

Topics and Techniques for Forensic DNA Analysis

Capillary Electrophoresis Fundamentals and Troubleshooting

**Florida Statewide
Training Meeting**

Indian Rocks Beach, FL
May 12-13, 2008



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Standards and Technology
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Questions?

- What are your biggest challenges with keeping your ABI 310/3100/3130xl running?
- What kind of signal intensity variation are you seeing between your different instruments?
- Have anyone seen uneven injection across a sample plate? (We believe this to be an autosampler calibration issue...e.g., position G10 or H12 does not inject properly)

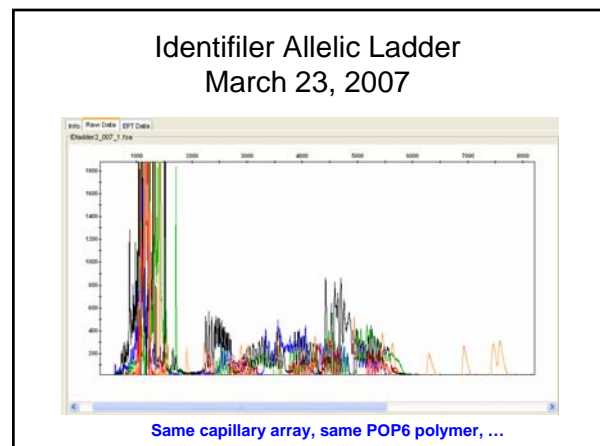
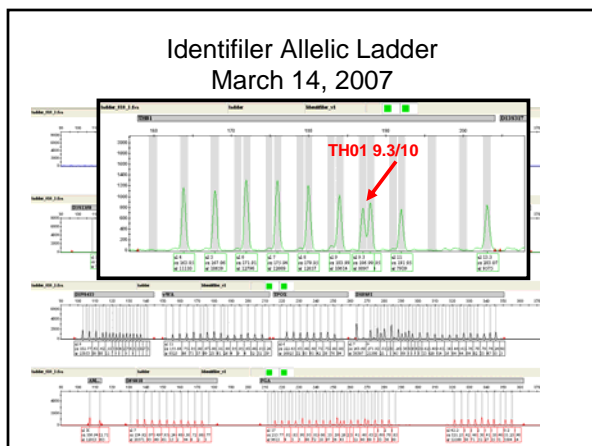
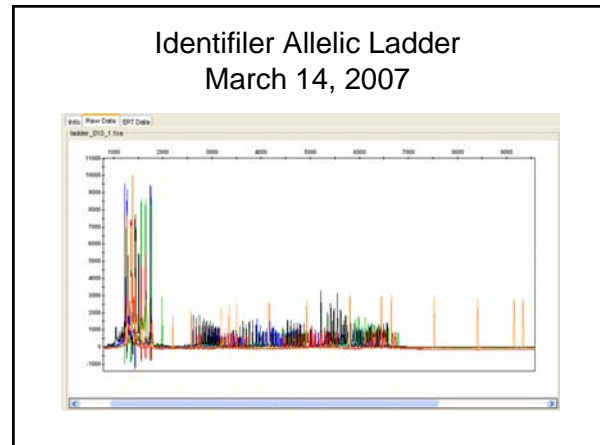
Planned Promega 2008 Meeting Troubleshooting Workshop

- Title: **"Principles of Interpretation and Troubleshooting of Forensic DNA Typing Systems"**
- Instructors: **John Butler (NIST) and Bruce McCord (FIU)**
- Date: **October 16, 2008** with Promega Int. Symp. Human ID

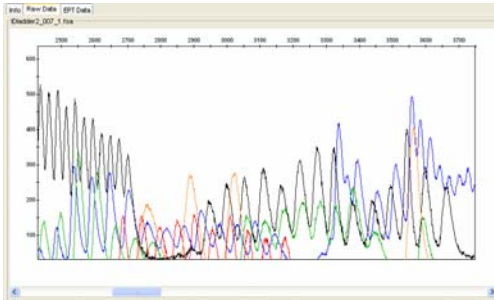
The workshop will consist of three parts:

- (1) a through examination of **theoretical issues with capillary electrophoresis** PCR amplification of short tandem repeat markers
- (2) a discussion of **how to properly set instrument parameters to interpret data** (including mixtures), and
- (3) **a review of specific problems seen by labs** submitting problematic data and commentary on possible troubleshooting solutions.

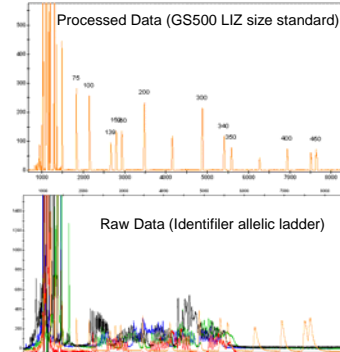
Seeking input of problems observed with CE systems



Examination of Resolution in TH01 Region

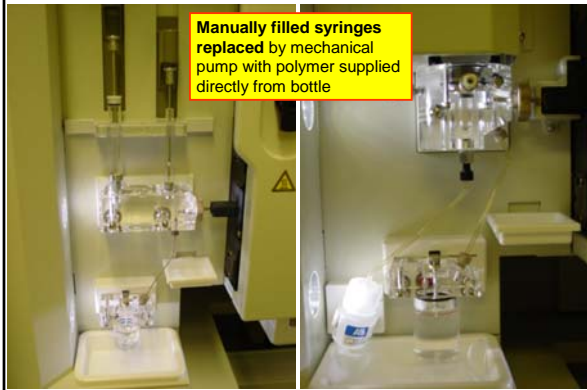


Examine the Size Standard...

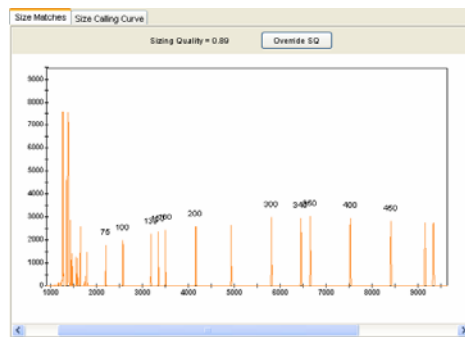


ABI 3100

ABI 3130xl
(upgraded from 3100)



The Size Standard Provides an Excellent Indicator of Performance on Every Sample



Review Article on STRs and CE

pdf available from <http://www.cstl.nist.gov/biotech/strbase/NISTpub.htm>

Electrophoresis 2004, 25, 1397-1412

Review

John M. Butler¹
Eric Blue²
Federica Crivellente^{3*}
Bruce R. McCord²

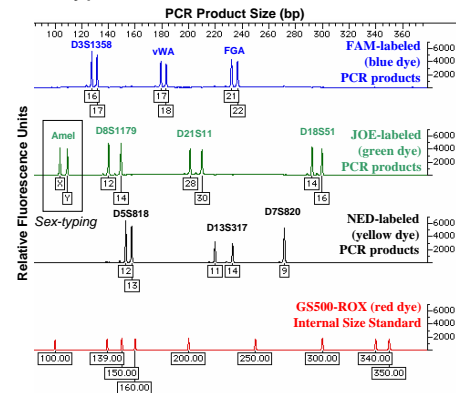
Forensic DNA using the ABI for STR anal

DNA typing with short applications include such as the ABI Pres for many laboratories ing sample prepar results using CE sys in the context throughput and ease

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Genotype Results with Profiler Plus™ kit



Analytical Requirements for STR Typing

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

- Fluorescent dyes must be **spectrally resolved** in order to distinguish different dye labels on PCR products
- PCR products must be **spatially resolved** – desirable to have single base resolution out to >350 bp in order to distinguish variant alleles
- High **run-to-run precision** – an internal sizing standard is used to calibrate each run in order to compare data over time

Steps in STR Typing with ABI 310

Butler, J.M. (2005) *Forensic DNA Typing, 2nd Edition*, Figure 13.8, © Elsevier Science/Academic Press

Detection with Multiple Capillaries (Irradiation for Capillary Arrays)

ABI 3100, 3130, 3100Avant ABI 3700 MegaBACE

Process Involved in 310/3100 Analysis

- Separation**
 - Capillary – 50um fused silica, 43 cm length (36 cm to detector)
 - POP-4 polymer – Polydimethyl acrylamide
 - Buffer - TAPS pH 8.0
 - Denaturants – urea, pyroldinone
- Injection**
 - electrokinetic injection process (formamide, water)
 - importance of sample stacking
- Detection**
 - fluorescent dyes with excitation and emission traits
 - CCD with defined virtual filters produced by assigning certain pixels

Ohm's Law

- $V = IR$ (where V is voltage, I is current, and R is resistance)
- Current, or the flow of ions, is what matters most in electrophoresis
- CE currents are much lower than gels because of a higher resistance in the narrow capillary
- CE can run a higher voltage because the capillary offers a higher surface area-to-volume ratio and can thus dissipate heat better from the ion flow (current)

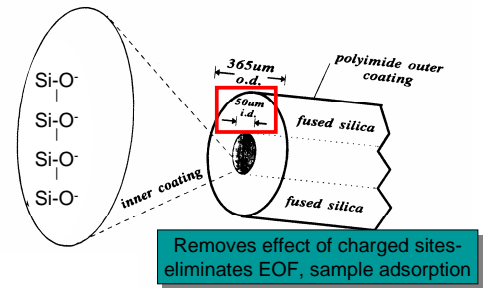
Separation Issues

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

Why TAPS instead of Tris-borate (TBE) buffer?

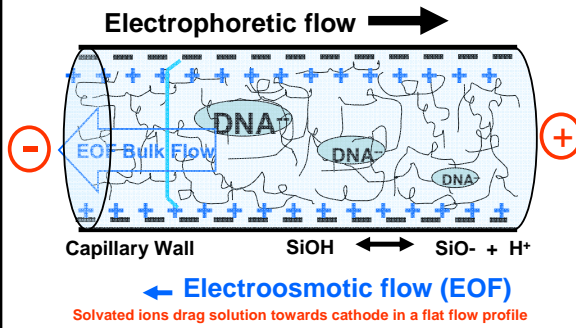
- TBE is temperature/pH sensitive
 - as temperature increases, the pH decreases (0.02 pH units with every 1 °C); this is the principle by which TaqGold activation works
- At lower pH, fluorescence emission of dyes decreases
 - see Singer and Johnson (1997) *Proceedings of the Eighth International Symposium on Human Identification*, pp. 70-77
- Thus when running at 60 °C on the ABI 310, if Tris-borate was used, fluorescent intensity of PCR products would be lower

Capillary Coating



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

Capillary Wall Coatings Impact DNA Separations

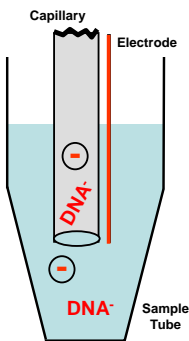


How to Improve Resolution?

1. Lower Field Strength
2. Increase Capillary Length
3. Increase Polymer Concentration
4. Increase Polymer Length

All of these come at a cost of longer separation run times

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
 E is the electric field applied
 t is the injection time
 r is the radius of the capillary
 μ_{ep} is the mobility of the sample molecules
 μ_{eof} is the electroosmotic mobility

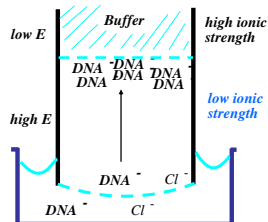
$[DNA_{sample}]$ is the concentration of DNA in the sample
 λ_{buffer} is the buffer conductivity
 λ_{sample} is the sample conductivity

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

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 = $\mu_{ep} = \text{velocity} / \text{electric field}$



Steps Performed in Standard Module

See J.M. Butler (2005) *Forensic DNA Typing*, 2nd Edition; Chapter 14

- **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

Comments on Sample Preparation

- Use high quality formamide (<100 $\mu\text{S}/\text{cm}$)!
 - ABI sells Hi-Di formamide
 - regular formamide can be made more pure with ion exchange resin
- Deionized water vs. formamide
 - Biega and Duceaman (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...

January 6, 2005 Letter from Applied Biosystems to ABI 310 Customers

- “Testing has shown that Hi-Di Formamide denatures DNA **without the need to heat samples...**”
- In other words, no heat denaturation and snap cooling needed!

Applied Biosystems Okays Use of Deionized Water for DNA Sequencing

Technical Bulletin #1 Issued August 2006

Applied Biosystems 3730/3730xl DNA Analyzer

Subject: Influence of Sequencing Injection Solution on 3730/3730xl DNA Analyzer Performance

In this Bulletin:

- Three Loading Solutions Tested on Page 1
- Loading Solution Test Data on Page 2
- Recommendations on Page 6
- Guidelines for Use on Page 6

Three Loading Solutions Tested

Loading Solution Background

Applied Biosystems presently recommends the use of Hi-Di™ Formamide as the sample-loading solution for all Applied Biosystems DNA sequencers to ensure sample preservation and resistance to evaporation. However, many users of the 3730 choose either deionized water or dilute EDTA solutions. These choices are driven largely by cost and safety/hazardous material considerations.

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)

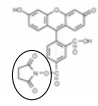
Filters determine which wavelengths of light are collected onto the CCD camera

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

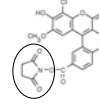
Amine Reactive Dyes used in Labeling DNA

FAM (Blue)



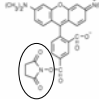
Emission 520

JOE (Green)



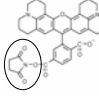
Emission 548

TAMRA (Yellow)

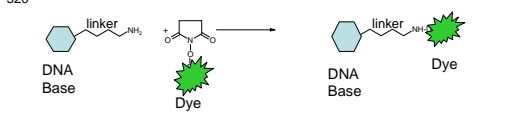


Emission 580

ROX (Red)



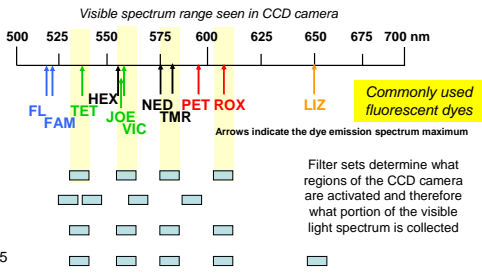
Emission 605



The succinimidyl ester reacts rapidly with amine linkers on DNA bases

Virtual Filters Used in ABI 310

Visible spectrum range seen in CCD camera



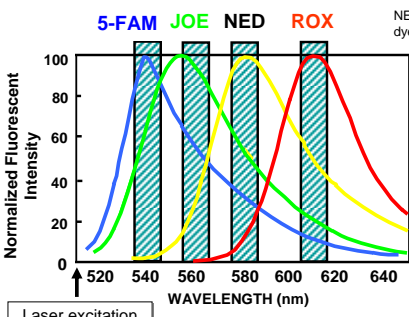
Commonly used fluorescent dyes

Arrows indicate the dye emission spectrum maximum

Filter sets determine what regions of the CCD camera are activated and therefore what portion of the visible light spectrum is collected

Filter	Blue	Green	Yellow	Red	Orange	Used with These Kits
Filter A	FL	JOE	TMR			PowerPlex 16
Filter C	6FAM	TET	HEX	ROX		in-house assays
Filter F	5FAM	JOE	NED	ROX		Profiler Plus
Filter G5	6FAM	VIC	NED	PET	LIZ	Identifier

Fluorescent Emission Spectra for ABI Dyes



5-FAM JOE NED ROX

Normalized Fluorescent Intensity

WAVELENGTH (nm)

Laser excitation (488, 514.5 nm)

ABI 310 Filter Set F

Butler, J.M. (2001) Forensic DNA Typing, Figure 10.4, ©Academic Press

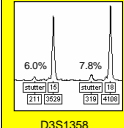
Please Note!

- 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

Deciphering Artifacts from the True Alleles

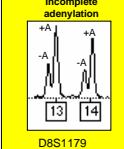
Biological (PCR) artifacts

Stutter products

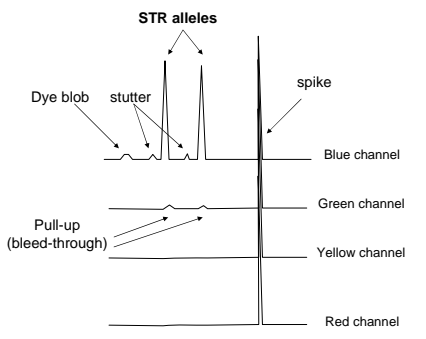


D3S1358

Incomplete adenylation



D8S1179



STR alleles

dye blob

stutter

spike

Blue channel

Green channel

Yellow channel

Red channel

Pull-up (bleed-through)

Butler, J.M. (2005) Forensic DNA Typing, 2nd Edition, Figure 15.4, © Elsevier Science/Academic Press

Dye Blobs (“Artifacts”)

- Free dye (not coupled to primer) can be injected into the CE capillary and interfere with detection of true STR alleles
- Dye blobs are wider and usually of less intensity** than true STR alleles (amount depends on the purity of the primers used)
- Dye blobs usually appear at an apparent size that is unique for each dye (e.g., FAM ~120 bp, PET ~100 bp)

Poor primer purity

Dye Blob Problems with Some PCR Primers

Individual Y-STR Locus Amplifications

Removal of Dye Artifacts Following PCR Amplification

Butler, J.M., Shen, Y., McCord, B.R. (2003) The development of reduced size STR amplicons as tools for analysis of degraded DNA. J. Forensic Sci 48(5): 1054-1064.

Conclusions

DNA typing by capillary electrophoresis involves:

- 1) The use of entangled polymer buffers
- 2) Injection by sample stacking
- 3) Multichannel laser induced fluorescence
- 4) Internal and external calibration

Practical Aspects of ABI 310/3100 Use

ABI Genetic Analyzer Usage at NIST

- ABI 310 x 2 (originally with Mac, then NT)
 - 1st was purchased in 1996
 - 2nd was purchased in June 2002
- ABI 3100 (Data collection v1.0.1)
 - Purchased in June 2002
 - Original data collection software retained
- ABI 3130xl upgrade (Data collection v3.0)
 - Purchased in April 2001 as ABI 3100
 - Upgraded to ABI 3130xl in September 2005
 - Located in a different room

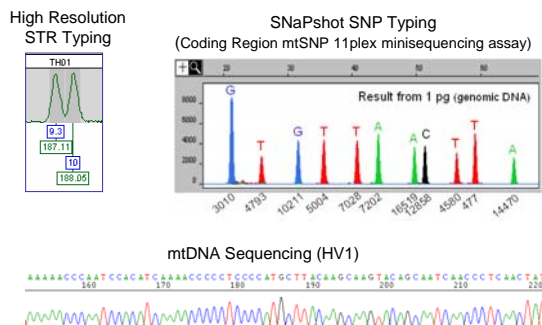
Jan 2007 – upgraded to 3130xl with data collection v3.0

Our Use of the ABI 3100

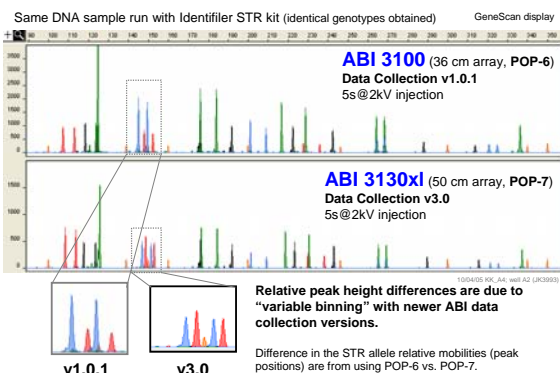
- Data collection software, version 1.0.1
- **POP-6** with 36 cm capillary array
- STR kits and in-house assays for autosomal STRs, Y-STRs, and miniSTRs
- SNaPshot assays for mtDNA SNPs, Y-SNPs, and autosomal SNPs
- DNA sequencing for mtDNA and STR repeat sequencing

We can routinely get more than 400 runs per capillary array by not changing the polymer between applications

NIST ABI 3100 Analysis Using POP-6 Polymer



Comparison of ABI 3100 Data Collection Versions



Consumables for ABI 310/3100

What we use at NIST

- A.C.E.™ Sequencing Buffer 10X (Amresco)
 - \$155/L = \$0.0155/mL 1X buffer (**costs 20 times less!**)
 - <http://www.amresco-inc.com>
- 3700 POP-6 Polymer (Applied Biosystems)
 - \$530 / 200 mL = \$2.65/mL (**costs 20 times less!**)

What ABI protocols suggest

- 10X Genetic Analyzer Buffer with EDTA
 - \$78/25 mL = \$0.312/mL 1X buffer (ABI)
 - 3100 POP-4 Polymer
 - \$365 / 7 mL = \$52/mL
- 2004 prices**

Maintenance of ABI 310/3100/3130

- Syringe – leaks cause capillary to not fill properly
- Capillary storage & wash – **it dries, it dies!**
- Pump block – cleaning helps insure good fill
- Change the running buffer regularly

YOU MUST BE CLEAN AROUND A CE!

Overall Thoughts on the ABI 310/3100/3130

- Settling on a common instrument platform has been good for the forensic DNA community in terms of data consistency (this is also true with the use of common STR kits)
- I am concerned that the community is very dependent primarily on one company...
- I really like using the instrument and can usually get nice data from it
- Like any instrument, it has its quirks...

Bruce McCord's Profiles in DNA Article

PROFILES IN DNA Volume 6 (2), Sept 2003, pp. 10-12

TECH TIPS

Troubleshooting Capillary Electrophoresis Systems

By Bruce McCord
Associate Professor of Forensic Chemistry, Ohio University, Athens, Ohio

INTRODUCTION
The development of capillary electrophoresis (CE) has played a key role in bringing about the modern application of DNA typing. Forensic laboratories are the beneficiaries of this new technology, but many practitioners are not fully aware of the underlying principles of the CE system. This article attempts to address the important issues in CE separations to aid analysts in troubleshooting problematic separations. The key to producing good DNA separations is to understand the principles underlying the injection, separation and detection of each allele. These points are addressed below.

SEPARATION
DNA analysis by CE is performed using entangled polymer buffers (Figure 1). These buffers can be easily pumped into a capillary prior to a separation and pumped out at its conclusion, providing a fresh separation matrix for each run. A typical buffer for forensic DNA separation contains 4% polydimethyl acrylamide (pDMA), buffered to pH 8

The key to producing good DNA separations is to understand the principles underlying the injection, separation and detection of each allele.

CE Troubleshooting Bruce McCord, AAFS 2006 Workshop (Seattle, WA) February 20, 2006

Outline for This Section

1. Chemistry/molecular biology problems – stutter, -A, degradation, inhibition, low copy #
2. Sample and buffer problems – formamide, urea, water, salt concentration, free dye (“dye blobs”)
3. External factors – power supply, room temperature, cleanliness, voltage leaks
4. Instrument problems – optical system, capillary clogging, air bubbles, syringe leaks
5. Troubleshooting benchmarks/QC monitoring

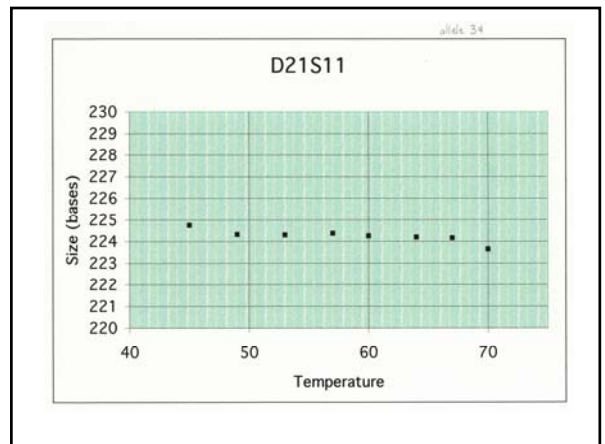
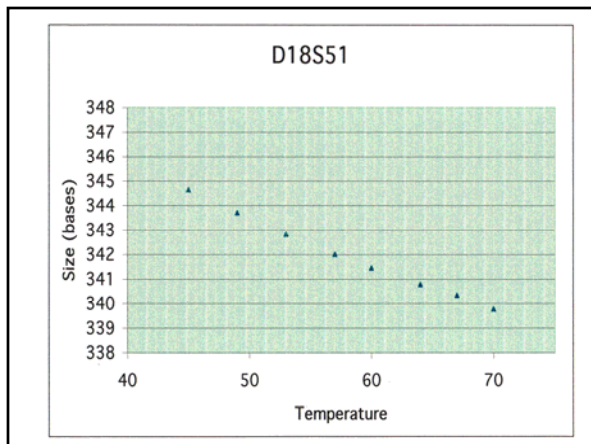
3. External Factors

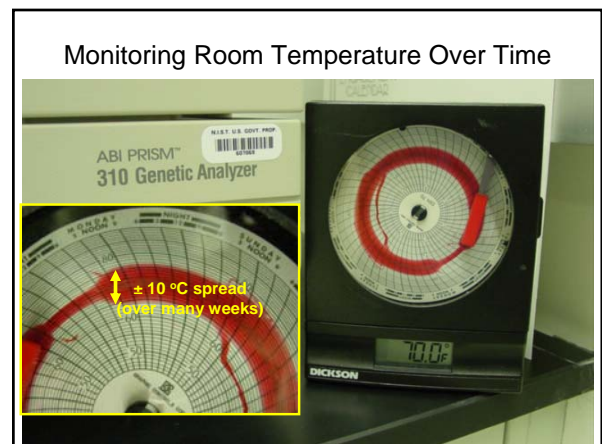
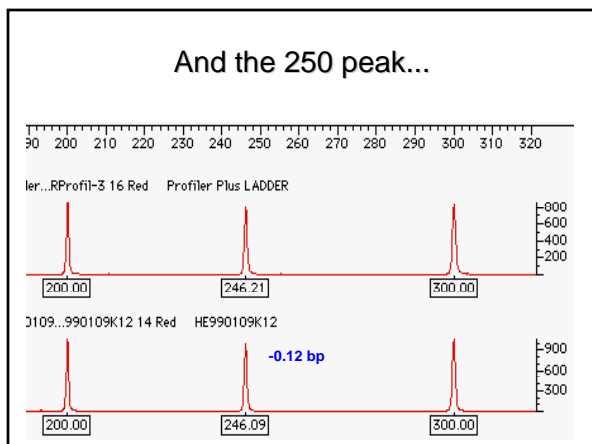
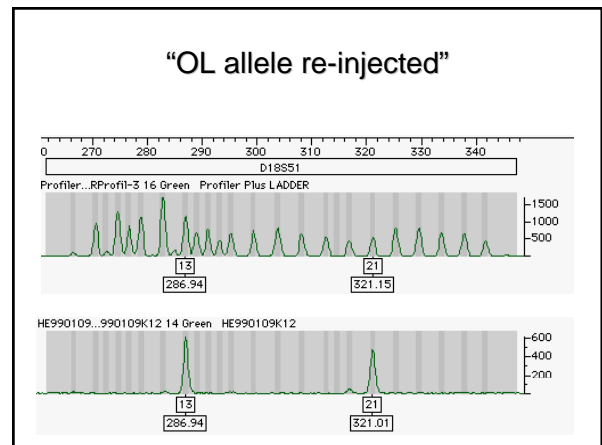
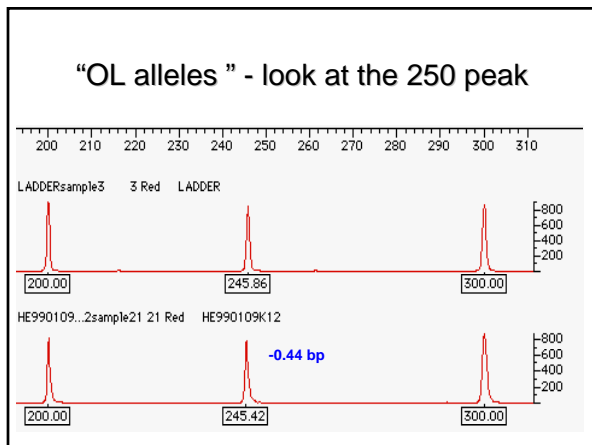
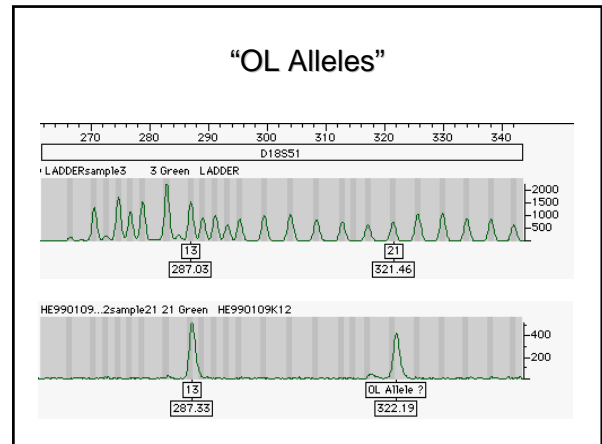
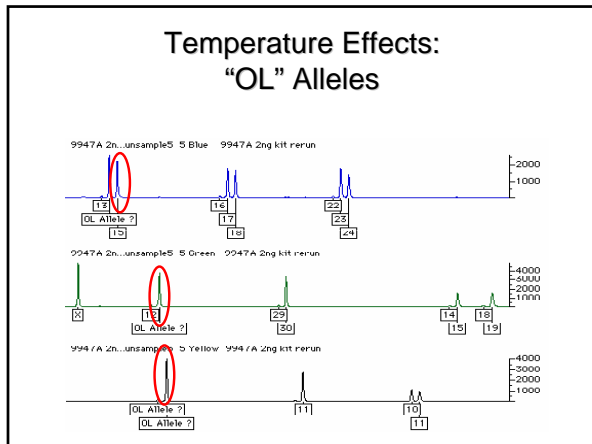
- Room temperature
 - Variations in room temperature can cause mobility shifts with band shifts and loss of calibration
 - Temperature is also important due to effects of high humidity on electrical conductance
- Cleanliness
 - Urea left in sample block can crystallize and catalyze further crystal formation causing spikes, clogs and other problems.
 - Best bet is to keep polymer in system and not remove or change block until polymer is used up.

Effect of Temperature on allele size


Temperature (°C)	Allele Size (bases)
45	267.5
50	266.5
55	265.5
60	264.5
65	264.0
70	263.5

Slope is 0.14 bases/degree centigrade
Therefore a small change in temperature has a big effect
(A 1-2 degree shift in temperature of the heat plate can produce an OL allele)






Temperature Probes




Refrigerator and freezer monitoring

Frig/Freeze Monitors \$240
 #DT-23-33-80 – USB Temperature Datalogger
 PLUS Software \$79.00 (#DT-23-33-60)

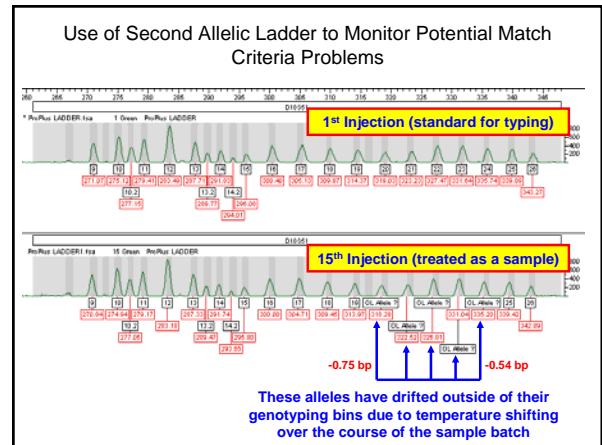
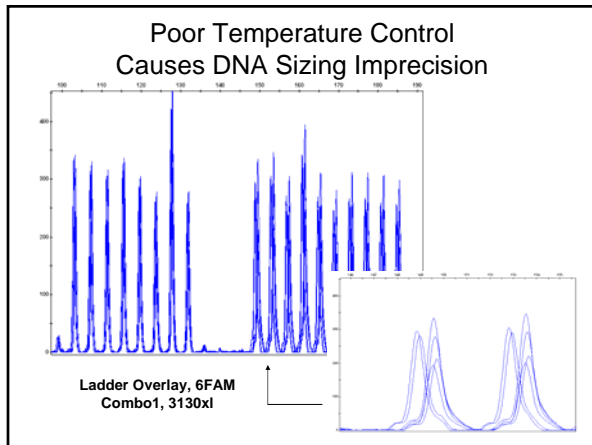
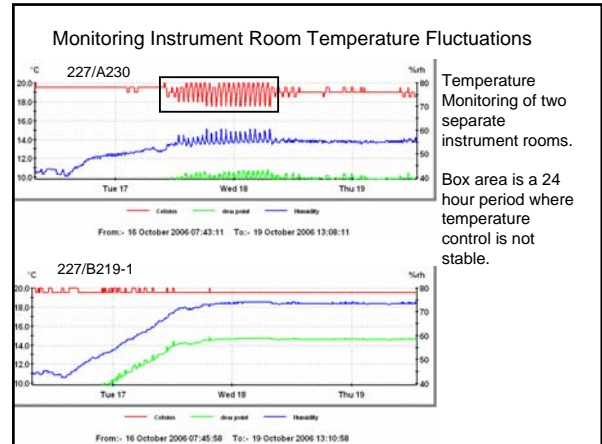
Room Monitors, # DT-23039-52 – USB
 Temperature-Humidity Datalogger \$91.00
 (Cole Parmer, Vernon Hills IL)



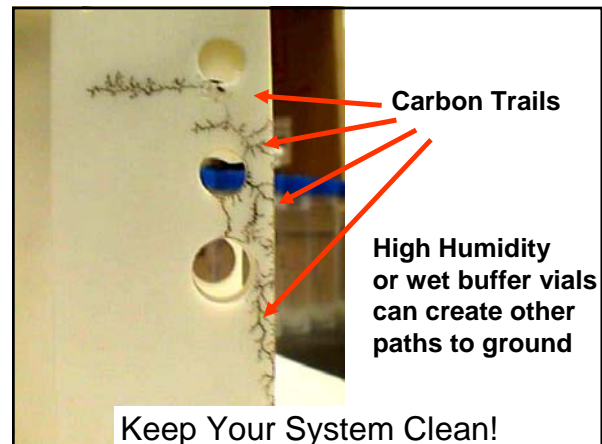
Room temperature monitoring



3130xl Genetic Analyzer



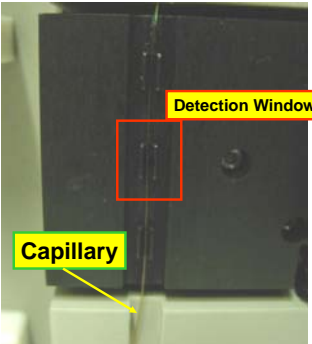
- ### Cleanliness
- Urea sublimates and breaks down to ionic components - these find a path to ground
 - Similarly wet buffer under a vial creates paths to ground
 - Capillary windows must be clear or matrix effects will occur
 - Laser will often assist in this process
 - Vial caps will transfer low levels of DNA to capillary



4. Instrumental Factors

- Optical System
 - Sensitivity changes with age, capillary diameter, capillary cleanliness, instrument calibration
- Fluidic System
 - Effects of bubbles, dust, urea crystals, leaks in syringe and capillary ferrule
- Matrix Calculations
 - Changes in buffer, optics, sample dye can alter the software calibrations
- Capillary Problems
 - Chemisorbed materials on capillary surface can produce osmotic flow, DNA band broadening and inconsistent resolution (meltdowns)

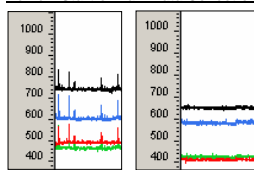
The Detection Window



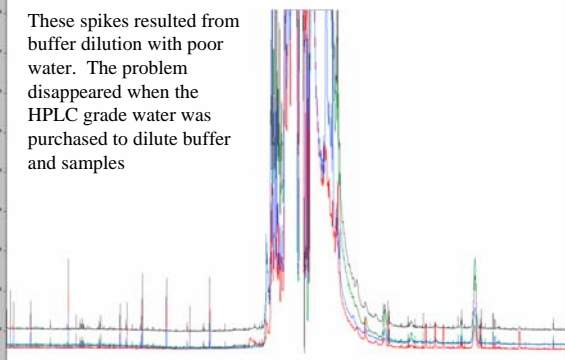
Make sure that the capillary window is lined up (if it is not, then no peaks will be seen)

Window may need to be cleaned with ethanol or methanol

Review Start of Raw Data Collection




Little spikes indicate need to change buffer... check current



These spikes resulted from buffer dilution with poor water. The problem disappeared when the HPLC grade water was purchased to dilute buffer and samples

Beware of Urea Crystals



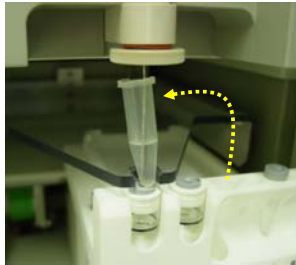
Urea crystals have formed due to a small leak where the capillary comes into the pump block

Urea sublimates and can evaporate to appear elsewhere

Use a small balloon to better grip the ferrule and keep it tight

Pump block should be well cleaned to avoid problems with urea crystal formation

Storage when ABI 310 is not in use

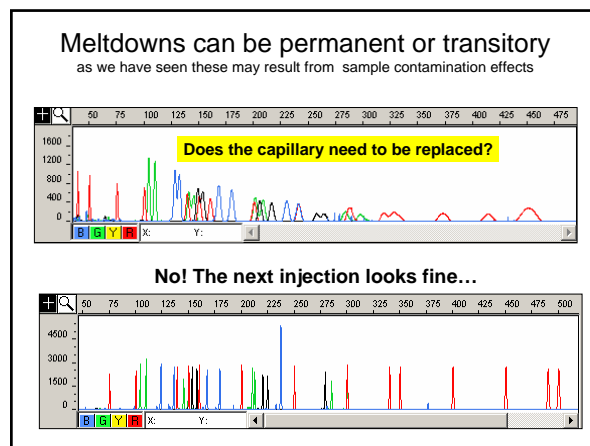
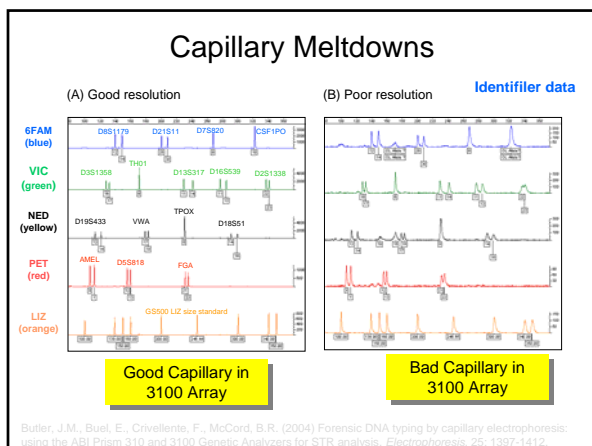


- Keep inlet of capillary in water...if it dries out then urea crystals from the polymer will clog the opening
- The waste vial (normally in position 3) can be moved into position
- A special device can be purchased from Suppelco to rinse the capillary off-line
- Store in distilled water
- Note that the laser is on when the instrument is on

Remember that the water in the open tube will evaporate over time...

Buffer Issues

- The buffer and polymer affect the background fluorescence- affecting the matrix
- Urea crystals and dust may produce spikes
- High salt concentrations may produce reannealing of DNA
- High salt concentrations affect current
- Low polymer concentrations affect peak resolution

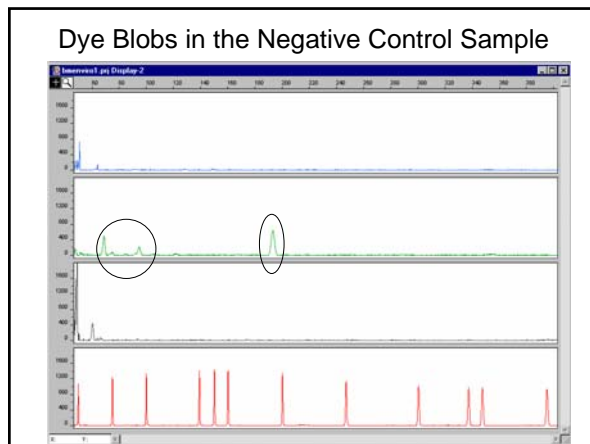
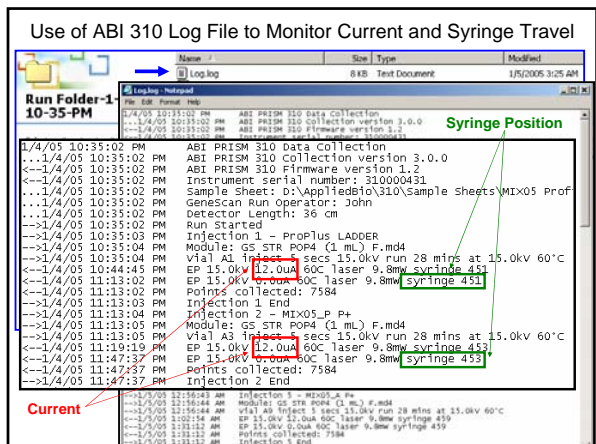


- ### Meltdowns may be the result of
- Bad formamide
 - Excess salt in sample/renaturation
 - Water in the polymer buffer
 - **Syringe leak** or bottom out
 - Poisoned capillary
 - Conductive polymer buffer due to urea degradation
 - Crack/shift in capillary window
 - Detergents and metal ions

- ### 5. Troubleshooting benchmarks
- **Monitor run current**
 - Observe syringe position and movement during a batch
 - Examine ILS (ROX) peak height with no sample
 - Observe "250 bp" peak in GS500 size standard
 - Monitor resolution of TH01 9.3/10 in allelic ladder and size standard peak shapes
 - **Keep an eye on the baseline signal/noise**
 - Measure formamide conductivity
 - Reagent blank – **are any dye blobs present?**
 - See if positive control DNA is producing typical peak heights (along with the correct genotype)

- ### Measurement of Current
- $V/I = R$ where R is a function of capillary diameter, [buffer], and buffer viscosity
 - In a CE system the voltage is fixed, thus changes in resistance in the capillary will be reflected in the current observed
 - Air bubbles, syringe leaks, alternate paths to ground, changes in temperature, changes in zeta potential, and contamination, will be reflected in the current
 - A typical current for a CE system with POP4 buffer is **8-12 μ A** (microamps)

- ### Syringe Travel
- The ABI 310 instrument also keeps track of the position of the syringe (in the log file)
 - Depending on the resistance to flow, the syringe will travel different lengths
 - Syringe leaks may be reflected in a longer distance traveled prior to each injection
 - These leaks occur around the barrel of the syringe and at the connection to the capillary block



Measuring Formamide Conductivity

(not this way)

The key is to measure the bottle when it comes in or buy the good stuff and immediately pipette it out into small tubes with or without ROX already added. Then freeze the tubes.

Do not ever open a cold bottle of formamide. Water will condense inside and aid in the formation of conductive formic acid.

Conclusion:

Troubleshooting is more than following the protocols

It means keeping watch on all aspects of the operation

1. Monitoring conductivity of sample and formamide
2. Keeping track of current and syringe position in log.
3. Watching the laser current
4. Watching and listening for voltage spikes
5. Monitoring room temperature and humidity

Multiplex_QA Article Published

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Research Article

Multiplex_QA: An exploratory quality assessment tool for multiplexed electrophoretic assays

Multiplex_QA is a data analysis tool for visualizing short- and long-term changes in the performance of multiplexed electrophoretic assays, particularly the commercial short tandem repeat (STR) kits used by the human forensic identity community. A number of quality metrics are calculated from the signal collected for the internal size standard included in nearly all multiplex assays. These quality metrics are related to the signal intensity, symmetry, retention, resolution, and noise of data collected by capillary electrophoresis systems. Interlocking graphical displays enable the identification of changes in the quality metrics with time, evaluation of relationships among the metrics, and detailed examination of electrophoretic features of particularly interesting analyses. While primarily intended for exploring which metrics are most useful for documenting data quality, the current version of the tool is sufficiently robust for use by forensic scientists with an interest in data analysis and access to a fast desktop computer.

Keywords: Electropherograms / Exploratory data analysis / Quality assessment / Resolution
DOI 10.1002/elps.200600116

User manual (127 pages) available for download from STRBase

Multiplex_QA Overview








The graph shows quality metrics (Y-axis) over time (X-axis) for three lanes: Lane-03, Lane-05, and Lane-01. The data points are color-coded: X (Unknowns), G (Blanks), and L (Ladders).

- Research tool that provides quality metrics to review instrument performance over time (e.g., examines resolution and sensitivity using internal size standard peaks)
- Runs with Microsoft Excel macros. Requires STR data to be converted with NCBI's BatchExtract program into numerical form.

Available for download from STRBase:
<http://www.cstl.nist.gov/biotech/strbase/software.htm>

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Leading the Way in Forensic DNA...

						
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Thank you for your attention...

Our team publications and presentations are available at:
<http://www.cstl.nist.gov/biotech/strbase/NISTpub.htm>

Questions?





See also <http://www.dna.gov/research/nist>
<http://www.cstl.nist.gov/biotech/strbase>
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