

# 2012 PulseNet PFGE Laboratory Update: Everything Old Is New Again

Grand Hyatt, Atlanta, CA  
August 27<sup>th</sup> – 30<sup>th</sup>, 2012

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Enteric Diseases Laboratory Branch, CDC

# OBJECTIVES

- ❑ **Identify how pattern quality may be negatively affected by variations in enzyme quality**
- ❑ **Recognize how gel quality may be negatively impacted by variations in agarose quality**
- ❑ **Describe the network-wide impact of reagent quality on gel and image quality**

# PFGE Reference Laboratory

- ❑ Efrain Ribot, PhD – Team Lead
- ❑ Molly Freeman, PhD – Unit Chief
- ❑ Maurice Curtis, BS
- ❑ Jessica Halpin, MS



# PFGE Reference Laboratory – What do we do?

## □ In the lab...

- Test difficult to type isolates, triage during outbreaks
- PFGE training – annual lab training in Spring
- PFGE protocol development, reagent testing, validation studies
- Provide technical support, customer service
- Projects – epi studies, WGS studies, etc...
- PFGE troubleshooting – [PFGE@cdc.gov](mailto:PFGE@cdc.gov)



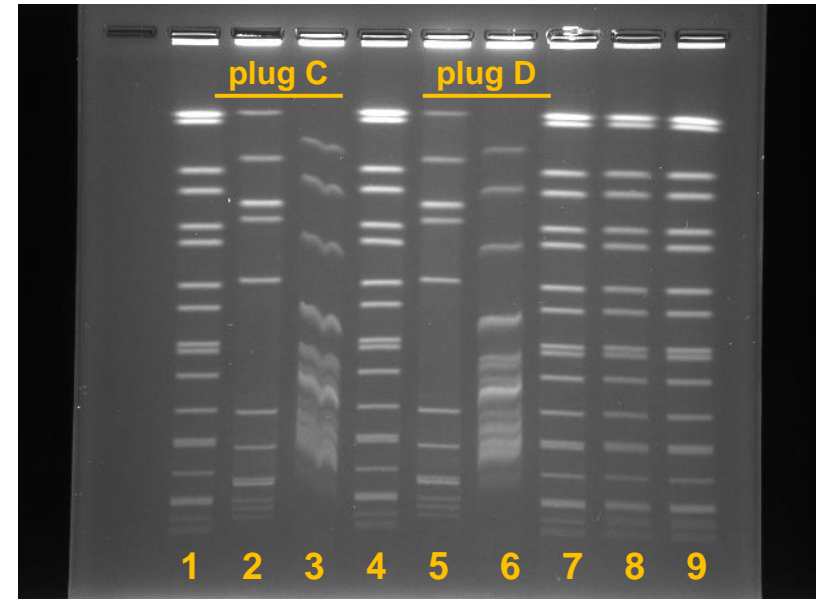
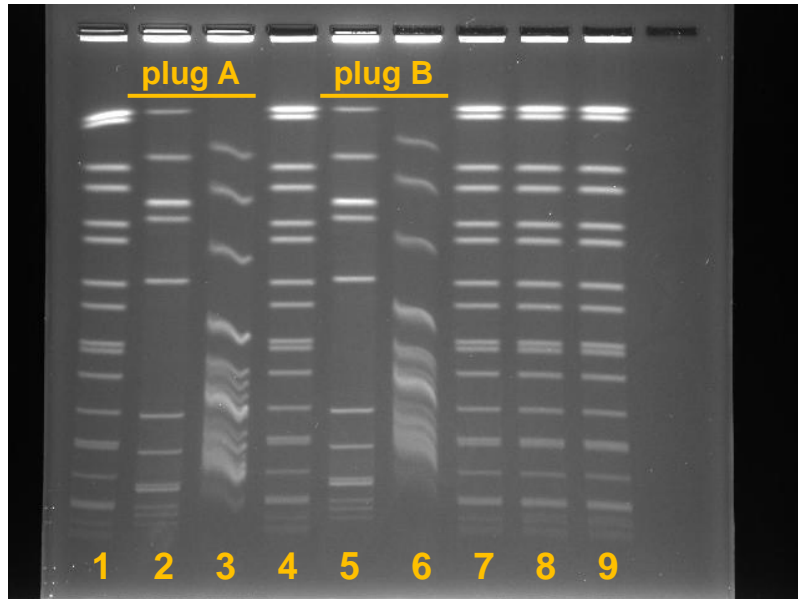
# Troubleshooting approaches

- ❑ **Study the protocol**
- ❑ **What has changed since the last “good” gel?**
  - Equipment, reagent, person, isolate / strain, etc...
  - Change in lab procedures / new SOPs, new vendor, contamination / decontamination event, etc...
- ❑ **Examine the gel closely**
  - Is the problem:
    - in all lanes, only the isolates, only the standards, only one enzyme?
    - apparent with multiple lab personnel?
    - apparent with multiple enzymes or multiple organisms?
    - associated with a specific mapper?
- ❑ **Maintain notebook of troubleshooting findings, gels**
  - ➔ **What is in common and what has changed?**

“Wavy” bands

# **CASE STUDY #1**

# *Listeria* Gels – NH Lab, Winter 2012



- same isolates in lanes 2, 3, 5 and 6
- different techs, different plugs, different master mixes
- *Ascl* (NEB) – lanes 2 and 5
- *Apal* (NEB lot# 0421103) – lanes 3 and 6, 125units (2.5 $\mu$ L of 50u/ $\mu$ l)
- BSA included

# What's going on?

## ❑ **Wavy bands:**

- Only in slices digested with *Apal*
- Occur in presence of BSA
- Present for multiple lab personnel

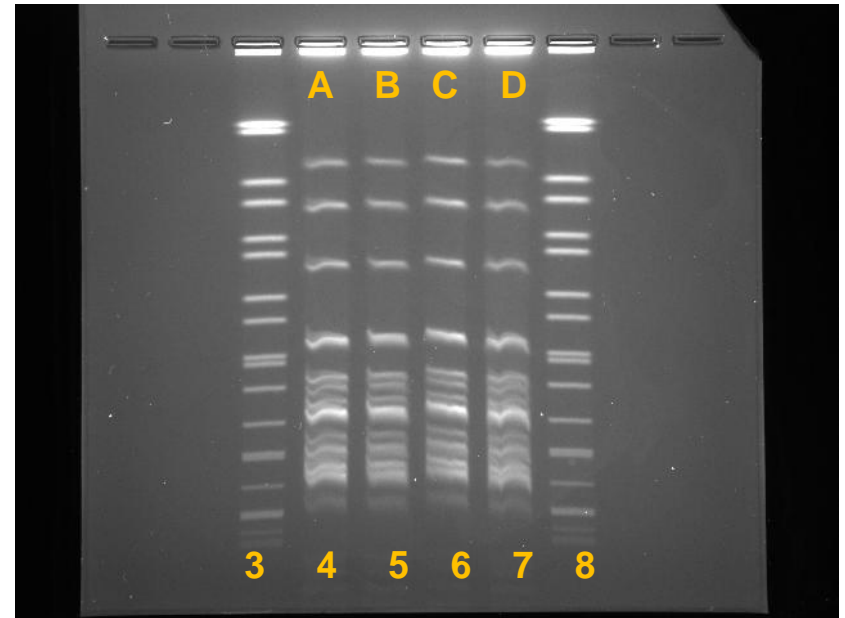
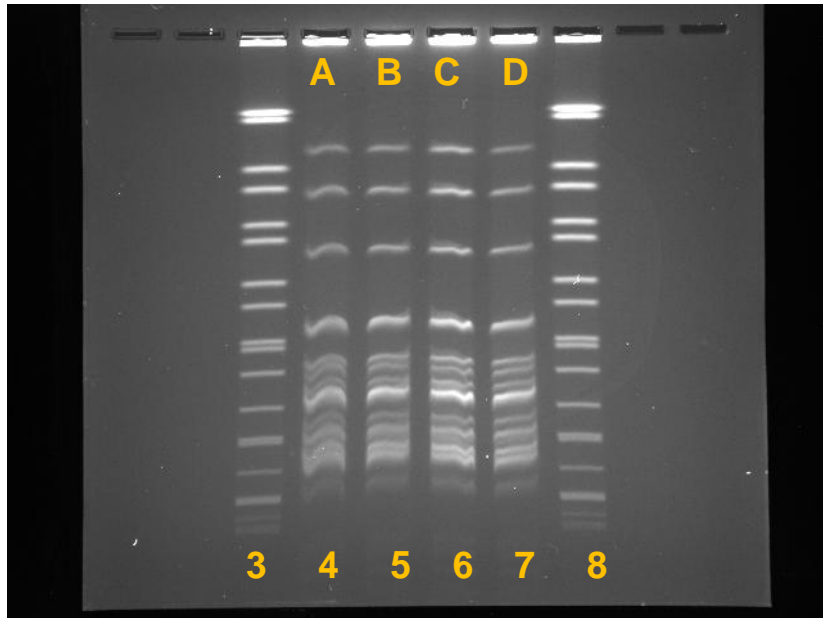
## ❑ **Ideas:**

- Damaged plug slice – bad spatula, bad razor, alcohol wipes, etc...
- *Apal* contaminant
- Change in PulseNet protocol

➔ **What is in common and what has changed?**

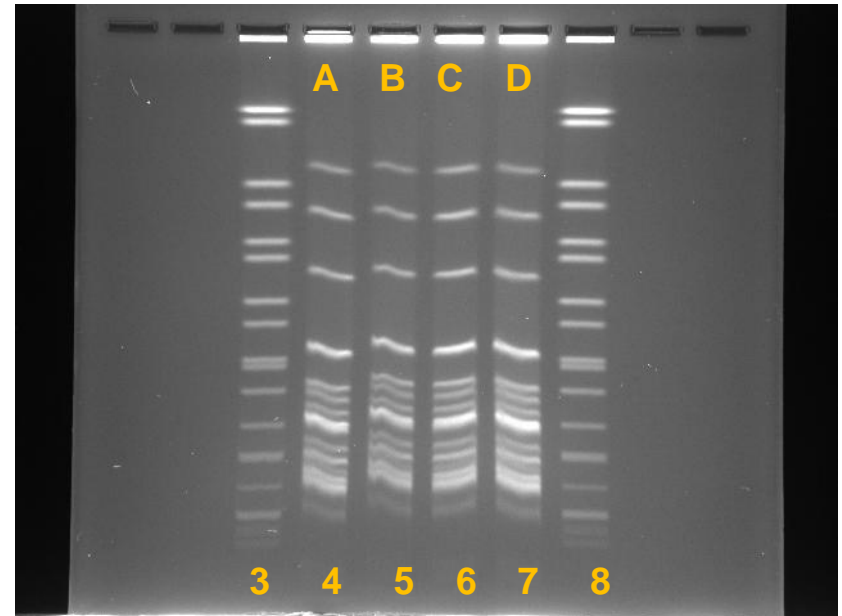
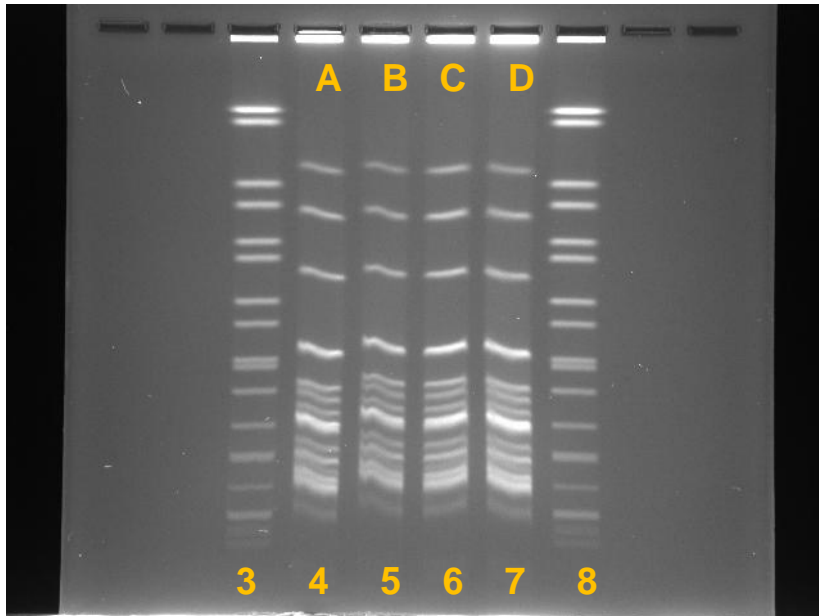


## *Listeria* Gels – damaged plug slices?



- 
- different razors, petri dishes, spatulas
  - same scientist
  - same 4 plugs from previous slide
  - *Apal* (NEB lot# 0421103) – lanes 4 – 7, 125units (2.5 $\mu$ L)
  - same master mix, BSA included

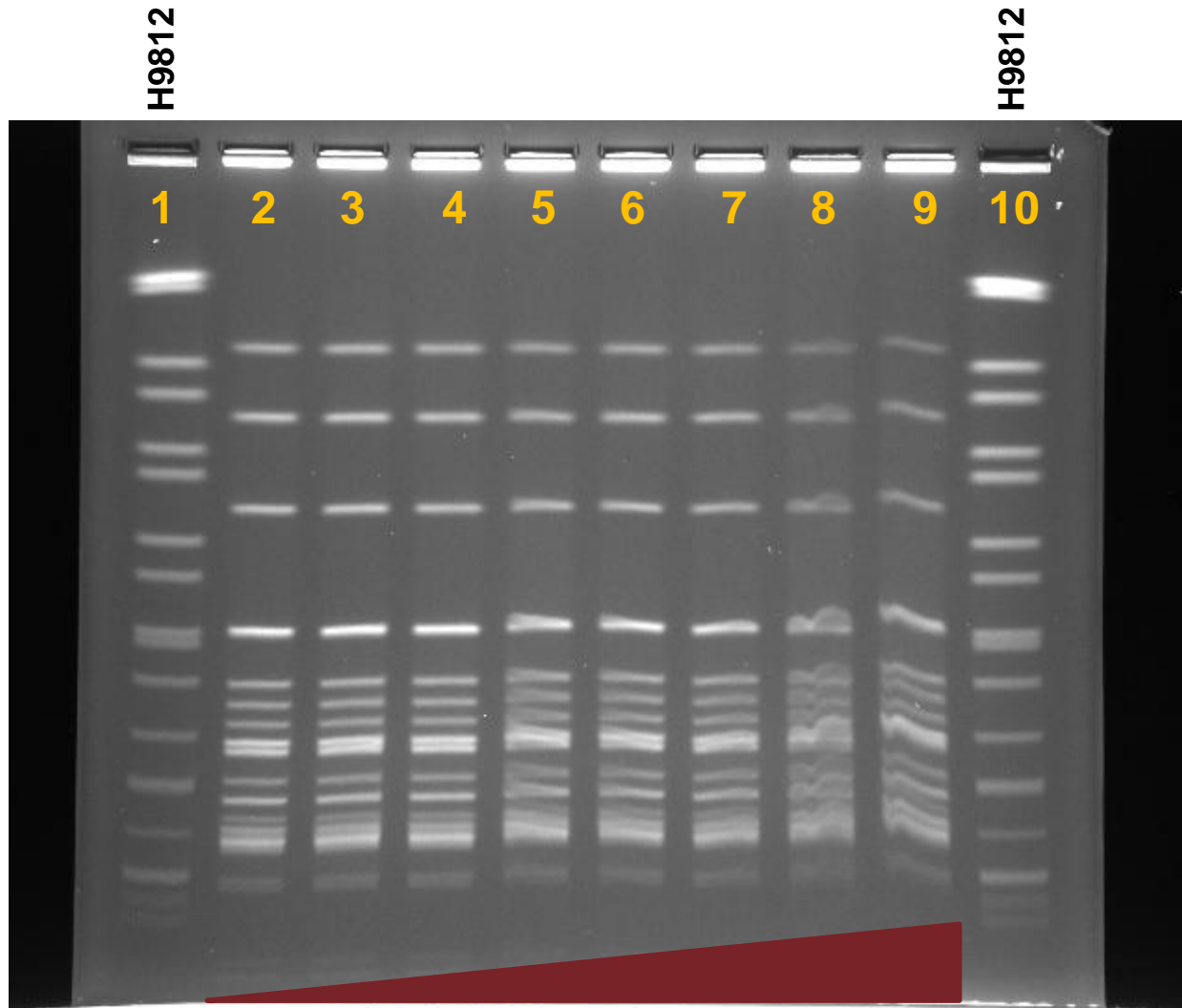
## *Listeria* gels – *Apal* enzyme?



- same scientist
- same 4 plugs as gels from previous slides
- • *Apal* (NEB lot# 0431110) – lanes 4 – 7, 125units (2.5 $\mu$ L)
- same master mix for both gels BSA added

**“The *Apal* product manager thought it may be protein binding causing the issue.”**

# *Listeria* and *Apal* – protein binding?



<i>Apal</i> (units):	25	50	75	100	125	150	200	250
<i>Apal</i> (uL):	0.5	1	1.5	2	2.5	3	4	5

## Result:

- band distortion directly correlated with units of enzyme added to master mix

# What's going on?

## ❑ Ideas:

- ~~Damaged plug slice – bad spatula, bad razor, alcohol wipes, etc...~~
- *Apal* contaminant / protein binding
  - wavy bands appeared with **new lots** in January
  - wavy bands directly correlated with increasing units of enzyme
- Change in PulseNet protocol
  - carrying out pre-digestion step (1X NEBuffer 4) and post-digestion step (0.5X TBE) had a small improvement (data not shown)
  - *Ascl*: 40 units / sample, 2hrs at 37°C
  - *Apal*: 50 units / sample, 2hrs at 25°C
  - ***Ascl*: 25 units / sample, 2hrs at 37°C (new recommendation)**
  - ***Apal*: 25 units / sample, 2hrs at 25°C (new recommendation)**

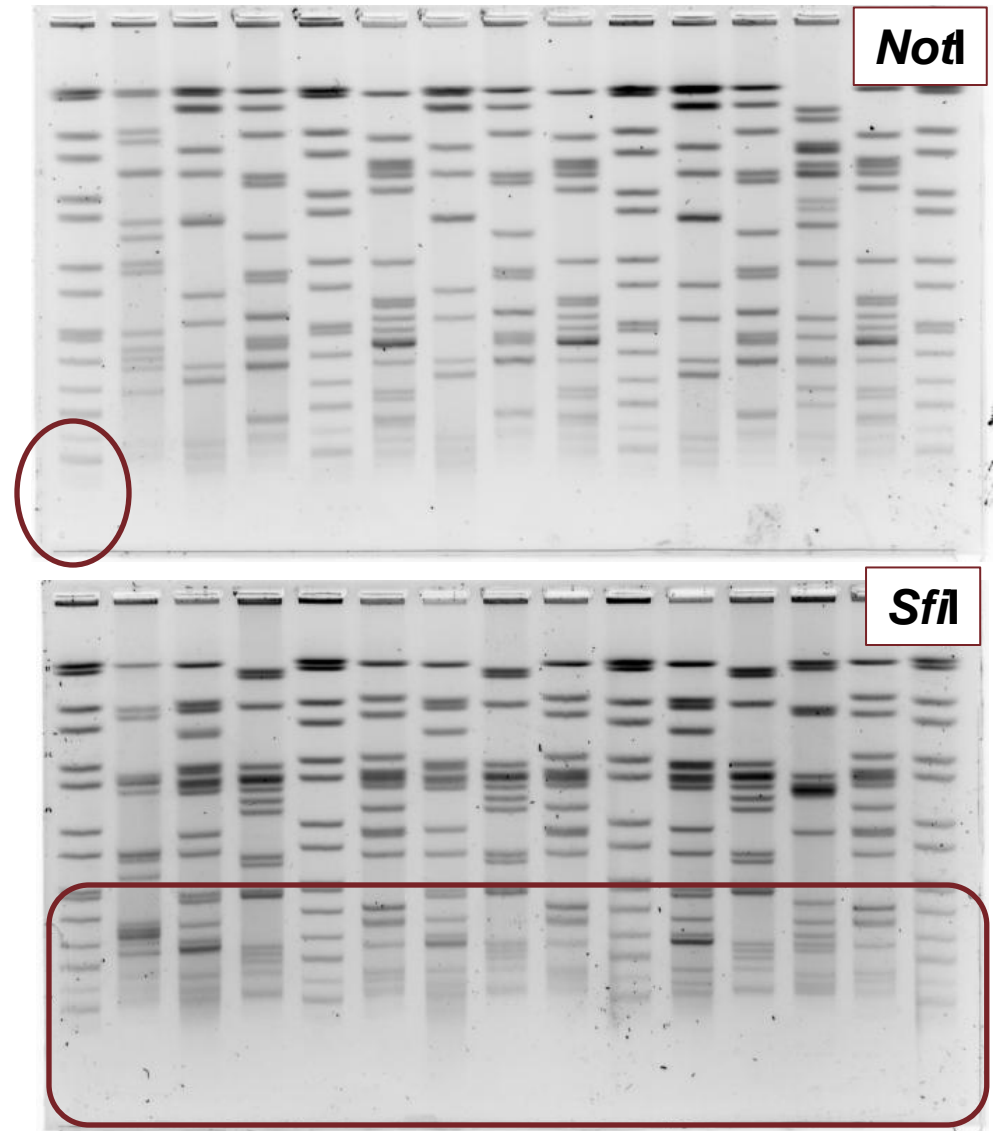
➔ **What is in common and what has changed?**

Diffuse, faint and/or disappearing low molecular weight bands

## **CASE STUDY #2**

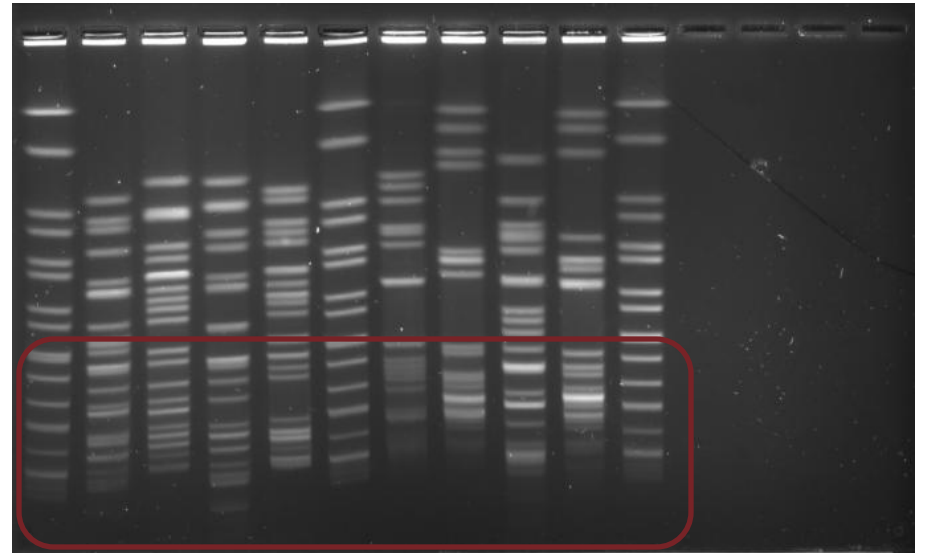
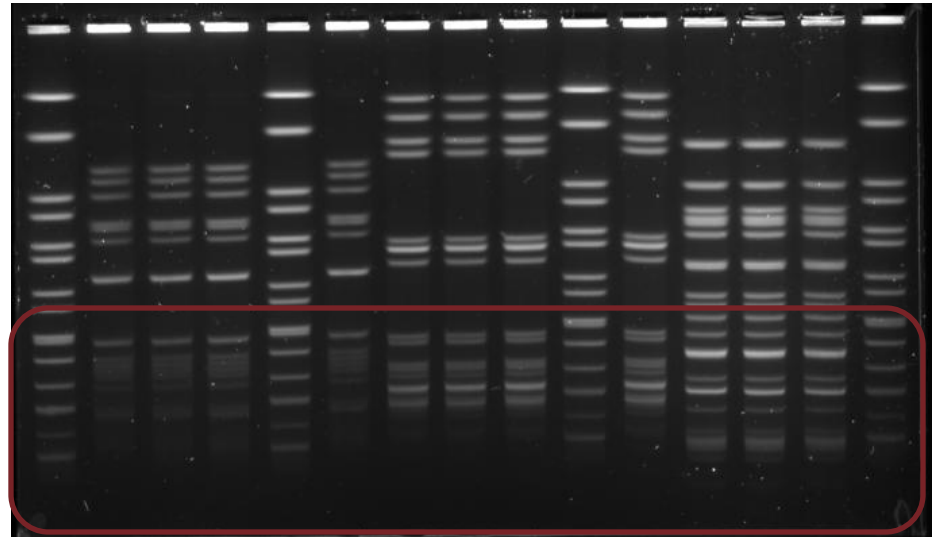
# “Fuzzy” Gels – CDC, Fall 2011

- SeaKem Gold (SKG)  
lot # 242048
- *Vibrio parahaemolyticus*  
conditions (10s – 35s)
- run time = 20 hours
- **could not be analyzed**
  - 1 hour short
  - bottom bands of standard  
(28.8 and 20.5kb) missing
  - some “tracking”



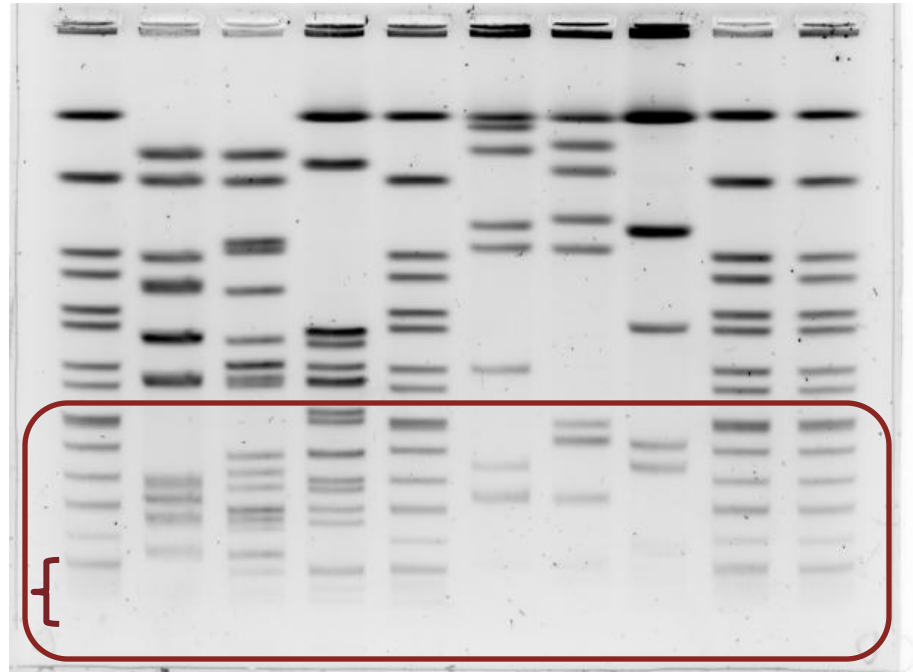
# “Fuzzy” Gels – CDC, Fall 2011

- SeaKem Gold (SKG)  
lot # 242048
- *E. coli* O157 conditions  
(2.16s – 54.17s)
- run time = 20.25 hours
- **could not be analyzed**
  - 1 hour short
  - bottom bands of standard  
(28.8 and 20.5kb) missing
  - some “tracking”



# “Fuzzy” Gels – CDC, Fall 2011

- SeaKem Gold (SKG)  
lot # 242048
- *Salmonella* conditions  
(2.16s – 63.8s)
- run time = 18 hours
- **could not be analyzed**
  - ~30 minutes short
  - bottom bands of standard  
(28.8 and 20.5kb)  
indistinct
  - no “tracking”
  - stained with GelRed





# What's going on?

## ❑ Indistinct, diffuse low-molecular weight bands

- Observed in multiple PulseNet protocols
  - *E. coli*, *Salmonella*, *Vibrios*, etc...
  - *Xba*I, *Bln*I, *Sfi*I, *Not*I, etc...
  - Ethidium bromide, GelRed
- Observed for different technicians in multiple labs

## ❑ Ideas

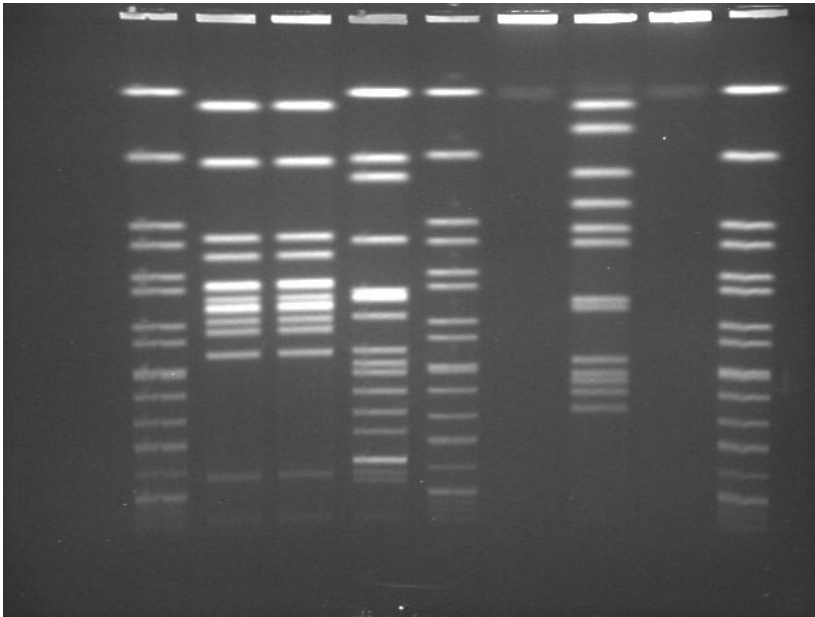
- Contaminant in water
- Contaminant or formulation change in TBE of one of the components
- Contaminant or formulation change in agarose

➔ **What is in common and what has changed?**

# Comparison of lots of SKG (Lonza)

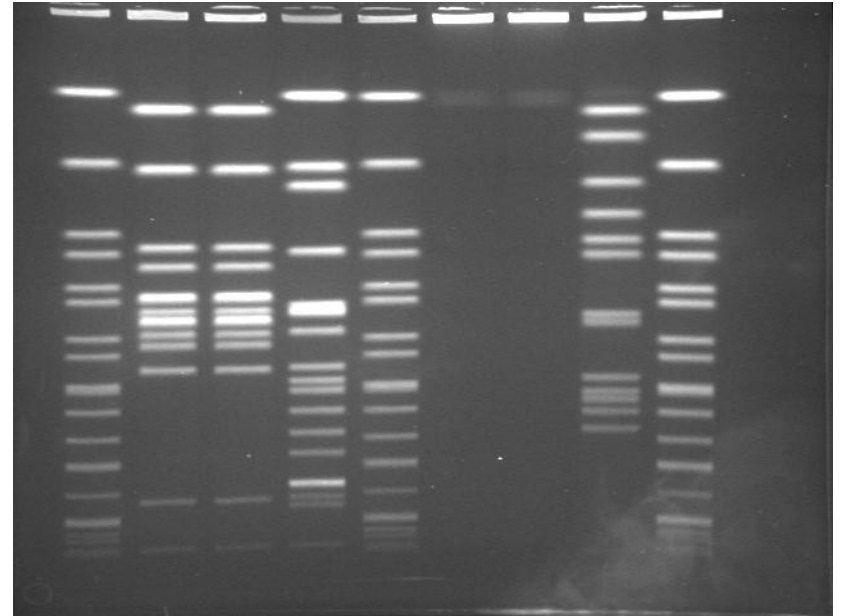
***Salmonella* conditions (2.16s – 63.8s)**

SKG lot# 252573



- SKG agarose clumped in the flask before microwaving, took longer to melt
- length is ~1hr short
- lower MW bands are faint and diffuse
- whole image is not as sharp as usual

SKG lot# 209453



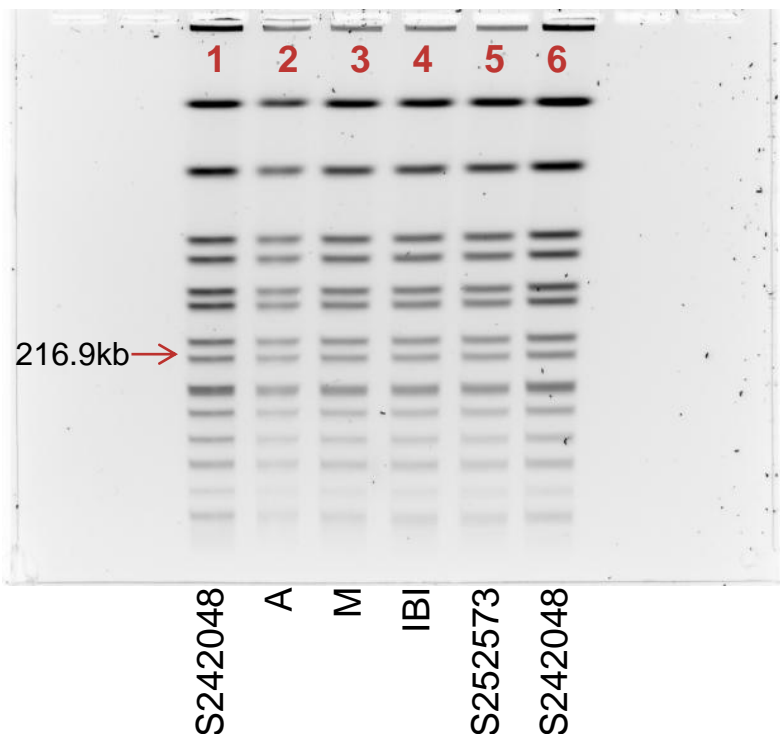
- SKG agarose appearance and melting was normal
- good length
- all bands are crisp and distinct

# Agarose Testing

- ❑ **Hypothesis:** Poor gel quality is due to bad lot(s) of SeaKem Gold used for running agarose
  
- ❑ **Plug agarose**
  - Amresco LF™, Bio-Rad Megabase, IBI (not validated), SKG lot # 242048, SKG lot # 252573
  - All plugs made on the same day with the same suspension of H9812 cells
  
- ❑ **Running agarose**
  - Amresco LF™, Bio-Rad Megabase, IBI (not validated), SKG lot # 242048, SKG lot # 252573, SKG Lot# 169896
  - All gels run by the same scientists using the same instrument for 18 hours, 14°C, *Salmonella* switch times

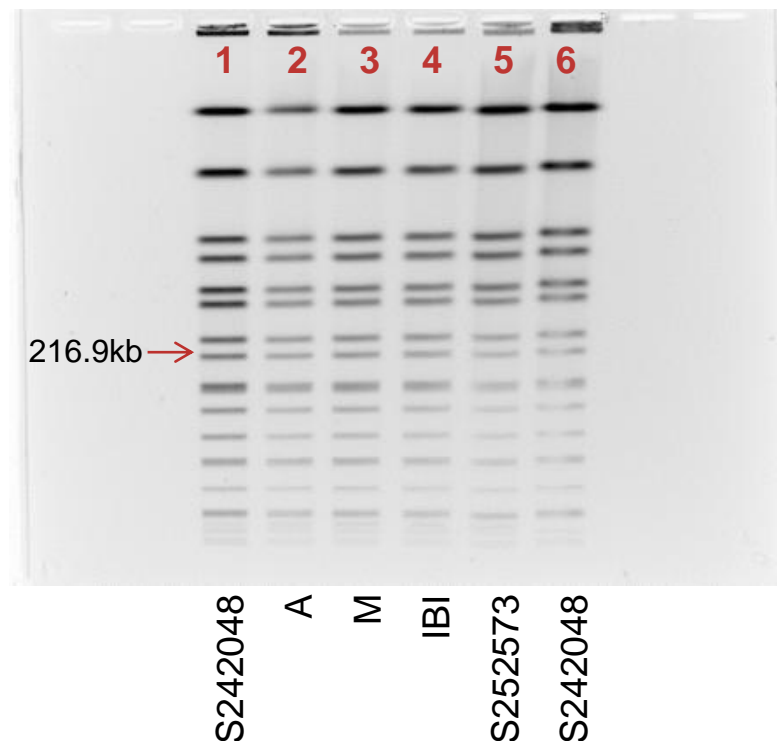
# Comparison of brand and lots of plug and running agarose

SKG lot # 252573



- indistinct, faint bands observed regardless of plug agarose used

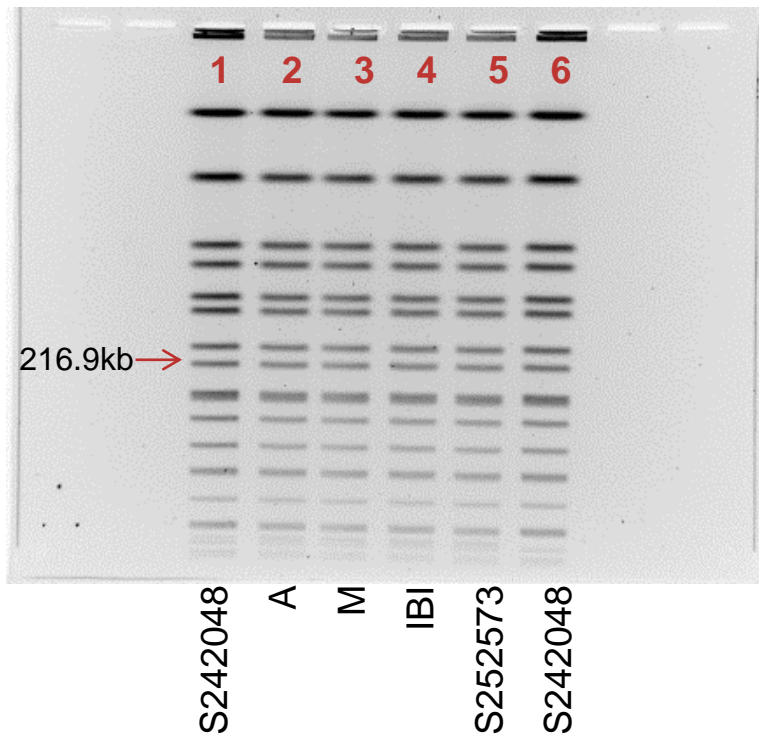
Amresco LF™ lot# 2899B228



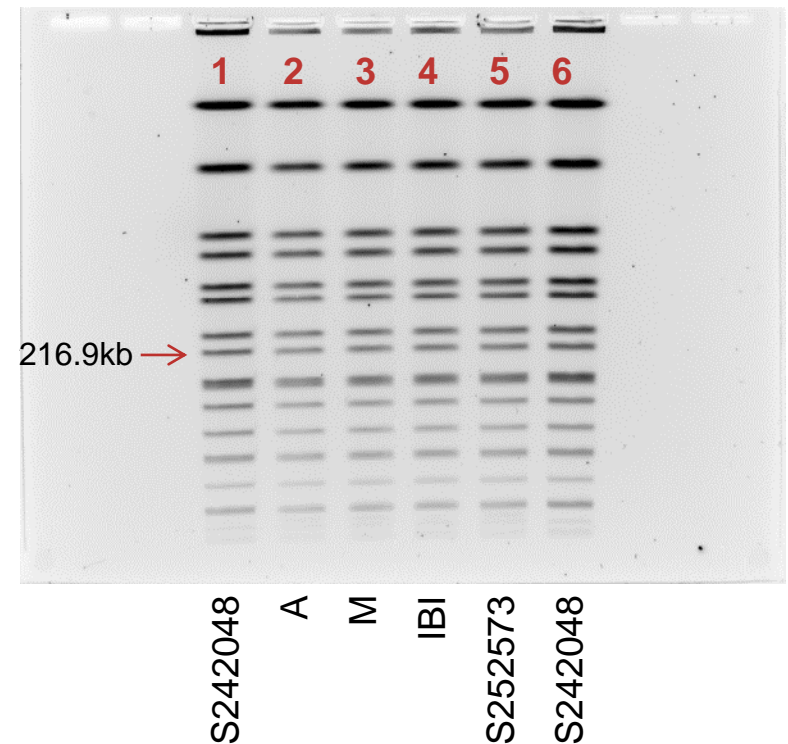
- crisp bands at all molecular weights, all plug agaroses
- some “tracking” observed

# Comparison of brand and lots of plug and running agarose

SKG lot # 169896 – “old”



SKG lot # 267744 – “replacement”



- crisp bands at all molecular weights regardless of plug agarose used

# Conclusions and Recommendations – SeaKem Gold (Lonza)

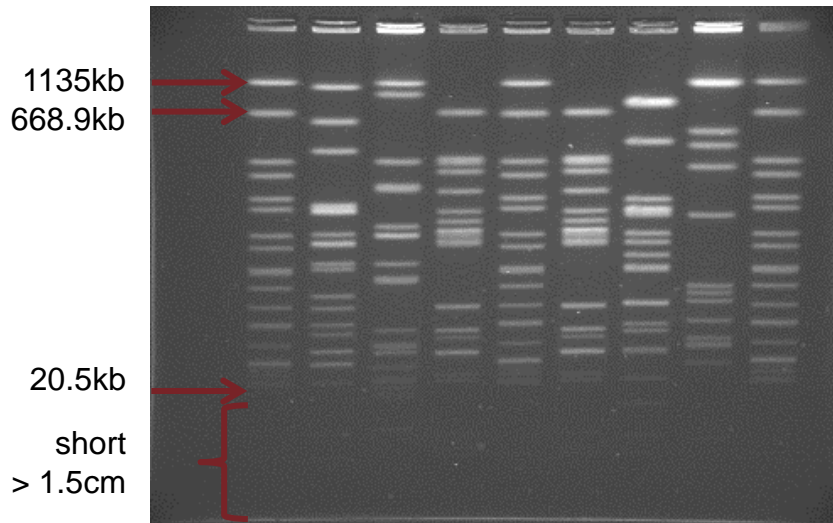
- ❑ **SeaKem Gold (SKG) lots #s 242048, 252573, 255693, and 262031 should not be used for plug or running agarose**
  - All 4 lots were linked to a single “bulk batch” at Lonza
  - Bulk batch and individual lots passed all QA/QC tests at Lonza
    - ✓ **Smallest band on QA/QC gels was 200kb**
  - No change in production process or formulation of SKG was identified
- ❑ **Please contact Lonza for a possible replacement**
  - Will replace at their discretion
  - New, replacement lots produce good quality gels
- ❑ **Continue to monitor agarose characteristics and gel quality**

Short gels / long run times, bad normalization

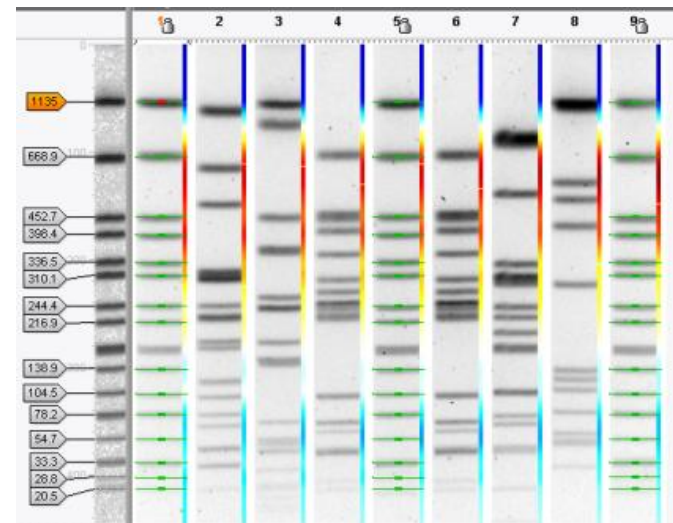
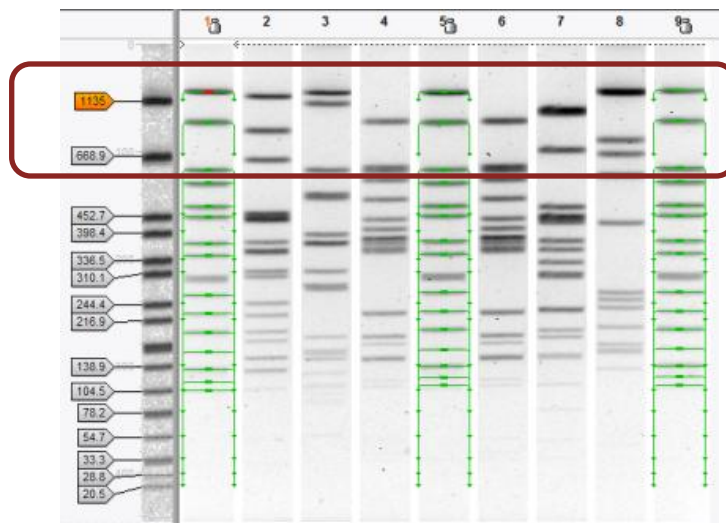
## **CASE STUDY #3**



# Characterization of poor quality gels run with Megabase (Bio-Rad)

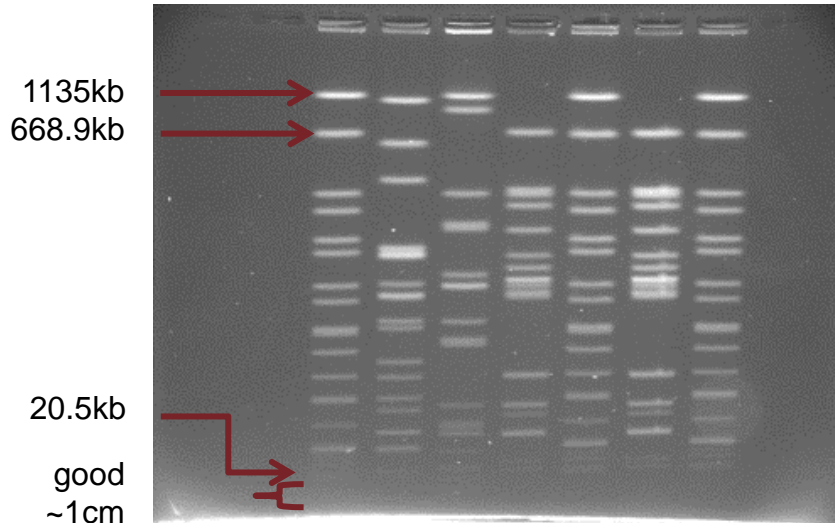


- Megabase lot# 45100031
- *Salmonella* conditions (2.16s – 63.8s)
- run time – 18.25 hours, 4hrs short
- failed normalization

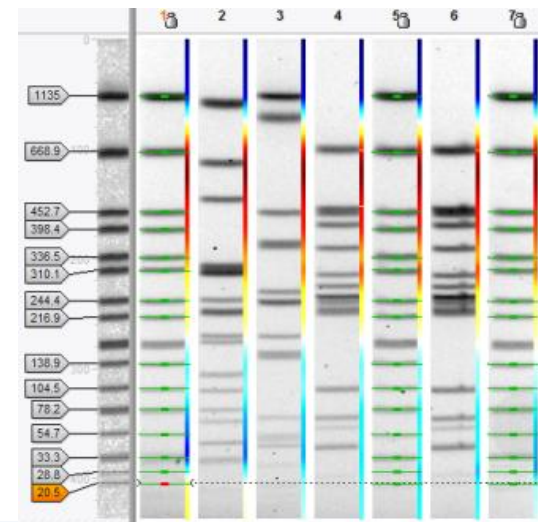
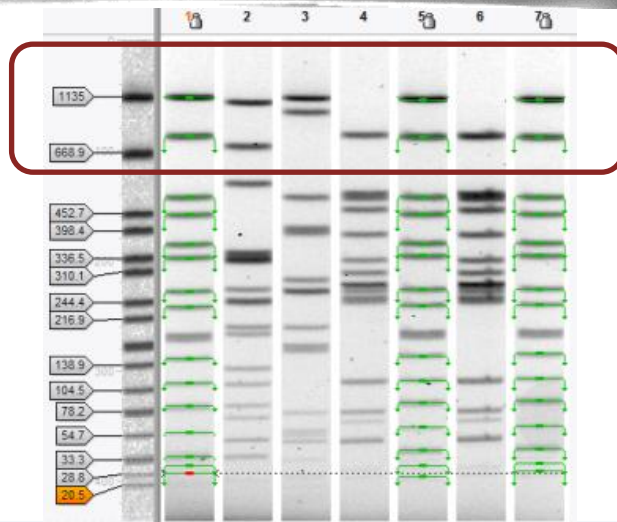




# Characterization of poor quality gels run with Megabase (Bio-Rad)



- Megabase Lot# 45100031
- *Salmonella* conditions (2.16s – 63.8s)
- run time – 22.25 hours, unreasonably long time for proper length
- failed normalization



# What's going on?

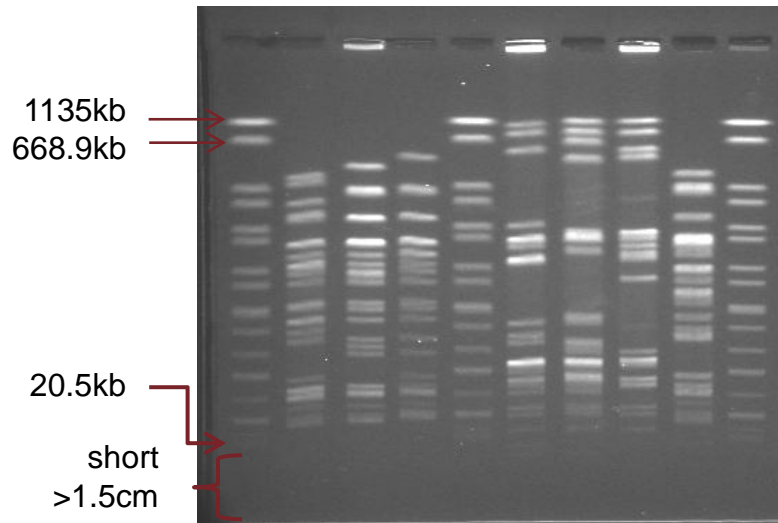
- ❑ **Failing normalization – gap between 1135kb and 668.9kb bands was smaller than expected**
- ❑ **Increased running time for bottom band of standard (20.5kb) to reach 1cm from bottom of gel**
  - Observed in multiple PulseNet protocols
  - Observed for different technicians in multiple labs on multiple instruments
  - Additional run time did not improve normalization
- ❑ **Ideas**
  - Machine malfunction
  - Contaminant or formulation change in agarose

**→ What is in common and what has changed?**

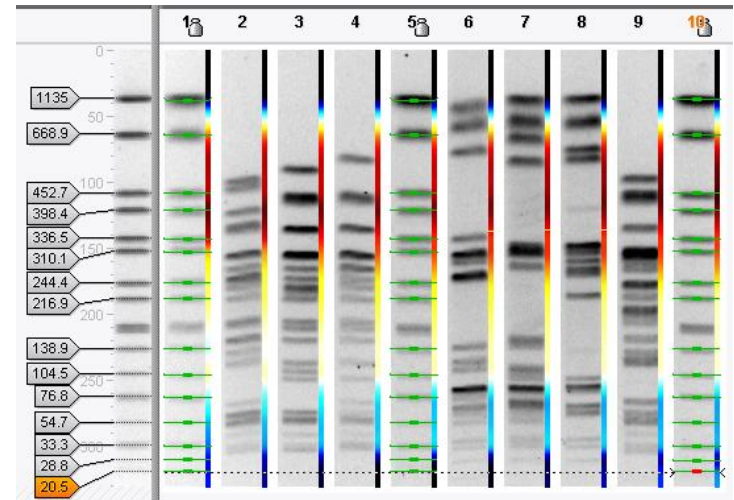
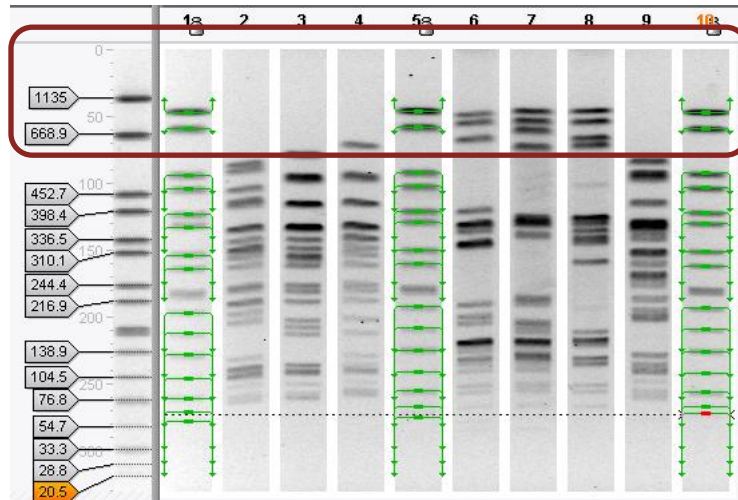
# Agarose Testing

- ❑ **Hypothesis:** Poor gel quality is due to bad lot(s) of Megabase, performing similarly to “old” formulation
  
- ❑ **Running agarose**
  - Megabase, “old” formulation
  - Megabase 037, “test lot”
  - Megabase lot# 45100031, “new” formulation

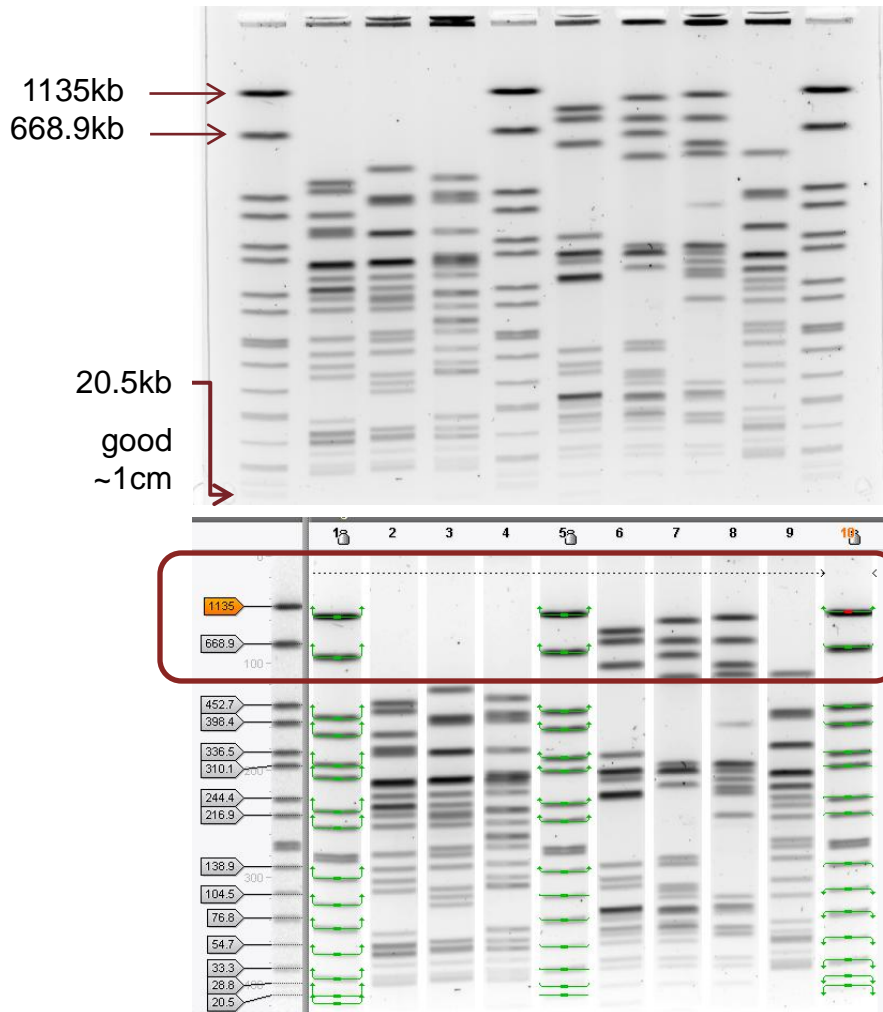
# Characterization of gels run with “old” formulation of Megabase (Bio-Rad)



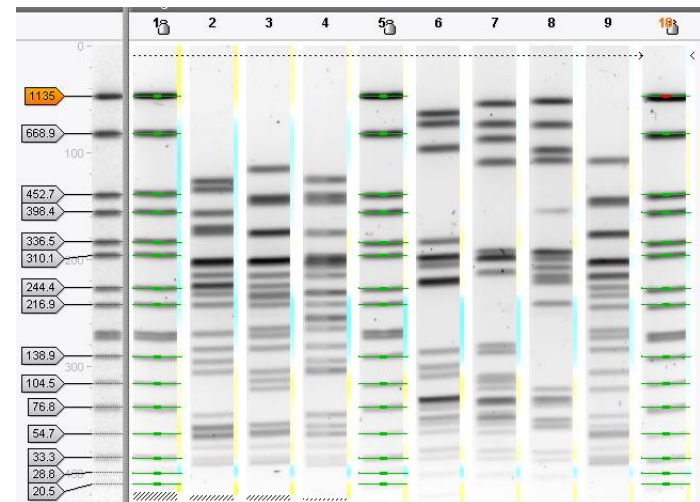
- Megabase – “old” formula
- tested in 2008
- *E. coli* switch times (2.16s – 54.17s)
- run time = 19 hours, ~2hrs short
- failed normalization



# Characterization of gels run with “new” formulation of Megabase (Bio-Rad)

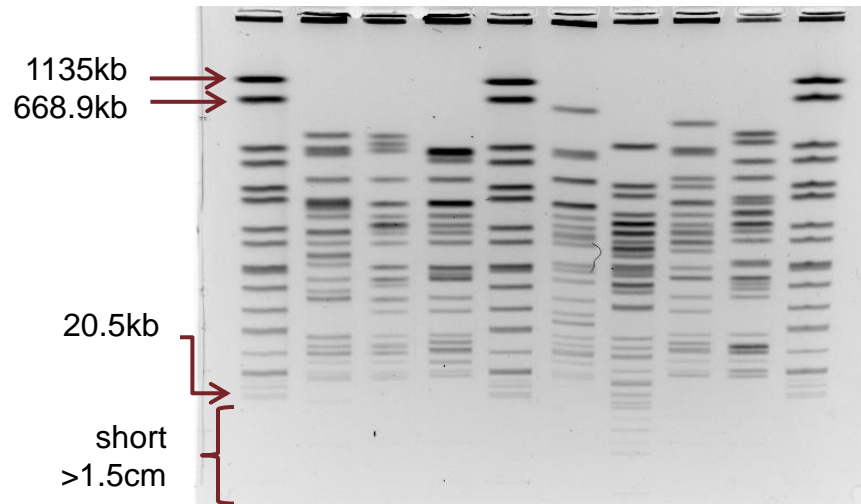


- Megabase – 037 “test lot”
- tested in 2009
- *E. coli* switch times (2.16s – 54.17s)
- run time = 18.25 hours; good length in a reasonable run time
- good normalization

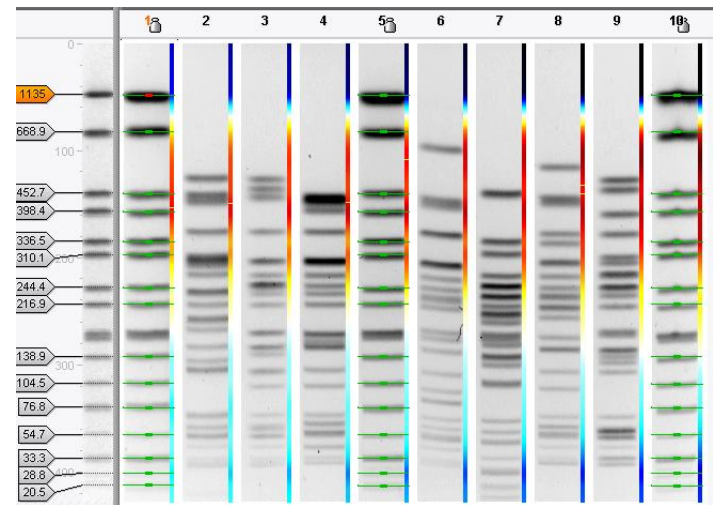
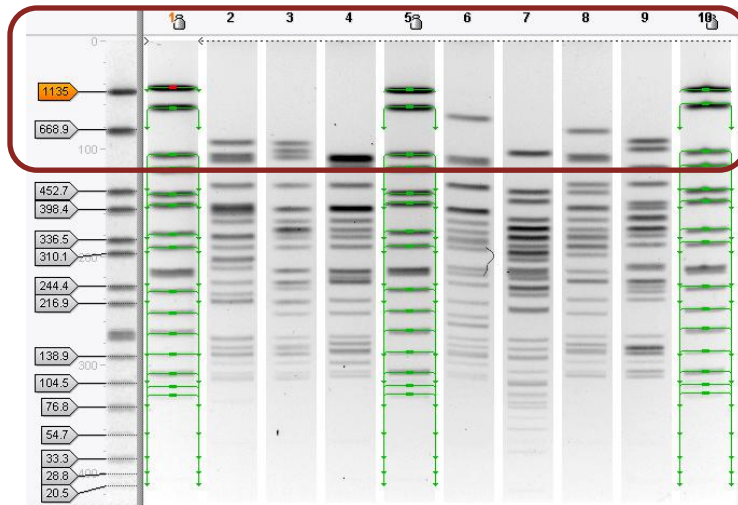




# Characterization of gels run with “new” formulation of Megabase (Bio-Rad)



- Megabase lot# 45100031
- *E. coli* switch times (2.16s – 54.17s)
- run time = 18 hours, >2 hours short
- failing normalization



# Conclusions and Recommendations – Megabase (Bio-Rad)

- ❑ **Megabase from all lots should not be used for plug or running agarose**
  - No change in production process or formulation of Megabase was identified, low end of specification range
  - PulseNet protocols will be updated to remove language suggesting Megabase as an acceptable alternative to SKG
  - Additional lots or new formulations will be tested and recommendations regarding using Megabase in PulseNet protocols revised
- ❑ **Continue to monitor agarose characteristics and gel quality**

# Sources of Variation within Standardized Protocols

## ❑ Interpretation of instructions

- legacy – “but we’ve always done it that way...”
- protocol drift vs. protocol shift

## ❑ Reagents

- lot-to-lot from one vendor
- vendor-to-vendor
- in-house prepared buffers vs. commercial, purchased buffers
- water quality

## ❑ PFGE equipment

- models, electrical supply, etc...

## ❑ Image acquisition

- staining, imaging system, camera, etc...



# Conclusions

- ❑ **Communication was key to identifying the cause of poor quality gels in these examples**
  - Reagents used across the entire network will impact data network-wide
  - Difficult to isolate variable or identify cause of poor quality gels if look at these phenomenon individually
  - Time to identify and resolve issues reduced due to cooperation and network-wide communication
- ❑ **Reagents will continue to have network-wide impact as production processes change, outsourcing, etc...**

# Acknowledgements

**All PulseNet participants at CDC,  
FDA, USDA, and in the State Public  
Health Laboratories**

**The findings and conclusions in this presentation are those of the author  
and do not necessarily represent the views of the Centers for Disease  
Control and Prevention**

**For more information please contact Centers for Disease Control and  
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