996

Potential of CE to revolutionize DNA analysis recognized by Analytical Chemistry

Requirements for Reliable STR Typing

- Reliable sizing over a 75-500 bp size region
- High run-to-run precision between processed samples to permit comparison of allelic ladders to sequentially processed STR samples
- Effective color separations of different dye sets used to avoid bleed through between 4 or 5 different colors
- Resolution of at least 1 bp to >350 bp to permit reliable detection of microvariant alleles

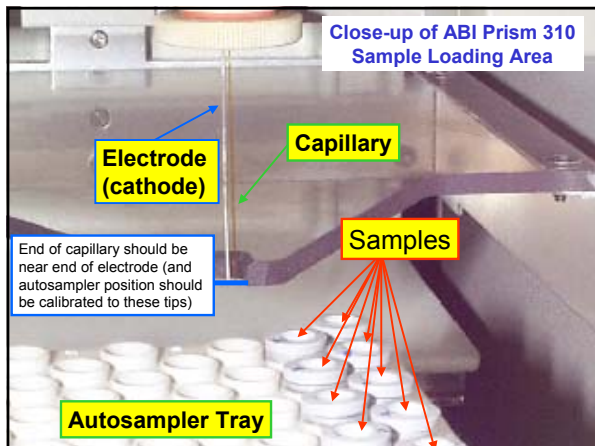
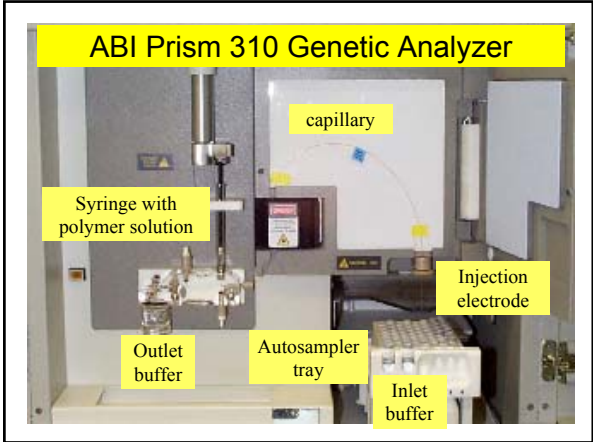
Butler *et al.* (2004) *Electrophoresis* 25: 1397-1412

McCord's Comments on this:

- I want
 - 1bp resolution from 100-350 bp
 - Resolution = $2d/(w_1 + w_2)$
 - Resolution in bp = (difference in size)/R
 - Precision of better than 0.17 bp for any given allele
 - $3 \times 0.17 = 0.51\text{bp}$ 3 standard deviations = 99.7% of data
 - Stable results from the first run to the last run
 - Calibration with one ladder reliable for 48 hours worth of runs
 - No recalibration needed

Components of ABI 310

- Chemistry
 - STR kits, fluorescent dyes, matrix samples, capillary, buffers, polymer, formamide
- Hardware
 - CCD camera, laser, electrodes, pump block, hot plate for temperature control, autosampler
- Software
 - Data collection, color separation, peak sizing & calling, genotyping, stutter removal

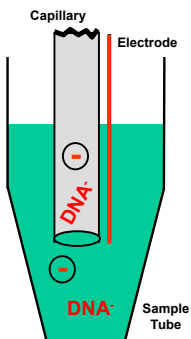


Process Involved in 310 Analysis

- Injection
 - electrokinetic injection process (formamide, water)
 - importance of sample stacking
- Separation
 - Capillary – 50um fused silica, 43 cm
 - POP-4 polymer – Polydimethyl acrylamide
 - Buffer - TAPS pH 8.0
 - Denaturants – urea, pyridinone
- Detection
 - fluorescent dyes with excitation and emission traits
 - CCD with defined virtual filters produced by assigning certain pixels

Injection

Electrokinetic Injection Process



Amount of DNA injected is inversely proportional to the ionic strength of the solution

Salty samples result in poor injections

Sample Conductivity Impacts Amount Injected

$$[DNA_{inj}] = \frac{Et(\pi r^2) (\mu_{ep} + \mu_{eof}) [DNA_{sample}] (\lambda_{buffer})}{\lambda_{sample}}$$

$[DNA_{inj}]$ is the amount of sample injected

$[DNA_{sample}]$ is the concentration of DNA in the sample

E is the electric field applied

λ_{buffer} is the buffer conductivity

t is the injection time

λ_{sample} is the sample conductivity

r is the radius of the capillary

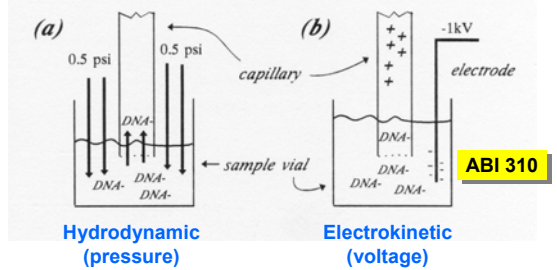
μ_{ep} is the mobility of the sample molecules

μ_{eof} is the electroosmotic mobility

Cl⁻ ions and other buffer ions present in PCR reaction contribute to the sample conductivity and thus will compete with DNA for injection onto the capillary

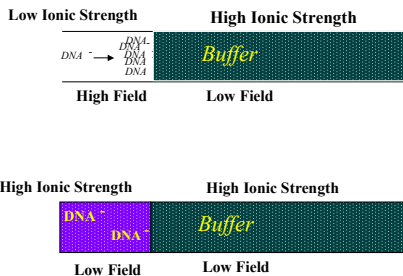
Butler et al. (2004) Electrophoresis 25: 1397-1412

CE Injection Methods



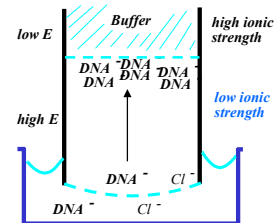
Ulfelder K. J.; McCord, B. R. Capillary Electrophoresis of DNA, In *Handbook of Capillary Electrophoresis*, Landers, J., Ed., CRC Press: NY, 1996; pp. 347-378.
Butler, J.M. (1997) Effects of sample matrix and injection on DNA separations. *Analysis of Nucleic Acids by Capillary Electrophoresis* (Heller, C., ed.), Vieweg: Germany, Chapter 5, pp. 125-134

Sample Stacking Process



Two Major Effects of Sample Stacking

1. Sample is preconcentrated. Effect is inversely proportional to ionic strength
2. Sample is focused. Ions stop moving in low electric field
3. Mobility of sample = $\mu_{ep} = \text{velocity} / \text{electric field}$



Typical Sample Preparation for ssDNA

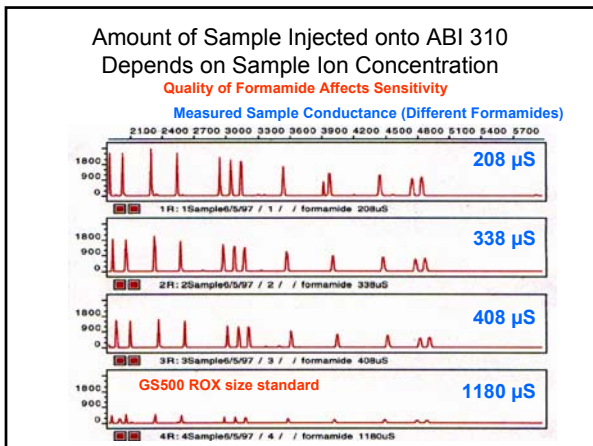
1. Perform PCR with dye-labeled primers
2. Dilute 1 μ L PCR product with 24 μ L deionized formamide; add 1 μ L ROX-labeled internal sizing standard
3. Denature 2 minutes at 95 $^{\circ}$ C with thermocycler
4. Cool to 4 $^{\circ}$ C in thermocycler or ice bath
5. Sample will remain denatured for at least 3 days

Injection Study

Evaluate the effects of sample injection on electrophoretic separations by CE.

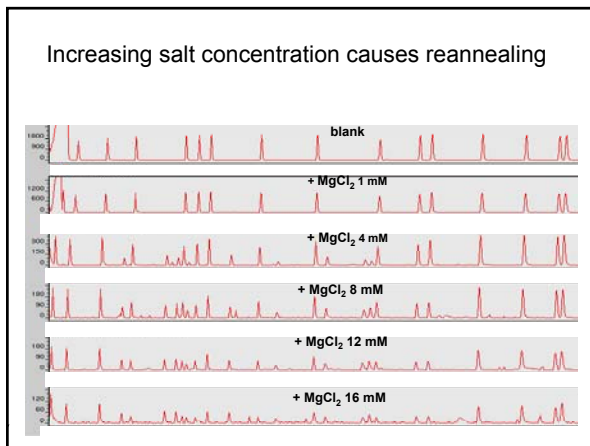
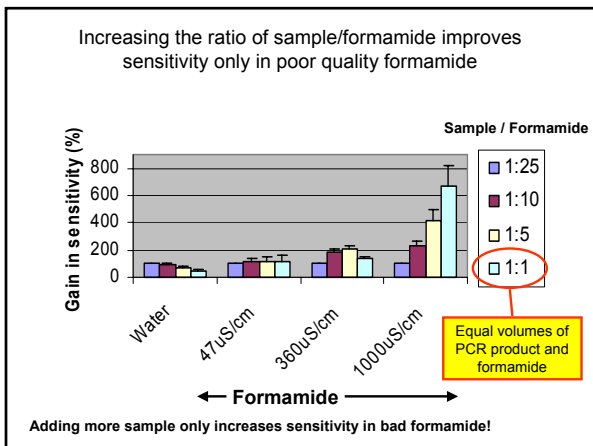
- different solvents (water and formamide of varying purity);
- different concentration of the sample;
- addition of salts;
- sample stacking

Crivellente, F.; McCord, B. R. The application of pH mediated sample stacking in the analysis of multiplexed short tandem repeats, *Journal of Capillary Electrophoresis* 2002, 7 (3-4), 73-80.



Effect of Formamide on Peak Resolution and Sensitivity (GS500 ROX Internal Standard)

Solvent	Resolution	Peak Height
Water	1.19+/- 0.01	2700+/- 300
Formamide (27μS)	1.15+/- 0.05	2960+/- 30
Formamide (360μS)	1.20+/- 0.08	879 +/- 4
Formamide (1000μS)	1.20+/- 0.06	290 +/- 14



- ### Comments on Sample Preparation
- Use high quality formamide (<100 μS/cm!)
 - ABI sells Hi-Di formamide
 - regular formamide can be made more pure with ion exchange resin
 - Deionized water vs. formamide
 - Biega and Ducean (1999) *J. Forensic Sci.* 44: 1029-1031
 - Crivellente, *Journal of Capillary Electrophoresis* 2002, 7 (3-4), 73-80.
 - water works fine but samples are not stable as long as with formamide; water also evaporates over time...
 - Denaturation with heating and snap cooling
 - use a thermal cycler for heating and cold aluminum block for snap cooling
 - heat/cool denaturation step is necessary only if water is substituted for formamide...

Separation

ELECTROPHORESIS THEORY

“Ok here’s my recipe idea called the electric pickle
 Attach the hot lead to a screw and shove it in. The neutral
 lead goes in the other end. Turn out the lights and plug it in
 It glows and sizzles. The juicy ones work best”
 www.voltnet.com/cook

$P = VI = I^2R$ pickle cooks

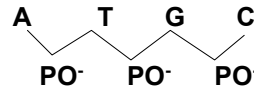
$v_{ep} = \mu_{ep}V$ Ions move through pickle faster
 at high voltage

$\mu_{ep} = q/6\pi\eta r$ small ions with high charge
 move fastest

DNA and Electrophoresis

“From a practical point of view it is disappointing that
 electrophoresis cannot be used to fractionate or analyze
 DNA’s on the basis of size” Olivera, Biopolymers 1964, 2, 245

$\mu_{ep} = q/6\pi\eta r$

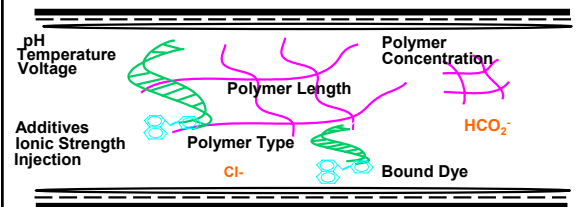


As size increases so does charge!

How to Sieve DNA in a Capillary?

- Cross linked gels
 - The familiar approach
 - High resolution, but not so reusable
- Entangled Linear polymers
 - Refillable, reusable
 - Lower Resolution
 - Osmotic Flow

Inside a CE Capillary



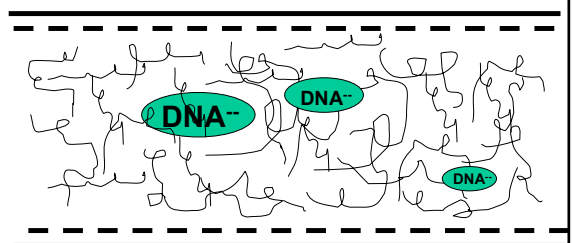
Which Factors Affect Resolution and Migration
 of DNA Fragments ?

Variables

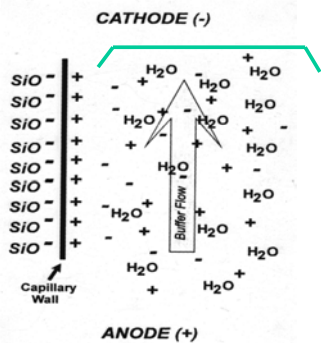
- pH – osmotic flow and DNA resolution
- Buffer concentration/ionic strength – osmotic flow and current/joule heating
- Polymer concentration – wall coating and DNA resolution
- Capillary diameter – resolution
- Polymer type- wall coating
- Field strength – resolution, run time, current, EOF
- Temperature- DNA conformation

Separation Mechanism

Electrophoretic flow →



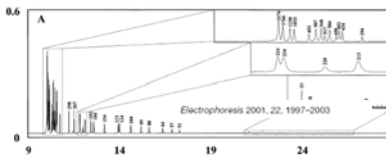
Electro Osmotic Flow



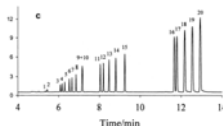
Solvated ions drag solution towards cathode in a flat flow profile

Issues with CE separations

- Big problem – electroosmotic flow

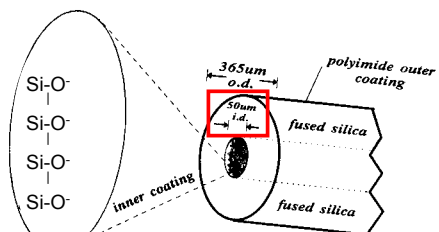


Polyethylene oxide separation of pBR 322 HAE III digest (EOF present)
Electrophoresis 2001, 22, 763-770



PDMA separation of DNA (EOF not present)
Electrophoresis 2001, 22, 1997-2003

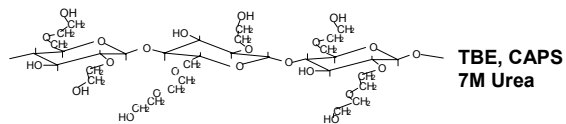
Capillary Coating



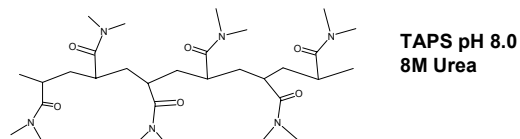
Removes effect of charged sites-eliminates EOF, sample adsorption

Dynamic coating of charged sites on fused silica capillary is accomplished with POP-4 polymer

Hydroxyethyl Cellulose (HEC)



Polydimethyl Acrylamide (POP)



So what are sieving buffers?

They are gels - very similar to polyacrylamide

They are not gels - they flow

Actually these are known as entangled linear polymers and there are many common applications



Entangled Polymer Solutions

- Polymers are not cross-linked (above entanglement threshold)
- "Gel" is not attached to the capillary wall
- Pumpable -- can be replaced after each run
- Polymer length and concentration determine the separation characteristics
- Examples:
 - 1% HEC (hydroxyethyl cellulose)
 - 4% linear polyacrylamide
 - POP-4 and POP-6

What is in POP-4 and Genetic Analyzer Buffer?

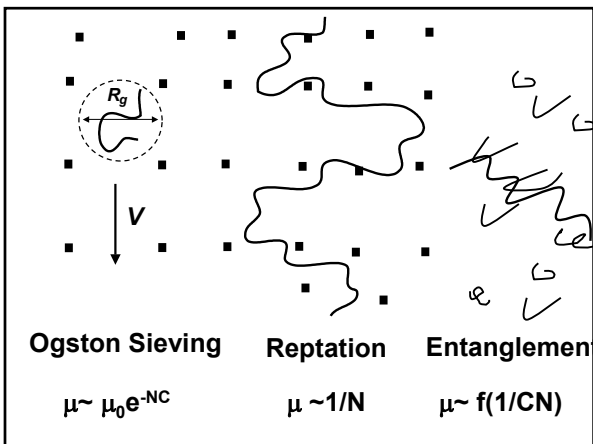
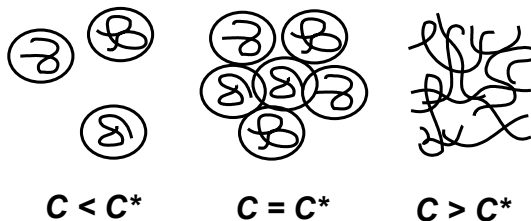
© 1997 Oxford University Press *Nucleic Acid Research*, 1997, Vol. 25, No. 19 3925-3929
Improved single-strand DNA sizing accuracy in capillary electrophoresis
 Barnett B. Rosenblum*, Frank Oaks, Steve Menchen and Ben Johnson
 PE Applied Biosystems, 850 Lincoln Centre Drive, Foster City, CA 94404, USA
 Received May 29, 1997; Revised and Accepted August 6, 1997

See also Wenz *et al.* (1998) *Genome Research* 8: 69-80

POP-4 (4% poly-dimethylacrylamide, 8 M urea, 5% 2-pyrrolidinone)
 US Patent 5,552,028 covers POP-4 synthesis

Running buffer contains 100 mM TAPS and 1 mM EDTA (adjusted to pH 8.0 with NaOH)
 TAPS = *N*-Tris-(hydroxymethyl)methyl-3-aminopropane-sulfonic acid

Transient Pores are Formed above the Entanglement Threshold

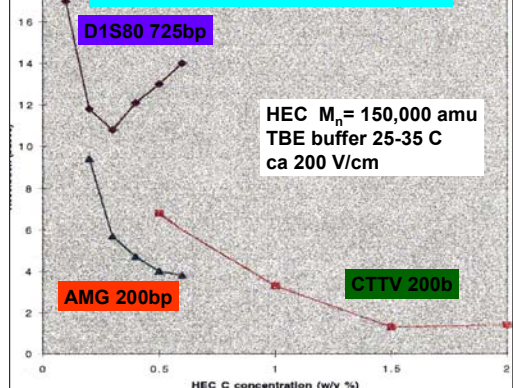


Large DNA undergoing Electrophoresis in an HEC sieving buffer (confocal microscopy)
 Oxazole yellow dimer labeled

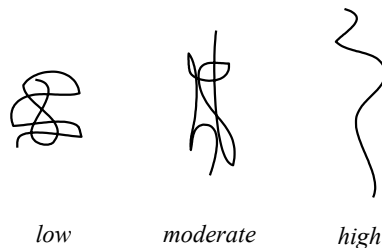


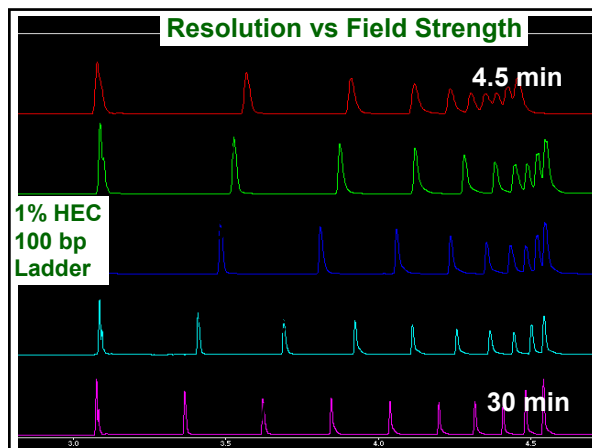
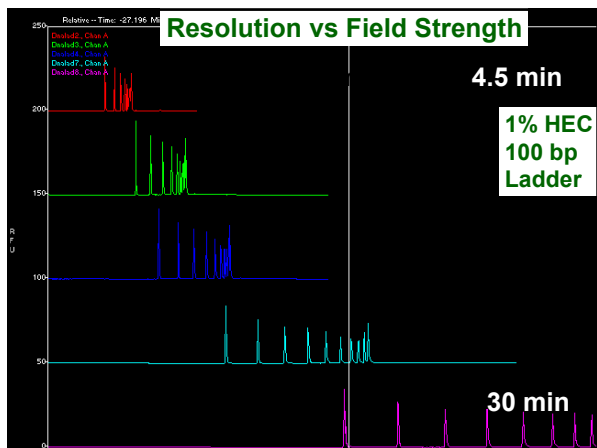
Michael Morris - U Michigan

Effect of Polymer Concentration on Resolution:



The electric field strength can influence the shape of the DNA molecule.

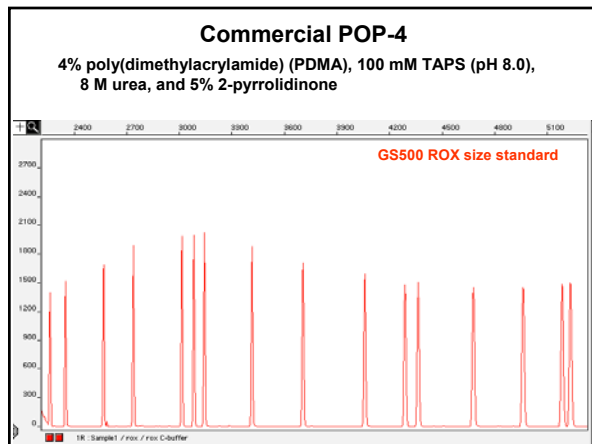




- ### The Keys
1. Polymer strand interactions create pores
 2. Average pore size~ average DNA volume
 3. Viscosity should be minimized
 4. Field strength optimized

- ### Separation Issues
- **Capillary wall coating** -- dynamic coating with polymer
 - Wall charges are masked by methyl acrylamide
 - **Electrophoresis buffer** --
 - Urea for denaturing and viscosity
 - Buffer for constant pH
 - Pyrrolidinone for denaturing
 - EDTA for stability and chelating metals
 - **Polymer solution** -- POP-4, (but others work also)
 - **Run temperature** -- 60 °C helps reduce secondary structure on DNA and improves precision. (Temperature control affects DNA sizing)

- ### How to Improve Resolution?
1. Lower Field Strength
 2. Increase Capillary Length
 3. Increase Polymer Concentration
 4. Change Polymer Length



NOTICE

US005567292A

United States Patent [19] Patent Number: **5,567,292**
Madabhushi et al. [45] Date of Patent: **Oct. 22, 1996**

[54] **POLYMERS FOR SEPARATION OF BIOMOLECULES BY CAPILLARY ELECTROPHORESIS**
 [75] Inventors: Ramakrishna S. Madabhushi, Foster City; Steven M. Meechen, Fremont; J. William Efevitch, San Mateo; Paul D. Grossman, Burlingame, all of Calif.
 [73] Assignee: The Perkin-Elmer Corporation, Foster City, Calif.
 [21] Appl. No.: 350,852
 [22] Filed: Dec. 6, 1994

Related U.S. Application Data

[63] Continuation-in-part of Ser. No. 170,078, Dec. 17, 1993, abandoned.
 [51] Int. Cl.⁶ C07K 1/36; C25B 7/00; C25B 15/00
 [52] U.S. Cl. 204/481; 204/454
 [58] Field of Search 204/180.1; 182.8; 204/299 R; 252/315.1

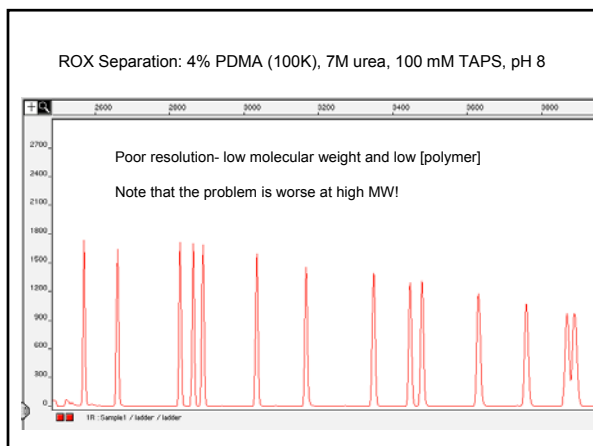
Smith et al. Methods in Enzymology vol. 155 pp. 260-301, Academic Press (1991). Methods in Enzymology 155:260-301, Academic Press (1991) (no month) The synthesis and use of fluorescent oligonucleotides in DNA sequence analysis.
 Grant et al., Chemical Dictionary, 5th ed., p. 122.
 Primary Examiner—John Niebling
 Assistant Examiner—Edna Wong
 Attorney, Agent, or Firm—Paul D. Grossman

[57] **ABSTRACT**
 The invention provides uncharged water-soluble silica-adsorbing polymers for suppressing electroosmotic flow and to reduce analyte-wall interactions in capillary electrophoresis. In one aspect of the invention, one or more of such polymers are employed as components of a separation medium for the separation of biomolecules, such as polynucleotides, polyaaromatics, proteins, and the like, by capillary electrophoresis. Generally, such polymers are characterized by (i) water solubility over the temperature range between about 20° C. to about 50° C., (ii) concentration in a separation medium in the range between about 0.001% to

Synthesis of PDMA (molecular mass ~100K)

- N,N- dimethylacrylamide (DMA) distilled under vacuum, middle portion of distillate collected
- 10 ml t-butanol used as a solvent
- 70% w/v monomer, 5 mmol/L 2,2'-azobisisobutyronitrile (AIBN)
- N₂ bubbled through solution for 10 min, RT
- Reaction performed at 55°C for 15 min
- Final product diluted by methylene chloride, precipitated by hexane and rotovapped to dryness

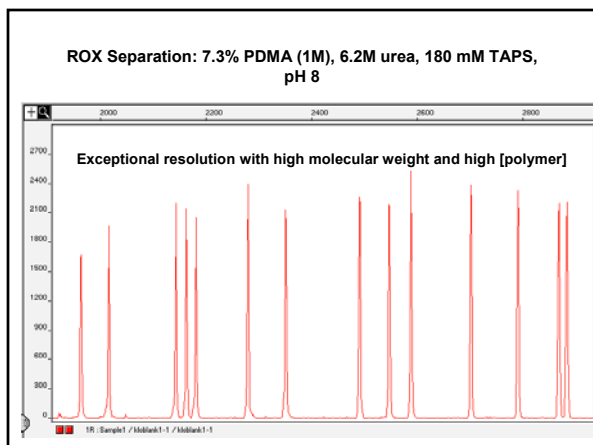
Liang, D. et al. *Electrophoresis* 2001, 22, 1997-2003.



Second Synthesis Procedure for PDMA (~1M)

- DMA distilled
- Added 16.3 ml of methanol to 46.3 ml dH₂O
- Added 6.3 g of DMA to mixture
- N₂ bubbled through for 1 h (covered flask to prevent excess methanol evaporation)
- Added 0.3 ml of ammonium persulfate stock solution (made by dissolving 0.2 g of APS in 1.8 ml of dH₂O) to the methanol/ H₂O mixture

Madabhushi, R.S. DNA Sequencing in Noncovalently Coated Capillaries Using Low Viscosity Polymer Solutions. In *Methods in Molecular Biology* 2001, Vol. 163.



DNA Separation Conclusions

- DNA molecules interact with entangled polymers – friction, sieving, entanglement
- Polymers are **not cross-linked** (as in slab gels)
- “Gel” is **not attached** to the capillary wall – not really a gel
- **Pumpable** -- can be replaced after each run
- Polymer length and concentration determine the separation characteristics – **too high and the capillary cant be refilled**

Section II: Issues

Precision and Resolution

How to Optimize and Maintain

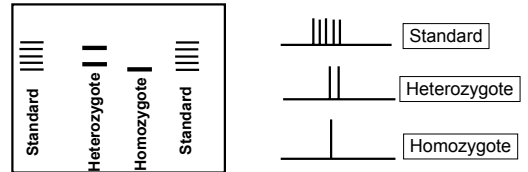
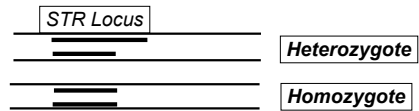
Stability

How to Characterize

Temperature

A Critical Parameter

Estimating Size



Gel Analysis

CE Analysis

Linearity of DNA Migration Times

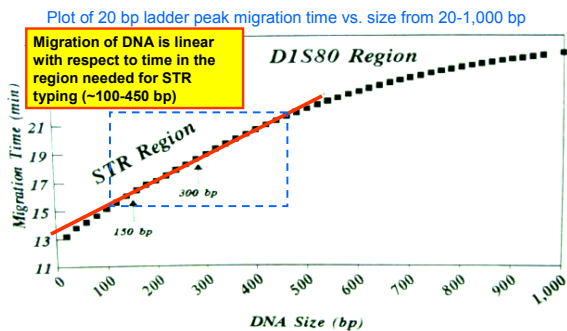


Figure 5-17 from John M. Butler Ph.D. dissertation (Aug 1995) "Sizing and quantitation of polymerase chain reaction products by capillary electrophoresis for use in DNA typing"

Use of Single Color and Dual Internal Size Standards for STR Typing

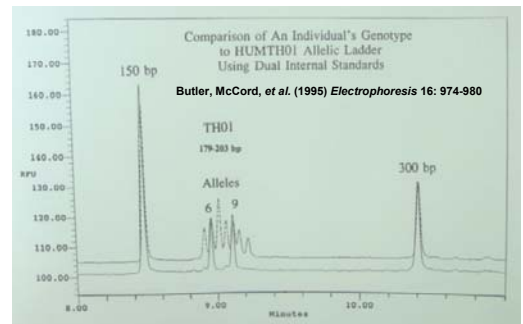
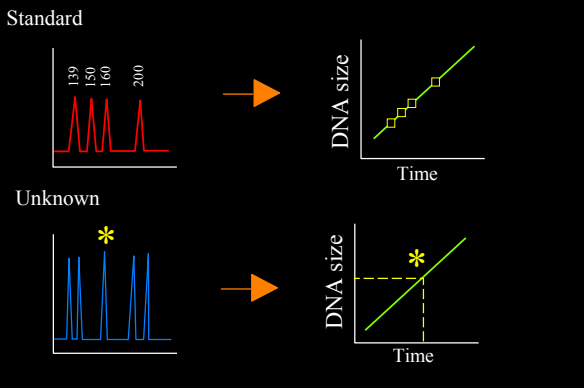


Figure 5-23 from John M. Butler Ph.D. dissertation (Aug 1995) "Sizing and quantitation of polymerase chain reaction products by capillary electrophoresis for use in DNA typing"

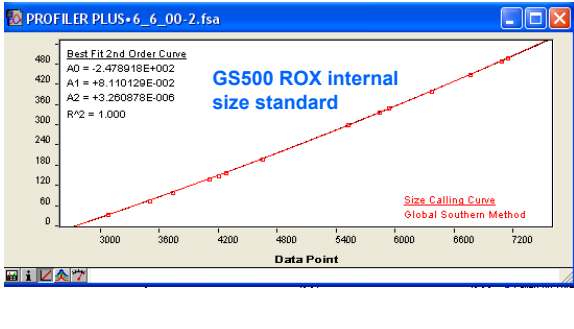
Sizing DNA Fragments using dye labeled DNA



DNA Size Estimation with ABI 310

1. Each sample is run with a ROX internal standard
2. An external standard is run with ROX as well
3. The unknown allele sequence is determined by comparison to the known ladder allele
4. Assumptions?

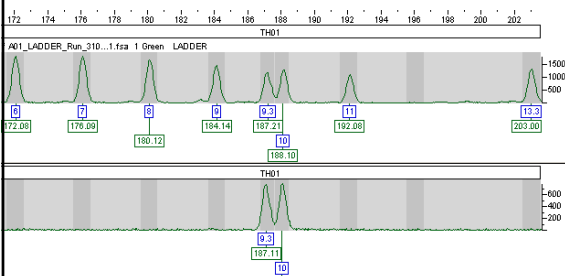
Note the exceptional linearity between 50-500 bp



Assumptions with ABI 310 Method

1. DNA is a sphere. (it is not)
2. The conditions for unknown run are the same as the ladder run. (they are not)
3. The ROX dye migrates relatively the same as the FAM dye. (It does not)
4. A calibration for one ladder is good for an entire run (sometimes)
5. Temperature is constant (to what precision?)

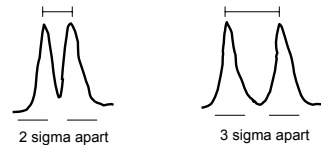
Single Base? Resolution with STRs? How is it defined?



NIST ABI 3100 with POP-6 and 36 cm array
Pop 6 is generally required for highest resolution
Higher concentration of polymer but difficult to pump with a 310

There is a definition for resolution

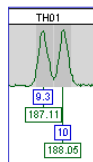
- $R = (t_2 - t_1) / [1/2(w_1 + w_2)]$
- At a resolution of 1 peaks are 2 standard deviations apart
- Baseline resolution occurs at an R value of 1.5 this is 3 standard deviations apart



Why is Resolution important ?

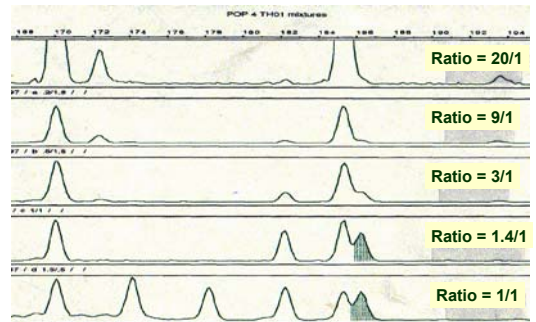
GenePrint CTTv

Locus	Seq.	Repeat #	Variants
CSF1PO	TAGA	6-15	10.3
TPOX	GAAT	6-13	
TH01	TCAT	5-10	9.3, 5.3, 6.1, 7.17.3 8.3, 10.3, 13.3
vWA	TCTG/TCTA	11-21	15.2, 18.2, 18.3, 19.2

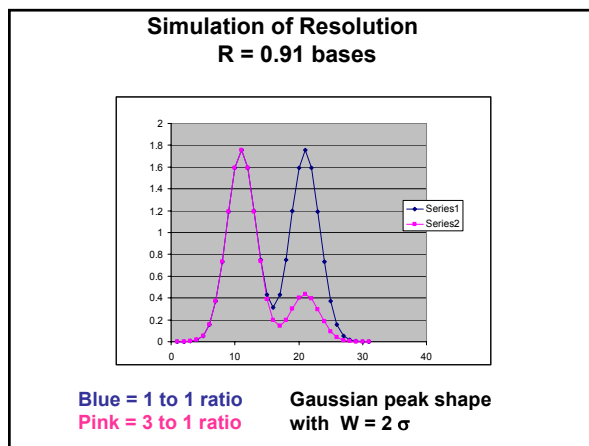
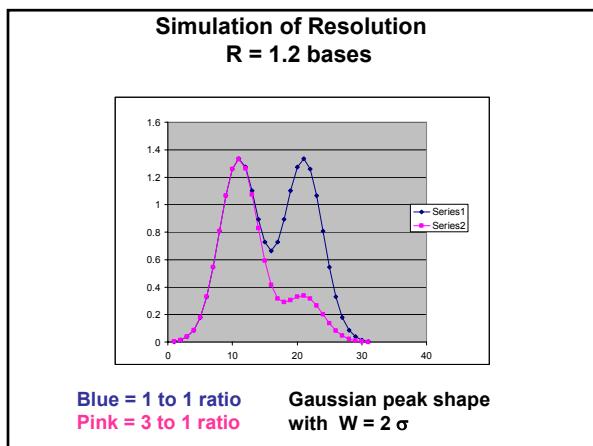
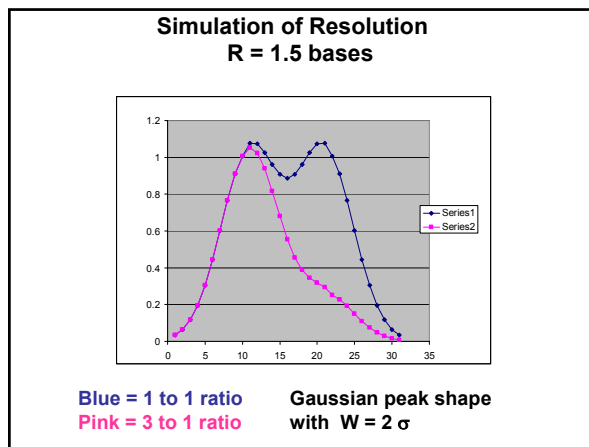
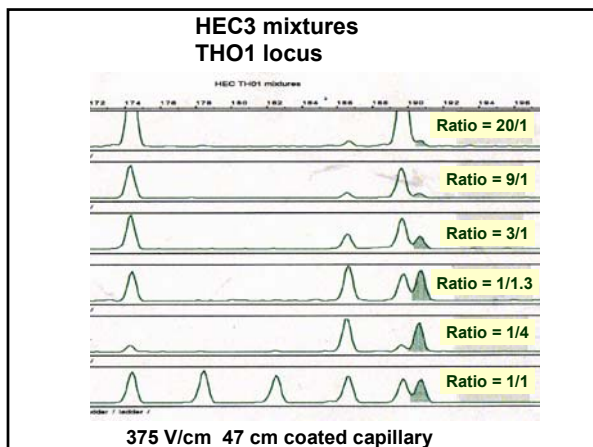


The presence of 1 and 2 bp variants places great constraints on the analysis!!

POP4 mixtures TH01 locus



320 V/cm 47 cm uncoated capillary
POP4 Polymer



Results

100 samples AmpFISTR Blue (D13S1358, vWA, FGA)
Consisting of Caucasian and Nigerian population samples

Average St. Dev.= 0.19 bases

Conditions:
3% Hydroxyethyl cellulose (MW approx. 40,000)
60C, 7.1M Urea, 100mM TBE, 40 cm 50um DB-17 capillary, 15kV, 30s @ 2.4 kV inj.

Other studies show Std Dev from 0.07 to 0.22 with typical results approx. 0.12 (polydimethyl acrylamide)

- ## Thoughts on Resolution
- There is a trade off between resolution and run time
 - Increased resolution can result in better mixture analysis
 - Resolution can degrade with time due to absorption on the inner capillary wall
 - Keep the column short and use thick polymers
 - Or use longer capillaries (but will have slower run times)

Precision (Resolution isn't enough!)

Microvariants can appear in the 4 base repeat motif present in these STRs

If 3 X Std. Dev. is greater than 0.5, then a certain number of 9.3 peaks will be labeled 10.

$\sigma = 99.7$

Precision

Current values in the literature range from 0.12 to 0.24 depending on the system and type of repeat. Most papers in the forensic literature report values under 0.15.

What affects precision?

Variants in STR sequence

Locus	allele	N	sequence
vWA	15	5	(XXXX) ₁₅ TCCA(TCTA)(TCCA) ₂
		1	(XXXX) ₁₅ (TCTA) ₂ TCCA(TCTA) ₂
		1	(YYYY) ₁₅ TCCA(TCTA)(TCCA) ₂

where XXXX = TCTA(TCTG)₄(TCTA)₁₀
 and YYYY = TCTA(TCTG)₃(TCTA)₁₁

These sequence variants mean that two alleles of the same size do not necessarily have the same sequence.

Lazaruk et al., Forensic Science International 119(2001)1-10

Casework vs Databasing

Quality of sample also affects precision

- PCR inhibitors
- Sample quantity
- Sample quality
- Ionic impurities

Temperature Effects and DNA Secondary Structure

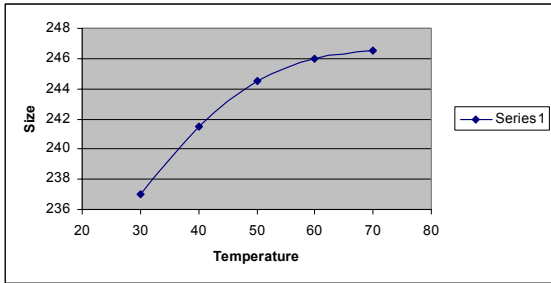
- Even under highly denaturing conditions DNA can self associate
- Differences in conformation can affect migration time
- Increase precision by limiting this effect?

How to avoid 2° Structure Effects

- Elevate Temperature to 60°
- Add Strong Denaturants
 - 7-8M Urea
 - Pyrrolidinone
- Examine response of 250, 340 peaks in ROX ladder

Rosenblum et al., Nucleic Acids Res., 1997

Change in size of GS 250 peak with Temperature (Tamra Std)



4% pDMA with 8M urea and 5% 2-pyrrolidinone
 *Rosenblum et al., Nucleic Acids Res.(1997) 25,19, 2925

Precision and Resolution

In addition to affecting the size of the DNA, temperature can affect the precision of the results. Elevated temperatures melt out DNA 2° structure, increasing the precision of the analysis. However, resolution is lost as a result of decreased viscosity. 100mM TBE 2% HEC, DB-17 Capillary

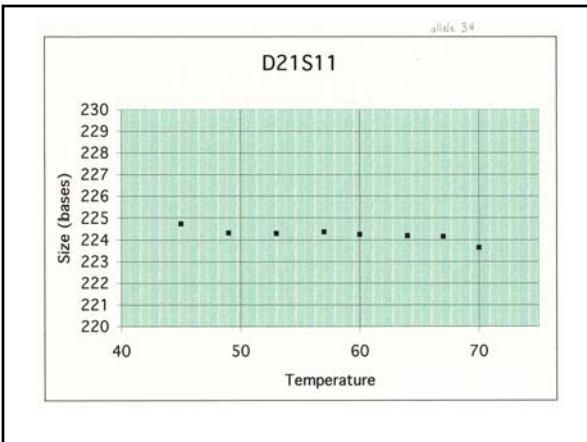
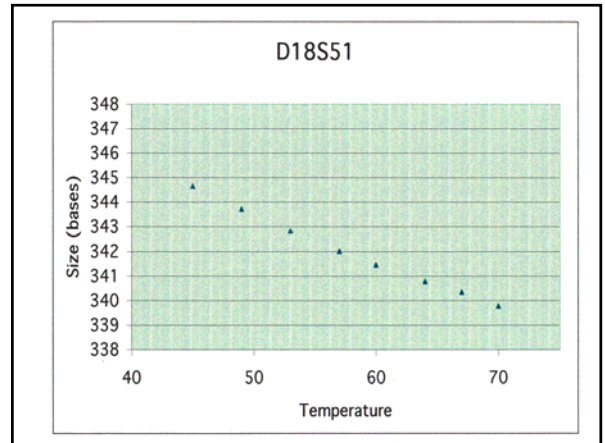
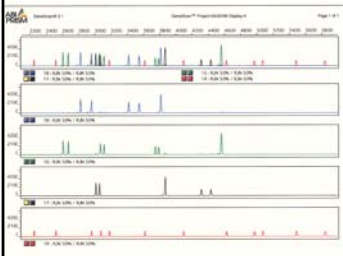
Temp	Resolution at 200bases	THO1 Allele 5	Std. Dev. (bases)
30	1.3	197.4	0.2
45	1.6	196.0	0.08
60	1.7 (n=7)	195.6 (n=7)	0.07 (n=200+)

So what is the effect of Temperature on Profiler+?

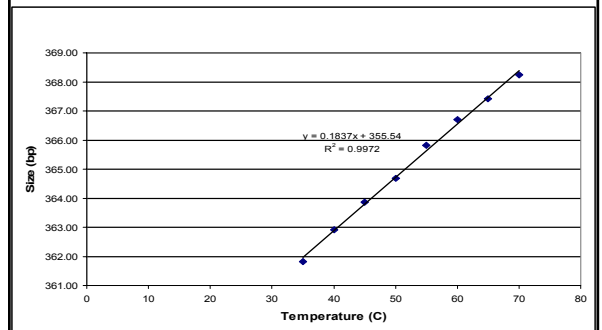
Could This affect precision?

Why do some band shifts occur at only one locus

Examine the allelic ladder at temperatures from 40-70 C



DYS 385-14 POP4 Size vs Temperature



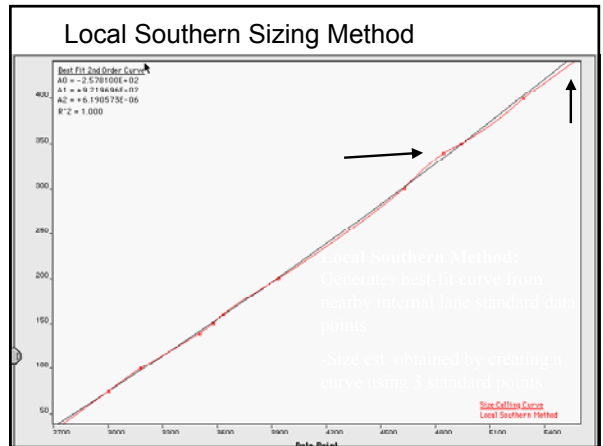
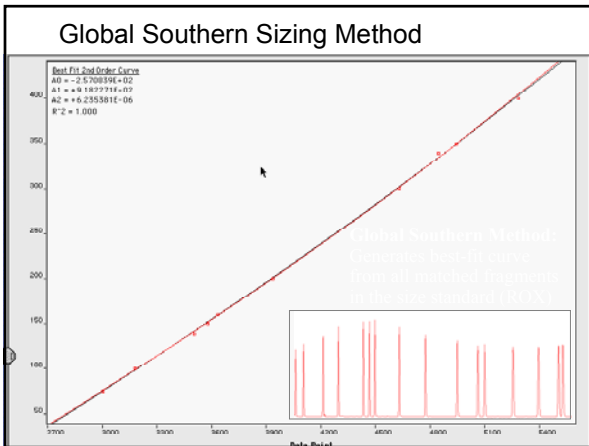
Effect of Temperature on Allele Size

POP4, pH 8, 350V/cm, 45-70°C

STR	Allele	Size*	Slope*	Std. Dev
D3S1358	12	111.2	-0.10	0.01
vWA	21	194.9	-0.07	0.02
FGA	30	264.7	-0.14	0.02
Amel	X	103.5	-0.13	0.01
D8S1179	19	170.4	-0.16	0.02
D21S11	36	232.4	-0.03	0.01
D18S51	26	341.9	-0.18	0.01
D5S818	7	131.2	-0.09	0.01
D13S317	8	205.0	-0.12	0.01
D7S820	15	292.8	-0.09	0.01

*Estimated size at 61°C
* °C/base, ave. of 4 measurements

- ### Effect of Operator Chosen Sizing Method
- **Global Southern Method:** Generates best-fit curve from all matched fragments in the size standard
 - **Local Southern Method:** Generates best-fit curve from only nearby internal lane standard data points



Effect of Operator Chosen Sizing Method

Global Southern Sizing			Local Southern Sizing		
allele #	average slope	SD of ave.	allele #	average slope	SD of ave.
CSF1PO			CSF1PO		
7	-0.052	0.01	7	-0.027	0.01
10	-0.050	0.01	10	-0.135	0.02
12	-0.047	0.01	12	-0.103	0.02
14	-0.048	0.01	14	-0.156	0.01
VWA			VWA		
14	-0.085	0.002	14	-0.060	0.009
15	-0.087	0.004	15	-0.059	0.009
17	-0.089	0.006	17	-0.097	0.008

- **Global Southern:**
 - Similar slopes within a locus
 - Differential response in slopes between loci
- **Local Southern:**
 - Differential response between and within loci
 - Many slopes significantly larger (-0.156 vs. -0.104)

Hartzell, Muncy, McCord, Forensic Science International, 2003, 133, 228-234.

- ### Implications of Temperature Studies
- Temperature affects precision through sample denaturation**
- New studies indicate there is a variable response to temperature especially between loci**
- The effect is far more pronounced in local southern**
- Temperature control is important because it affects both precision and resolution.**
- Band shifts are a natural consequence of differential response to temperature**

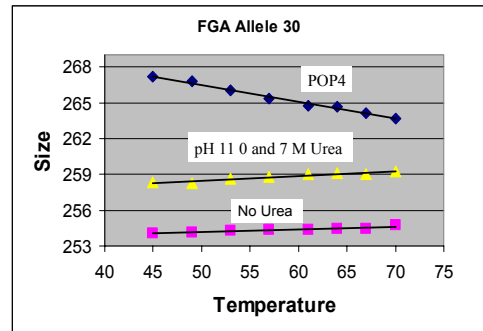
High pH Studies

DNA analysis can proceed at elevated pH (11+)
At such pH values buffers highly denaturing

Will temperature stability improve if conditions are more denaturing?

Yes, but capillary lifetime suffers

Temperature vs estimated size for POP4 and CAPs buffer (pH 11) at 0 and 7M urea



Implications of pH & Temp. Studies

STR size and migration varies in response to temperature and sizing method - secondary structure!

Temperature response is minimized at high pH with fluorocarbon capillaries but capillary lifetime is limited at this pH

What does this tell us about day to day applications?

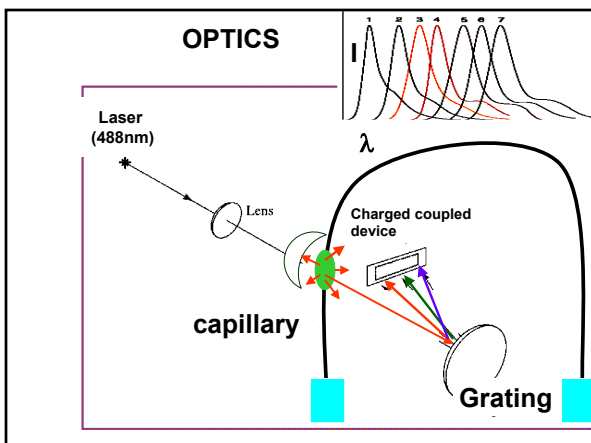
Bottom line:

Use global southern sizing.

Keep temperature constant !

Nock, McCord et al. *Electrophoresis*, 2001, 22, 755-762
Hartzell, Muncy, McCord, *Forensic Science International*, 2003, 133, 228-234.,

Detection



Detection Issues

- Fluorescent dyes
 - spectral emission overlap
 - relative levels on primers used to label PCR products
 - dye "blobs" (free dye)
- Virtual filters
 - hardware (CCD camera)
 - software (color matrix)

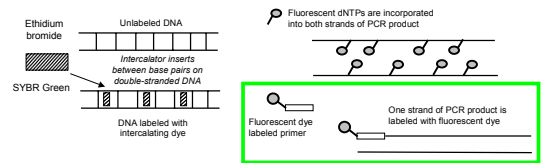
Filters determine which wavelengths of light are collected onto the CCD camera

Laser Used in ABI 310

- Argon Ion Laser
- 488 nm and 514.5 nm for excitation of dyes
- 10 mW power
- Lifetime ~5,000 hours (1 year of full-time use)
- Cost to replace ~\$5,500
- Leads to highest degree of variability between instruments and is most replaced part
- Color separation matrix is specific to laser used on the instrument

Methods for Fluorescently Labeling DNA

- Intercalating Dyes (post-PCR)
- Dye-labeled nucleotide insertion during PCR
- Dye-labeled primer insertion during PCR

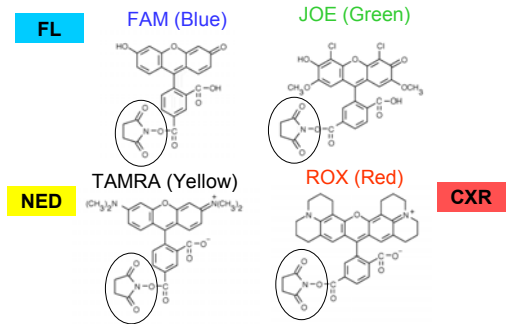


Butler, J.M. (2001) *Forensic DNA Typing*, Figure 10.2, ©Academic Press

Fluorescent Labeling of PCR Products

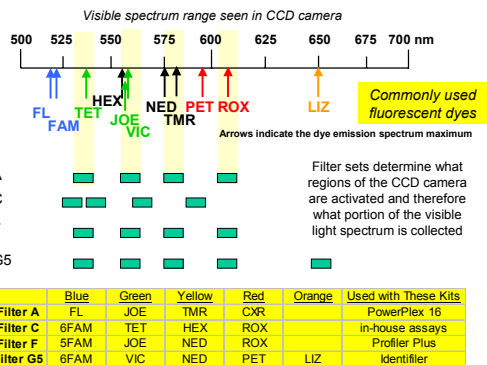
- Dyes are attached to one primer in a pair used to amplify a STR marker
- Dyes are coupled to oligonucleotides (primers) through NHS-esters and amine linkages on the 5' end of the primer: Dye-(CH₂)₆-primer
- Dye-labeled oligonucleotides are incorporated during multiplex PCR amplification giving a specific color "tag" to each PCR product
- PCR products are distinguished using CCD imaging on the 310

Fluorescent Dyes Used in 4-Color Detection

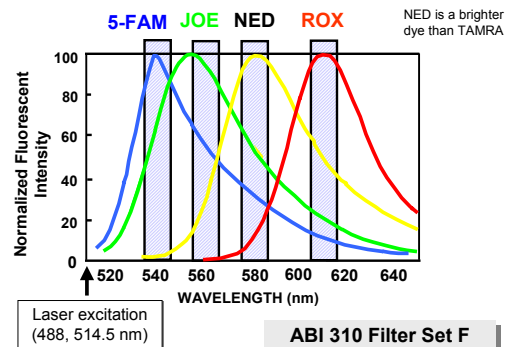


Butler, J.M. (2001) *Forensic DNA Typing*, Figure 10.3, ©Academic Press

Virtual Filters Used in ABI 310



Fluorescent Emission Spectra for ABI Dyes



Please Note!

- There are no filters in a 310
- Its just the choice of pixels in the CCD detector
- All the light from the grating is collected
- You just turn some pixels on and some off

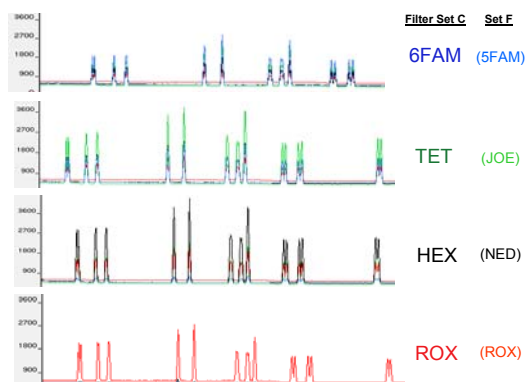
You then make a matrix

The matrix is the solution to a problem
what's the contribution at any given wavelength (filter set) from each dye ?

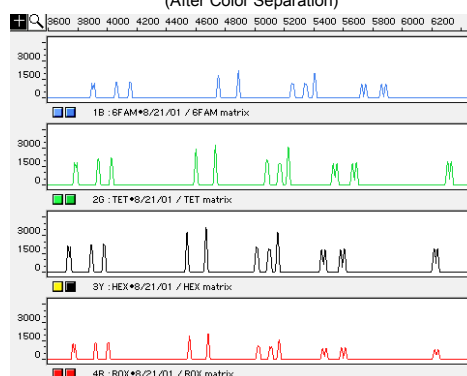
There are 4 dyes

- Remember algebra from high school?
- To solve a problem with 4 unknowns, you need 4 equations

Matrix Standards (Raw Data)



Matrix Standards (After Color Separation)



For Example

$$I_{540} = bx_b + gy_b + yz_b + rw_b \text{ intensity of blue}$$

$$I_{560} = bx_g + gy_g + yz_g + rw_g \text{ intensity of green}$$

$$I_{580} = bx_y + gy_y + yz_y + rw_y \text{ intensity of yellow}$$

$$I_{610} = bx_r + gy_r + yz_r + rw_r \text{ intensity of red}$$

Where
b is the %blue labeled DNA
g is the %green labeled DNA, etc.

x,y,z,w are the numbers in the matrix (sensitivity to each color)

POP4STRMODF				
Reactions				
	B	G	Y	R
B	1.0000	0.8502	0.1380	0.0009
G	0.8300	1.0000	0.7622	0.0051
Y	0.6416	0.8324	1.0000	0.1102
R	0.4493	0.6484	0.7851	1.0000

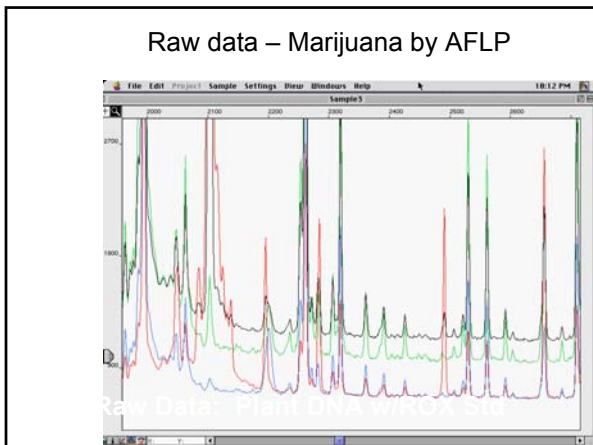
If you solve xyzw for each dye individually
Then you can determine any mixture

Matrix File Table from an ABI 310

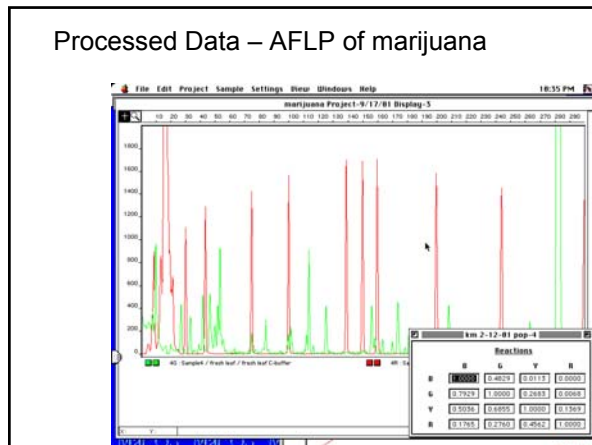
POP4STRMODF				
Reactions				
	B	G	Y	R
B	1.0000	0.8502	0.1380	0.0009
G	0.8300	1.0000	0.7622	0.0051
Y	0.6416	0.8324	1.0000	0.1102
R	0.4493	0.6484	0.7851	1.0000

These values are used by the GeneScan® Analysis Software to separate the various dye colors from one another. The letters B, G, Y, and R represent the dye colors Blue, Green, Yellow, and Red, respectively.

Raw data – Marijuana by AFLP



Processed Data – AFLP of marijuana



Comments on Matrices (Multi-Component Analysis)

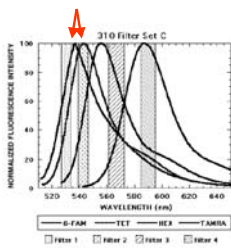
- Make sure that the right filter set and matrix are applied when collecting data
- You can always apply another matrix to a sample collected on the ABI 310 but it must be run with the right filter set (matrix must be run first with ABI 3100)
- It is important to update matrices on a regular basis (depending on use) due to differences in laser power over time
- A good indication of when to run a new matrix is the observation of pull-up between dye colors when peaks are smaller than ~4,000 RFUs

If Wrong Filter Is Used...



ABI 310 Filter C

	B	G	Y	R
B	1.0000	0.5194	0.0815	0.0032
G	0.0002	1.0000	0.4959	0.0049
Y	0.5487	0.5353	1.0000	0.0437
R	0.2515	0.3116	0.3887	1.0000

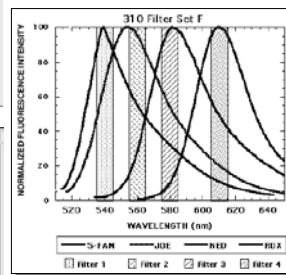


More color overlap between blue and green dyes

Matrices for Filter F with POP-4 vs. POP-6 (on the same ABI 310)

	B	G	Y	R
B	1.0000	0.6444	0.0487	0.0010
G	0.6027	1.0000	0.5554	0.0061
Y	0.3421	0.6146	1.0000	0.1060
R	0.1690	0.3478	0.5791	1.0000

	B	G	Y	R
B	1.0000	0.6416	0.0474	0.0006
G	0.6024	1.0000	0.5482	0.0063
Y	0.3436	0.6107	1.0000	0.1019
R	0.1686	0.3452	0.5827	1.0000



Values are very similar between the two matrices

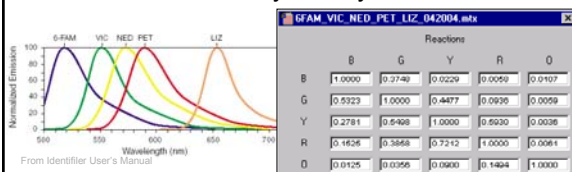
Same Dye Set and Filter F with Different ABI 310s

POP451R-MDF				
Reactions				
	B	G	Y	R
B	1.0000	0.9502	0.1380	0.0008
G	0.8500	1.0000	0.7622	0.0051
Y	0.6416	0.8324	1.0000	0.1102
R	0.4493	0.6484	0.7851	1.0000

SFAM_JDE_NED_NDR POP4				
Reactions				
	B	G	Y	R
B	1.0000	0.6444	0.0407	0.0010
G	0.6027	1.0000	0.5556	0.0061
Y	0.3421	0.6146	1.0000	0.1060
R	0.1690	0.3478	0.5791	1.0000

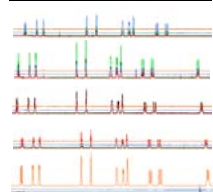
Instrument lasers make a big difference

5 x 5 matrix for 5-dye analysis on ABI 310

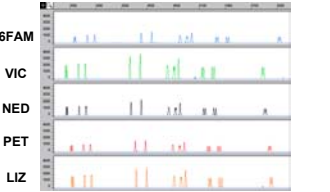


6FAM_VIC_NED_PET_LIZ_042004.mtx				
Reactions				
	B	G	Y	R
B	1.0000	0.3740	0.0229	0.0050
G	0.5323	1.0000	0.4477	0.0050
Y	0.2781	0.5408	1.0000	0.0030
R	0.1628	0.3868	0.7212	1.0000
D	0.0125	0.0395	0.0900	0.1404

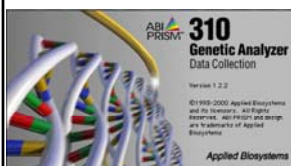
Raw Data for Matrix Standards



Processed Data (matrix applied with baselining)



310 Data Collection Software



Macintosh	Windows NT
1.0.2	1.0 (5-dye)
1.2.2	2.0 (5-dye)
2.1 (5-dye)	

- Controls 310 run conditions
- Translates light on CCD camera into electropherogram (raw data)
- Sample sheets and injection lists are created

ABI manual is P/N 904958B

Injection List in Data Collection Software

- Lists samples to be analyzed (repeats can be easily performed)
- Sets virtual filter on CCD camera
- Sets electrophoresis time and voltage
- Sets injection time and voltage
- Sets run temperature
- If desired, sample analysis can be set up for automatic matrix color separation and sizing with internal standards using defined analysis parameters

Steps Performed in Standard Module

- **Capillary fill** – polymer solution is forced into the capillary by applying a force to the syringe
- **Pre-electrophoresis** – the separation voltage is raised to 10,000 volts and run for 5 minutes;
- **Water wash of capillary** – capillary is dipped several times in deionized water to remove buffer salts that would interfere with the injection process
- **Sample injection** – the autosampler moves to position A1 (or the next sample in the sample set) and is moved up onto the capillary to perform the injection; a voltage is applied to the sample and a few nanoliters of sample are pulled onto the end of the capillary; the default injection is 15 kV (kilovolts) for 5 seconds
- **Water wash of capillary** – capillary is dipped several times in waste water to remove any contaminating solution adhering to the outside of the capillary
- **Water dip** – capillary is dipped in clean water (position 2) several times
- **Electrophoresis** – autosampler moves to inlet buffer vial (position 1) and separation voltage is applied across the capillary; the injected DNA molecules begin separating through the POP-4 polymer solution
- **Detection** – data collection begins; raw data is collected with no spectral deconvolution of the different dye colors; the matrix is applied during Genescan analysis

Conclusions

Current advances in DNA typing are the result of:

- 1) The application of short tandem repeats
- 2) The development of multiplex PCR
- 3) Multichannel laser induced fluorescence
- 4) Capillary electrophoresis with entangled polymer buffers

Conclusions II

1. Systems for DNA typing must combine precision and resolution for identification of STR alleles reliably
2. Differential response to temperature can affect precision
3. Denaturation is important in maintaining reproducibility
4. Separation efficiency and precision are controlled via the temperature, polymeric buffer and the injection media.

Acknowledgements

Funding: National Institute of Justice

People:

- Ohio University
- Tim Nock
- Jim Dove
- Sharon Williams
- Dr. Federica Crivellente
- Kylie Muncy
- Brittany Hartzel
- Dr. Yin Shen
- Kerry Opel
- Dr. Nancy Tatarek
- Dr. Denise Chung
- Elsewhere
- Dr. Alice Isenberg FBI
- Dr. John Butler NIST
- Dr. Ralph Allen UVA
- Dr. David Mao J&W Scientific
- Dr. John Petersen UT-Galveston