# Understanding and Evaluating Microbiological Sampling and Testing

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Science Staff Office of Public Health Science

# **Today's Presentation**

- Sampling methods
- Assessing sampling plans and testing methods
- Method validations and laboratory quality assurance
- FSIS testing programs, methods and pathogenspecific issues to consider
- Industry testing activities

# Sampling Methods

# **Sampling Methods**

- Destructive sampling grab, N60, excision
  - e.g., RTE, ground comminuted products, egg products, carcass excision
- Non-Destructive sampling
  - Typically chosen when destructive sampling not an option
  - Examples:
    - Carcass rinsates or sponge
    - Parts rinsate
    - Environmental sponge

## What is N60?

- N60 = number of samples (n) = 60
  - Multiple representative samples provides best option for detecting scattered contamination
  - Provides 95% confidence that no more than 5% of food pieces the size of each "n" in the entire lot are contaminated
- Keys to success
  - Must ensure that sampling is as representative as possible across the lot
  - Large composite "N60" samples typical need a larger test portion

## **Common Sampling Problems**

- Small sample or sampling method may not be ideal for detection
  - e.g., small swab device or environmental area sampled
- Sanitizer or residual antimicrobial chemicals might interfere with the test
  - Insufficient drip time prior to carcass sample collection
  - Excessive liquid carryover for parts sample collection
- Temperature abuse for the sample prior to testing
  - Holding under refrigeration for long periods allows competing bacteria to grow
  - Freezing can kill some pathogens (e.g., Campylobacter)

# **Assessing Sampling Plans**

## **Sampling Plans**

- All sampling plans have significant limitations
  - Relative rigor of the sampling program must be evaluated
- Best sampling plans provide the opportunity but no guarantee of detection

- i.e., scattered contamination is difficult to detect

- Frequent sampling and sampling multiple sites/time points provides a better opportunity for detection
  - Examples:
    - Multiple samples per day vs. once per month
    - N60 per lot vs. one grab sample per lot
- Does the type of sampling meet the intended need?
  - Destructive vs. non-destructive sampling

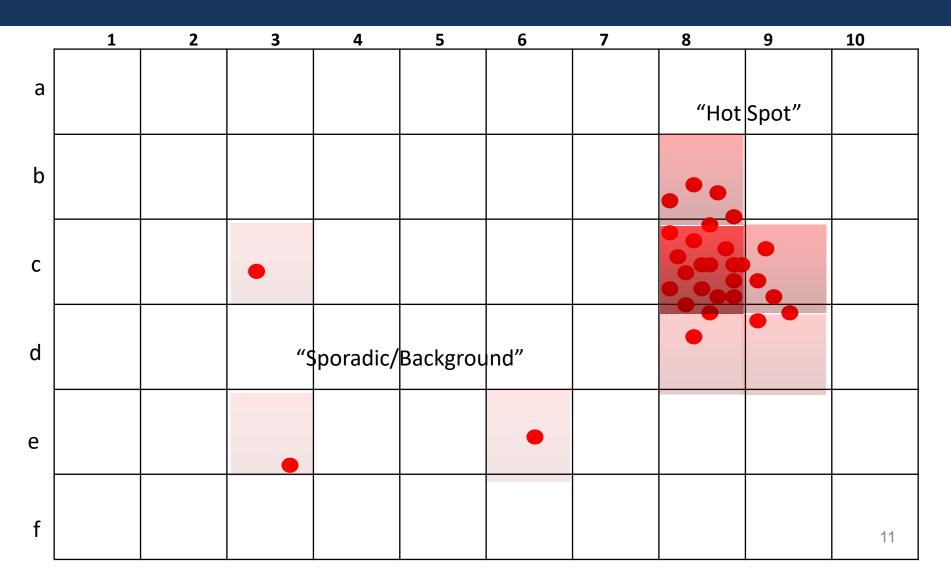
# **Sampling Plans**

- Statistical sampling plans assume
  - Uniform manufacturing conditions
  - Equal probability of contamination throughout the lot (homogeneous distribution)
  - Independent, random sampling (equal probability of sampling throughout the lot)

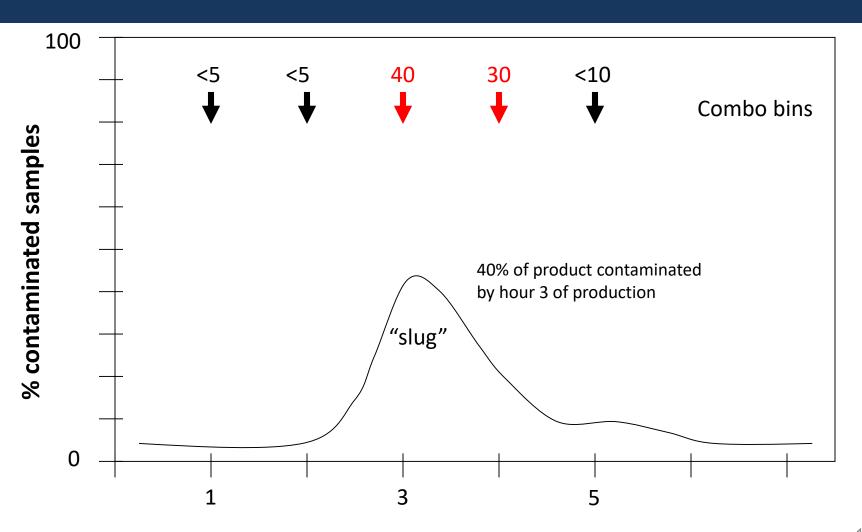
## Why are Pathogens Hard to Detect?

- They are typically not evenly distributed
- They occur at low levels
- They are often injured when found in the product
- Detection may be inhibited by material in the food product (food matrix)
  - Example: high amounts of fat may inhibit PCR assays; spices, salt, acidulants can affect isolation and detection

E. coli O157:H7 Contamination in a N60 Sampled Lot



## E. coli O157:H7 Contamination in Ground Beef



Time of production, hrs

# Assessing Testing Methods

**Key Players for Ensuring Robust Testing Methods** 

- The establishment that needs the testing
- The laboratory they hire
- The manufacturer of the screening test they use
- The organization validating the screening test

# **Steps in Detection Methods**

- Sample collection
- Sample preparation
- Enrichment for the pathogen
- Screening of the Pathogen
- Confirmation of the Pathogen

## **Considerations for Testing Methods**

- Is the method fit for the intended purpose of the analysis?
- Has the method been optimized and experimentally validated for sensitive detection of pathogens?
- Is the laboratory complying to the validated method protocol?

# **Assessing Fitness for Purpose**

- Is the test portion appropriate to meet the need?
- Is the method enrichment-based with the intent to detect the lowest possible numbers of stressed pathogen cells?
- Has the food matrix been validated for the method used?
- Are confirmation procedures appropriate for determining true negative samples?

## **The "Test Portion"**

- Laboratory sample preparation => "test portion"
  - "analytical unit" or "analytical portion"
  - Definition: the part of the "sample" that is actually tested by the laboratory
- The test portion determines the theoretical (*i.e.*, best possible) sensitivity of the test
  - e.g., 1 cell/test portion
  - 25-gram test portion: detecting 0.04 cells/gram is possible
  - 325-gram test portion: detecting 0.003 cells/gram is possible

## Enrichment

- Test portion is incubated 8-48 hours in a culture broth
  - Why?
    - Contamination levels are too low for detection without enrichment
    - Must grow to high levels so very small volumes have enough pathogen present for later detection steps
- Different pathogens require different enrichment media (broth)

One vs. two-stage enrichment

- Primary enrichment vs. secondary enrichment
  - Resuscitation vs. selective growth

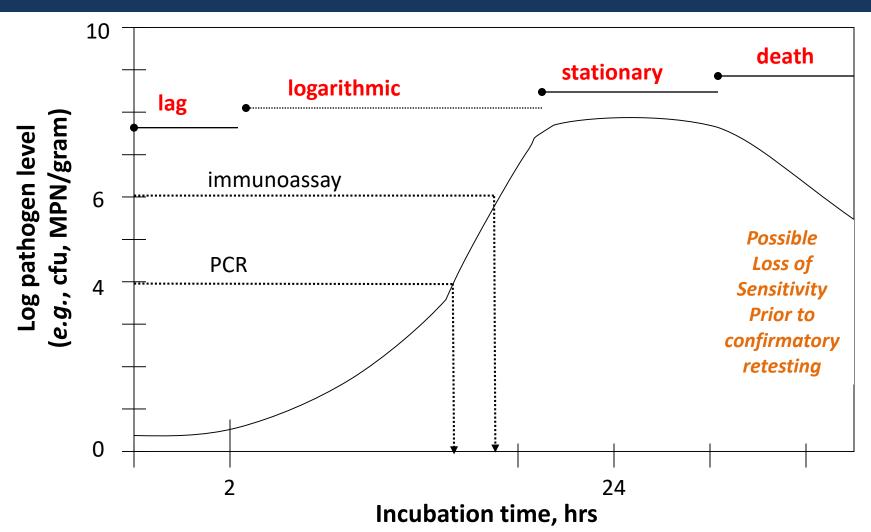
## **Considerations for Proper Enrichment**

 Resuscitation (lag phase) can require 2-3 hours before log-phase growth begins

Some samples support slower growth

- Has enrichment broth been tempered to warm temperature prior to incubation?
  - Particularly critical for large test portions or shorter incubation periods

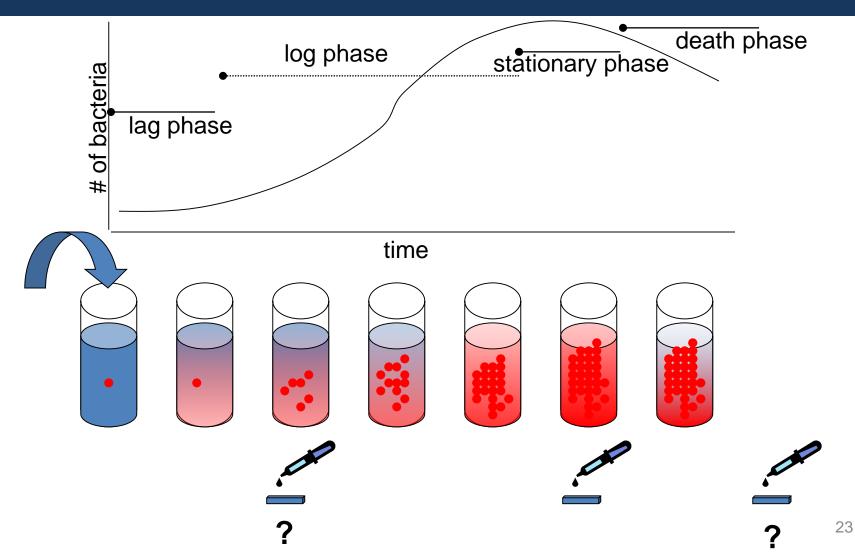
## **Pathogen Growth During Enrichment**



## **Enrichment Period**

- Different screening tests require different levels of enriched pathogen
- Shorter incubation periods (<15 hours) may warrant additional scrutiny of laboratory compliance to the validated protocol
- Has enrichment/screening combination been validated for a larger test portion?
  - Particular concern for large test portions incubated for shorter periods
  - *e.g.*, 375-gram test portion incubated for 8 hours
- Proposed incubations <8 hours may warrant OPHS review</li>

## **Role of Enrichment**



## **Confirmatory Testing**

- Non-culture confirmation (*e.g.*, PCR)
- Culture confirmation (*e.g.*, FSIS confirmation)
  - Plating the enrichment on selective and differential agar media
  - Immunomagnetic separation (IMS) necessary prior to plating for *E. coli* O157:H7 and non-O157 STECs
    - Suspect colonies = "presumptive positive"
  - Purification and confirmatory identification tests including:
    - Biochemical (*e.g.*, identifies "*E. coli*")
    - Serological (*e.g.*, identifies "O157" and "H7")
    - Genetic (*e.g.*, identifies "*stx*" = Shiga toxin genes)

## **Concerns for Confirmation**

- Do not re-sample the lot or sample reserve!
- Non-culture confirmation
  - Same considerations as the screening test
  - Used under validated conditions
  - Transport and storage of enrichment
- Culture confirmation- carefully assess!
  - Typically expect that methods comply with a validated procedure (*e.g.*, MLG, FDA-BAM, ISO)
  - Small changes can affect ability to recover pathogen of interest

## **Quantitative Testing**

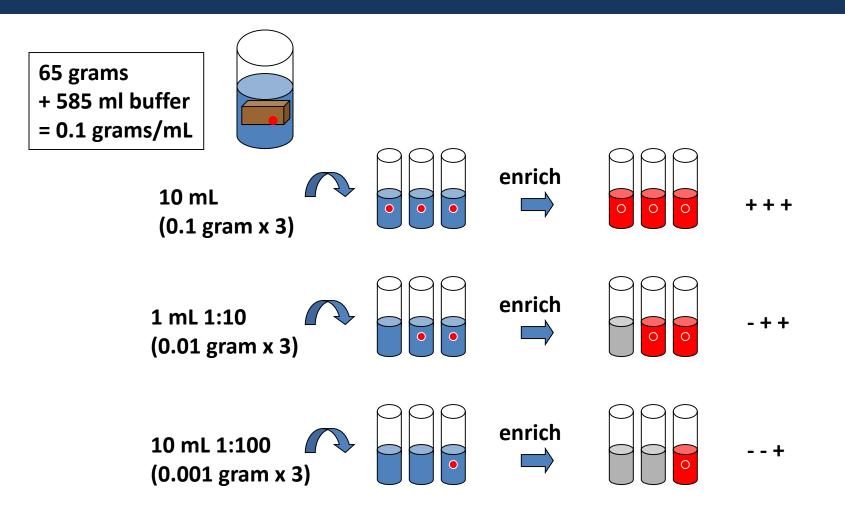
- Two options:
  - MPN
  - Direct plating

**NOTE:** Quantitative testing typically cannot accommodate larger test portions and provide the opportunity for detection that a qualitative test can provide

## Most Probable Number (MPN) Enumeration Analysis

- Traditional enrichment-based analyses are performed on three or more dilutions, each typically in triplicate, from a single sample homogenate (*i.e.*, MPN = method format, not a specific method per se)
- Advantages:
  - Better sensitivity (lower LOD) than direct plating
- Disadvantages:
  - Very resource intensive/expensive
- Application:
  - For quantifying low levels of pathogens (*e.g., Salmonella, E. coli* O157:H7, *L. monocytogenes*)

Quantitative Testing - MPN (most probable number)



## **Quantitative Testing - MPN (most probable number)**

|                     | 1 050 0    | <u></u>         | · · ·        |
|---------------------|------------|-----------------|--------------|
| Table 3. MPN Inde   |            | nfidence Limits | for Various  |
| Combinations of Pos |            |                 | Series Using |
| Inoculum Quantities | ,          |                 |              |
| Combination of      | MPN Index  | 95% Confide     |              |
| Positives           | per g (ml) | Lower           | Upper        |
| 0-0-0               | <3.0       |                 | 9.5          |
| 0-0-1               | 3.0        | 0.15            | 9.6          |
| 0-1-0               | 3.0        | 0.15            | 11.          |
| 0-1-1               | 6.1        | 1.2             | 18.          |
| 0-2-0               | 6.2        | 1.2             | 18.          |
| 0-3-0               | 9.4        | 3.6             | 38.          |
|                     |            |                 |              |
| 1-0-0               | 3.6        | 0.17            | 18.          |
| 1-0-1               | 7.2        | 1.3             | 18.          |
| 1-0-2               | 11.        | 3.6             | 38.          |
| 1-1-0               | 7.4        | 1.3             | 20.          |
| 1-1-1               | 11.        | 3.6             | 38.          |
| 1-2-0               | 11.        | 3.6             | 42.          |
| 1-2-1               | 15.        | 4.5             | 42.          |
| 1-3-0               | 16.        | 4.5             | 42.          |
| 2-0-0               | 9.2        | 1.4             | 38.          |
| 2-0-1               | 14.        | 3.6             | 42.          |
| 2-0-2               | 20.        | 4.5             | 42.          |
| 2-1-0               | 15.        | 3.7             | 42.          |
| 2-1-1               | 20.        | 4.5             | 42.          |
| 2-1-2               | 27.        | 8.7             | 94.          |
| 2-2-0               | 21.        | 4.5             | 42.          |
| 2-2-1               | 28.        | 8.7             | 94.          |
| 2-2-2               | 35.        | 8.7             | 94.          |
| 2-3-0               | 29.        | 8.7             | 94.          |
| 2-3-1               | 36.        | 8.7             | 94.          |
| 2 3 1               | 50.        | 0.7             | 54.          |
| 3-0-0               | 23.        | 4.6             | 94.          |
| 3-0-1               | 38.        | 8.7             | 110.         |
| 3-0-2               | 64.        | 17.             | 180.         |
| 3-1-0               | 43.        | 9.0             | 180.         |
| 3-1-1               | 75.        | 17.             | 200.         |
| 3-1-2               | 120.       | 37.             | 420.         |
| 3-1-3               | 160.       | 40.             | 420.         |
| 3-2-0               | 93.        | 18.             | 420.         |
| 3-2-1               | 150.       | 37.             | 420.         |
| 3-2-2               | 210.       | 40.             | 430.         |
| 3-2-3               | 290.       | 90.             | 1000.        |
| 3-3-0               | 240.       | 42.             | 1000.        |
| 3-3-1               | 460.       | 90.             | 2000.        |
| 3-3-2               | 1100.      | 180.            | 4100.        |
| 3-3-3               | >1100.     | 420.            |              |
|                     |            |                 |              |

Example: "3-2-1" = Y MPN/g (use MPN table to determine Y)

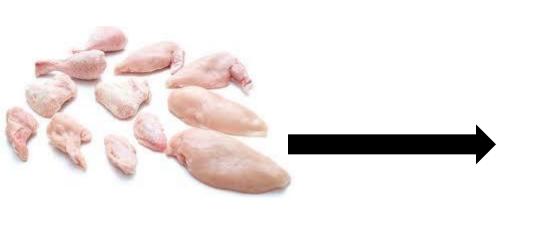
Total tested: 65 grams FSIS method

Level of Detection = <0.03 MPN/gram (0-0-0)

## **Direct Plating Enumeration Methods**

- Product is homogenized in diluent and small volume is directly dispensed onto agar media (*i.e.*, sometimes there is a 1-2 h "resuscitation" step, but enrichment is never used prior to plating)
- Advantages:
  - Allows easy inexpensive quantitative analysis
- Disadvantages:
  - Accommodates only a very small test portion
  - Higher LOD (*i.e.*, often 100 CFU/g) not suitable for detecting low levels of pathogens
- Application:
  - Expedient for higher level analytes (e.g., indicators, Campylobacter, S. aureus, C. perfringens, B. cereus)

## **Quantitative Testing: Direct Plating**





# Perform a 1:4 dilution using 325 grams of chicken

How much media do I need?

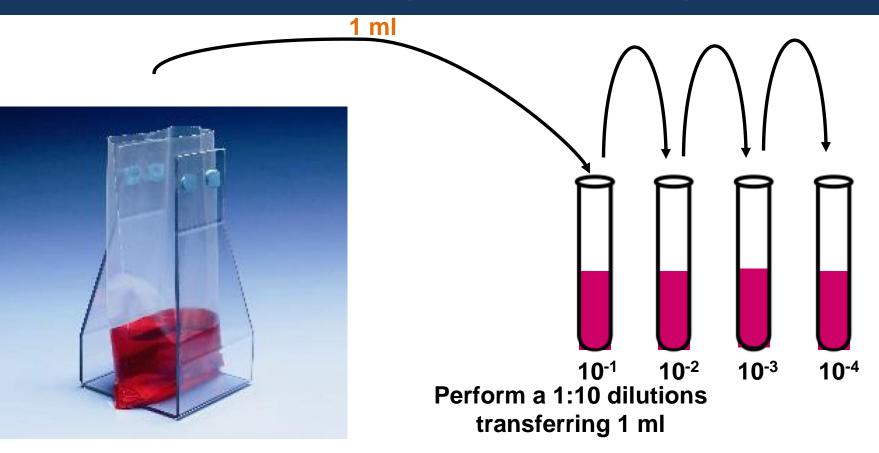
325 g x 4-fold dilution = 1300 g or ml (1 ml = 1g)

1300 ml – 325 g = 975 ml of media

Seal bag and shake

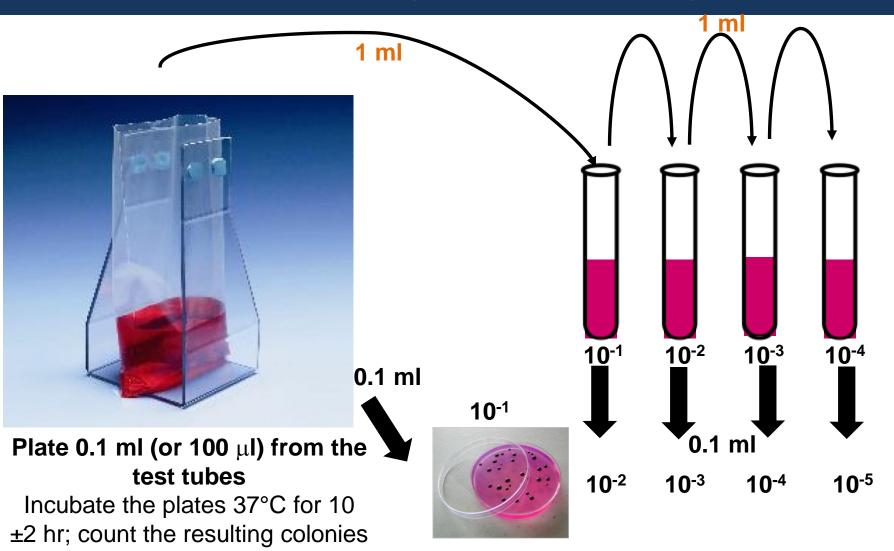
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## **Quantitative Testing: Direct Plating**

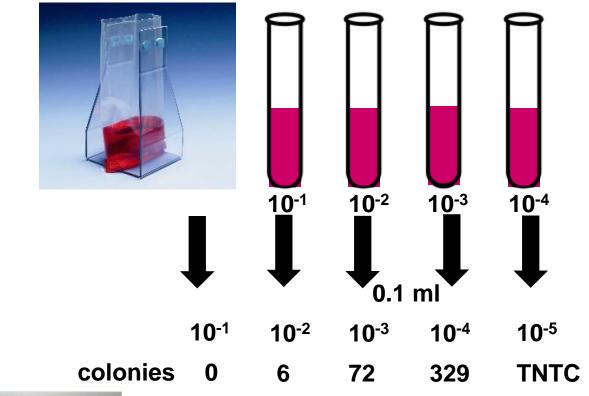


How much media do I need in each test tube? 1 ml x 10-fold dilution = 10 ml final volume 10 ml – 1 ml = 9 ml of media

## **Quantitative Testing: Direct Plating**



## **Quantitative Testing: Direct Plating**



 $4 \times 329 \div 1/10^{-4} = 1.32 \times 10^7 \text{ CFU/mI}$ 



# Method Validations

## Value of Validation

- Determines performance characteristics of the method in comparison to a gold standard (reference) method (*e.g.*, usually FSIS or FDA method)
- Independent evaluation provides credibility
- Rigor varies (multilab vs. single lab, # tests, etc)
- Still must consider fitness for purpose and how the method is applied
  - e.g., some AOAC-validated methods are not consistent with FSIS goals or Compliance Guidelines

# **Method Validation**

- Recognized independent method validation organizations:
  - Government: FSIS (MLG) and FDA (BAM)
  - AOAC International (U.S.A.)
    - AOAC Official Methods of Analysis (OMA) validations
    - AOAC-RI "Performance Tested Method" validations
  - AFNOR (France)
    - *e.g.*, bioMerieux Vitek biochemical confirmation tests
  - Others (ISO, MicroVal, NordVal, AENOR, etc.)
- However, past validations conducted by these organizations may not be relevant to larger test portions or other testing scenarios

### Food Safety and Inspection Service: Foodborne Pathogen Test Kits Validated by Independent Organizations

- FSIS maintains a list, updated quarterly, of methods that have been validated by independent organizations
  - <u>http://www.fsis.usda.gov/wps/portal/fsis/topics/regul</u> <u>atory-compliance/New-Technologies</u>
- None of the test kits listed are implicitly approved by USDA FSIS
  - A validated test kit must also be fit for purpose and appropriate for the specific application in a food safety program

**Process for Validating Qualitative Pathogen Methods** 

- Series of laboratory experiments using inoculated samples under controlled conditions
- Inoculate portions with pathogen strain at very low level where only 20-80% of samples are positive (*i.e.*, fractional recovery)
- Statistically compare percent of positive samples in alternative method to reference method (FSIS MLG)

# **Considerations for Validation Data**

- Was method compared to an appropriate reference method (*e.g.*, FSIS MLG)?
- If not performed by AOAC, AFNOR, etc., is supplemental validation data available?

May require additional scrutiny

# **Testing Method Specifications**

- <u>Sensitivity</u>: probability that truly positive samples are detected as positive by analytical test

   100 – false negative rate
- <u>Specificity</u>: probability that truly negative samples detected as negative by analytical test

   100 – false positive rate
- <u>Level of detection (LOD)</u>: lowest level of contamination reliably detected by analytical test
  - LOD expressed as ratio of organisms to quantity tested material (*e.g.*, CFU per gram, MPN per mL, CFU per square-ft) but definitions vary (e.g., LOD95, POD)

### Food Safety and Inspection Service: Factors Impacting Detection and Method Specifications

- Detection as measured by sensitivity, specificity, and LOD can vary based on:
  - Specific strains of pathogen
  - Intrinsic factors for the sample matrix
    - Levels of competing bacteria
    - Fat, salt, pH and additives
- Experimental design for the validation study (*e.g.*, cell stress, etc.)

# **Complying with the Validated Protocol**

- Do AOAC/AFNOR/ISO citations match the protocol in use?
  - Modifications are common, and some contribute to greater potential for false negative result
- Compare the lab procedure to the validated protocol (*i.e.*, package insert)
- If culture confirmation is used, verify that it follows validated method as well

Methods Not Validated by Recognized Organizations

- "Supplemental" or "extension" validations
- *E. coli* O157:H7 and non-O157 STEC testing for 325-375g test portions
  - Modifications required for AOAC validated procedures based on 25g
  - Instructions for sample preparation may not be clear for the lab

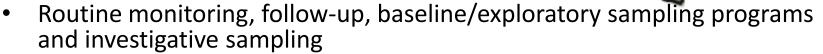
Laboratory Accreditation and Quality Assurance

- ISO 17025 = protocol for establishing and documenting a microbiology laboratory quality program (*i.e.*, "HACCP" for labs)
- Accrediting bodies = A2LA and others
- Accreditation implies robust quality program but does not necessarily indicate methods meet FSIS expectations
  - Laboratories are able to perform the methods they use as expected, but methods are not "accredited" to be fit for purpose
- Laboratories are not required to be ISO accredited, but should have quality assurance programs that ensure results are reliable and accurate

# **FSIS Testing Programs**

# **Microbiological Testing by FSIS Laboratories**

- Three Field Service Labs administer regulatory testing programs
  - Washington DC
     Executive Associate for Laboratory Services
  - Athens, Georgia
    - EFSL-routine/other testing
    - LQAS-quality assurance
    - FERN- biosecurity
  - St. Louis, Missouri
    - MWFSL-routine testing
  - Albany, California
    - WFSL-routine testing
    - Canning issues



#### ISO 17025 Accredited



**FSIS Microbiological Sampling Programs** 

| Sample Type | Number Collected |
|-------------|------------------|
| Domestic    | 136,944          |
| Import      | 10,207           |

Fiscal year 2018 data

# FSIS Micro Sampling Program Objectives

- Assess effectiveness of industry process controls
- Provide critical feedback to industry
- Monitor compliance with performance standards, zero-tolerance policies
- Allow FSIS to monitor industry-wide trends
- Serve as a strong incentive to reduce the occurrence of pathogens in products
- Capture pathogen characterization information (*i.e.*, serotype, speciation, PFGE, antimicrobial resistance, whole genome sequencing)

## **FSIS Sampling Programs**

- Sampling plans measure compliance with performance standards:
  - Salmonella and Campylobacter verification programs (raw poultry)
- Zero-tolerance policies for food pathogens
  - E. coli O157:H7 and non-O157 Shiga toxin-producing E. coli (non-O157 STEC) (raw non-intact beef or components of raw ground beef)
  - Listeria monocytogenes in RTE and pasteurized egg products and on food contact surfaces
  - *Salmonella* in RTE and pasteurized egg products

# FSIS Methods and Pathogenspecific Issues to Consider

Shiga toxin-producing E. coli (STEC) Testing

- Includes:
  - *Е. coli* О157:Н7
  - Six non-O157 Shiga toxin-producing *E. coli* (STEC) (non-O157 STEC) O26, O45, O103, O111, O121, and O145

# **O157 STEC Program**

- Strain must have:
  - 0157(+)
  - *stx*(+) OR *stx*(-) and H7(+)
  - biochemical(+)
  - Currently FSIS only analyzes beef manufacturing trimmings (MT60) for non-O157 STECs

# **Non-O157 STEC Program**

- Six non-O157 STEC = O26, O45, O103, O111, O121, O145
  - Strain must have:
    - *stx*(+) and *eae*(+) genes
    - one of the six O-groups
    - biochemical(+)
  - Currently FSIS only analyzes beef manufacturing trimmings (MT60) for non-O157 STECs
    - Phased rollout MT65 MT64 MT43

### E. coli Top Seven STEC Analysis (MLG 5C)

| Day 1  | Sample Prep and Primary Enrichment<br>42°C±1 for 15-24 hours   |  |  |
|--|--|--|--|
| Day 2  | Perform PCR<br>All samples that do not test PCR<br>negative are carried forward for<br>further analysis              | negative or Screening<br>potential (+) –<br>stx(+) & eae(+)                            |  |
|  | Immunomagnetic Bead Capture &<br>Rainbow Agar Plating  |  |  |
| Day 3  | Latex Agglutination & Sheep Blood<br>Agar Plating  | negative<br>presumptive (+) – agglutination(+) and<br>rapid screen (+) or inconclusive |  |
| Day 4 / 5                                      | Perform<br>-stx/eae gene analysis<br>-latex agglutination & genetic<br>serological test<br>-biochemical confirmation | Confirmation   |  |
| Non-O157 Conf<br>stx(+) &<br>biochen<br>O-grou | eae(+)<br>nical(+)   | O157 Confirmed Positive<br>O157(+); biochemical(+)<br>stx(+) OR stx(-) & H7(+)         |  |

Larger *E. coli* O157:H7 and Non-O157 STEC Test Portions

- Larger test portions (325-375 grams) are most important for N60 and other composite samples containing many samples
- Less important for single "grab" samples of ground beef final product testing when:
  - Trim and components have already been tested using robust sampling and 325-375-gram test portions
  - Multiple samples are collected throughout the production day
- Methods must be adapted, optimized and validated for effective use with 325-375 gram test portions

### E. coli O157:H7 and Non-O157 STEC Testing Concerns

- Supplemental validation and special instructions for testing larger test portions
  - For enrichment periods <15 hours</p>
  - 325-375g test portions typically often require longer minimum enrichment period than 25g
- Culture-based detection and confirmation requires immunomagnetic separation (IMS)

Listeria Testing

• Includes:

- *L. monocytogenes* testing (FSIS)

- Listeria-like or Listeria spp. testing (industry)

## Listeria monocytogenes (MLG 8.11)

| Day 1 | Sample Prep and Primary Enrichment<br>Stomach 25g sample + 225 ml UVM<br>Incubate 30°C for 20-26 hrs |                            | Enrichment   |
|-------|--|----------------------------|--------------|
| Day 2 | Plating, Secondary Enrichment & Rapid Screen<br>MOX & MOPS-BLEB<br>Incubate 35°C for 18-24hrs        | confirm (-)<br>possible(+) |              |
| Day 3 | Streak plates for next day<br>Horse blood and MOX plates   |                            | Screening    |
| Day 4 | GeneProbe and restreak<br>Incubate 35°C variable time  | presumptive(+)             |              |
| Day 5 | Biochemical analysis, restreak &<br>GenProbe   | presumptive (+)            | Confirmation |
| Day 6 | Further characterization,<br>morphological, and atypical isolate<br>analysis                         |                            |              |
| Day 7 | GenProbe   | confirm (-/+)              | 59           |

Expectations for *Listeria* Environmental Testing Equivalence

- Compliance Guidelines Controlling Lm in Postlethality Exposed RTE Meat and Poultry Products Jan 2014
- For optimal sensitivity of detection, method for food contact surface testing must:
  - Validated by a recognized body (*e.g.*, AOAC, AFNOR)
  - Be enrichment-based
  - Enrich the entire sponge/swab sample
    - *e.g.,* an aliquot from sponge/swab does not provide opportunity to detect bacteria trapped in the sponge

Analytes for Industry Food Contact or Environmental Surface Testing

- Establishment laboratories test for one of the following:
  - Listeria monocytogenes
    - Use internationally recognized **enrichment-based method** that biochemically confirms culture as *L. monocytogenes*
  - *Listeria* spp.
    - Use internationally recognized **enrichment-based method** that uses ELISA, PCR or other screening technology to provide more rapid but less specific *Listeria* spp. result
  - "Listeria-like" indicator bacteria
    - Use the first part of an internationally recognized enrichment-based method to find suspect *Listeria* colonies (*e.g.*, darkened colonies on MOX using the FSIS method)

## Salmonella Testing

- Raw products
  - Meat and turkey carcass sponge samples
  - Chicken carcass/parts rinsates
  - Raw meat and comminuted poultry
- Processed products
  - RTE (325g portion)
  - Pasteurized egg

# Salmonella (MLG Ch. 4.10)

| Day 1 | Sample Prep and Primary Enrichment<br>Stomach sample + BPW<br>Incubate 35°C for 20-24 hrs   |                                | Enrichment         |
|-------|---|--------------------------------|--------------------|
| Day 2 | Perform PCR<br>All samples that do not test PCR negative are<br>carried forward to RV and TT broth<br>Incubate 42°C for 22-24 hrs | confirm (-)                    | Screening          |
| Day 3 | Streak RV and TT on BGS and DMLIA<br>plates<br>Incubate 35°C for 18-24 hrs  |                                |                    |
| Day 4 | Pick suspect colony from Plate<br>medium to TSI and LIA slants.<br>Incubate 35°C for 22-26 hrs                                    |                                |                    |
| Day 5 | Streak on SBA for biochemical testing<br>Incubate 18-24 hrs at 35°C   | presumptive (+)<br>confirm (-) |                    |
| Day 6 | Perform biochemical testing and serology using colony from SBA plate.   | confirm (-/+)                  | Confirmation<br>63 |

**Campylobacter** Testing

- Qualitative
  - Enrichment-based (as opposed to direct plating) since Aug 27, 2018 - exception: "other raw chicken parts" (EXP\_CPT\_OT01 and LO\_CPT\_OT01)

- Targets
  - C. jejuni, C. lari or C. coli

### **Campylobacter (MLG 41.04) - Qualitative**

| Day 1-2 | Sample Prep and Primary Enrichment or Pl<br>Sample + BF-BEB or plate (Campy-Cefex)<br>Incubate 42°C for 48 hrs |               | Enrich or plate   |
|---------|--|---------------|-------------------|
| Day 3   | PCR Screen & Plating/isolation<br>Campy-Cefex<br>Incubate 42°C for 48 hrs                                      | confirm (-)   | Plating/isolation |
| Day 5   | Microscope examination for<br>morphology/motility  |               | Confirmation      |
|         | Latex agglutination  | confirm (-/+) |                   |

#### **RESULTS ARE NOT USED FOR REGULATORY PURPOSES**

# **Issues for** *Campylobacter***Testing**

- Campylobacter is highly vulnerable to freezing
   Do not freeze samples
- Can be a challenging test (inconsistent results across labs)

# Industry Testing Programs

### Food Safety and Inspection Service: Microbiological Testing by FSIS-Regulated Establishments

- Industry testing aims to:
  - Fulfill regulatory requirement (9 CFR 310.25, 381.94, 430.4, 590.580)
  - Support decisions made in hazard analysis (9 CFR 417.2 (a))
  - Provide on-going verification of HACCP plan (9 CFR 417.4 (a)(2))
  - Evaluate effectiveness of sanitary SOPs (9 CFR 416.14)
  - Fulfill purchase agreements
  - Respond to process deviations

### Food Safety and Inspection Service: Communication Between Establishment and Laboratory is Vital

- The communication challenge
  - The establishment may not understand the testing conducted on their behalf
  - The laboratory does not necessarily know what the establishment needs
  - The laboratory may not be aware of special validated procedures for larger test portions
- The establishment is ultimately responsible

**Establishment Documentation for Testing Methods** 

- Does the establishment have the necessary documentation?
  - Can the establishment provide the method used for microbial detection?
  - Can the establishment provide evidence that the method used was properly validated by an independent body?
  - Can the establishment explain why the method fits the need?

# **Issues for Industry Labs**

- On-site vs. off-site labs
  - Shipment of samples/handling during shipment
- Overarching concerns for on-site labs
  - Is testing effective?
  - Is testing safe in that facility?
    - Enrichment of pathogens in an establishments
- Evaluate the following:
  - Are personnel qualified?
  - Does the lab have proper equipment and materials for testing and disposal of contaminated media?
  - Do they follow the validated testing protocol?

### Food Safety and Inspection Service: Establishment Responsibilities for Laboratory Testing

- The establishment is ultimately responsible for the testing they request from private laboratories
- Has the establishment properly conveyed testing needs?
  - e.g., test portion equivalent to FSIS as opposed to the default 25g in protocols.
- Is the laboratory aware of FSIS expectations?
   Directives, Notices and guidance
- Establishment should provide documented detailed methodology and validation information for FSIS review

## Food Safety and Inspection Service: FSIS Verification of Establishment Sampling and Testing Programs

- Effectiveness verified by FSIS
  - Reviews/observations of EIAOs during FSA
  - Establishment provides supporting documentation
  - Technical and policy support provided through askFSIS
  - Establishment, not laboratory, is responsible for implementing effective program

## Food Safety and Inspection Service: FSIS Verification of Establishment Sampling and Testing Programs

- Focus of FSIS' evaluation
  - Is the method fit for the intended purpose?
  - Does the method support the hazard analysis decisions?
  - Is the method comparable to the appropriate FSIS method (or is there justification for an alternative)?
  - Is a comparable or appropriate test portion used?
  - Is the method validated and used under validated conditions?
  - Does the laboratory assure the quality of the results?

# Helpful Guidance

- RTE
  - Appendix A FSIS Salmonella Compliance Guidelines for Small and Very Small Meat and Poultry Establishments that Produce Ready-to-Eat (RTE) Products and Revised Appendix A (Jun 2017) – Being updated
  - Appendix B FSIS Compliance Guideline for Stabilization (Cooling and Hot-Holding) of Fully and Partially Heat-Treated RTE and NRTE Meat and Poultry Products Produced by Small and Very Small Establishments and Revised Appendix B (Jun 2017) – Being updated

- STEC
  - Compliance Guideline for Minimizing the Risk of Shiga Toxin-Producing *Escherichia coli* (STEC) and Salmonella in Beef (including Veal) Slaughter Operations (2017)
  - Compliance Guideline for Establishments Sampling Beef Trimmings for Shiga Toxin-Producing *Escherichia coli* (STEC) Organisms or Virulence Markers (August 2014)

- HACCP
  - Meat and Poultry Hazards and Controls Guide (Mar 2018)
  - FSIS Compliance Guideline HACCP Systems
     Validation
     (April 2015)

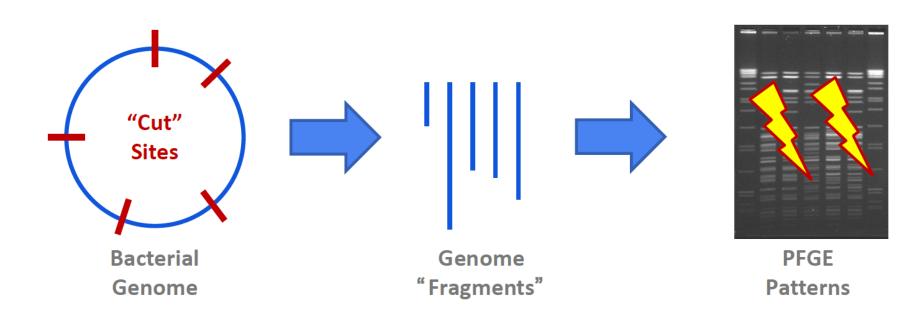
- Microbiological Test Methods and Laboratories
  - Establishment Guidance for the Selection of a Commercial or Private Microbiological Testing Laboratory (June 2013)
  - FSIS Guidance for Evaluating Test Kit Performance (October 2010)
  - Foodborne Pathogen Test Kits Validated by Independent Organizations

Food Safety and Inspection Service: Whole Genome Sequencing

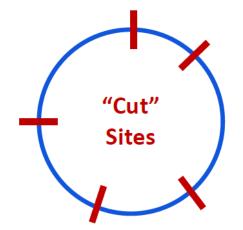
# Whole Genome Sequencing

# Whole Genome Sequencing – Background before WGS

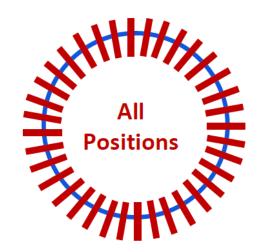
# PFGE



# Whole Genome Sequencing – PFGE-WGS Comparison



PFGE only gives information at a "cut" site via the banding pattern



WGS has the ability to give us information at nearly every position in the genome

# Food Safety and Inspection Service: Whole Genome Sequencing - Benefits

- WGS has a number of uses that benefit FSIS and its mission to protect public health.
- These uses include:
  - identifying harborage and cross-contamination of pathogens in FSIS-regulated facilities,
  - tracing human illness outbreak data to regulated food products, and
  - identifying unique genes related to virulence and pathogenicity, survival and adaptation, and resistance to biocides (sanitizers, metal, etc.) and antimicrobials.

Whole Genome Sequencing – The Transition

- FSIS began performing WGS in parallel with PFGE for *Lm* starting in FY13 and for all pathogens starting in early FY16.
- Centers for Disease Control and Prevention (CDC) PulseNet partners are transitioned away from using PFGE as the primary molecular characterization tool toward using WGS.
- In coordination with CDC PulseNet, FSIS suspended PFGE for *Lm* and as of January 15, 2018, now generates *Lm* characterization through WGS only.

Whole Genome Sequencing – How is WGS analyzed?

- FSIS uses different tools to analyze WGS information including:
  - Multi-locus Sequence Typing (MLST) resulting in
    - Public Sequence Type
    - Allele Code
  - High-quality Single Nucleotide Polymorphisms (hqSNP)

Whole Genome Sequencing – Sequence Typing

Multi-locus Sequence Typing (MLST) MLST can generate a **pattern name or designation** (similar to a PFGE pattern name) based on differences in a pre-defined set of genes.

MLST Results will be Provided by FSIS as Follows:

- **Public Sequence Type** ("MLST ST", "ST", or "pubST")
  - small number of genes (i.e., 6-12)
  - named using the publicly available database developed by Jolley & Maiden (2010) (e.g., publicST09)

# • Allele Code

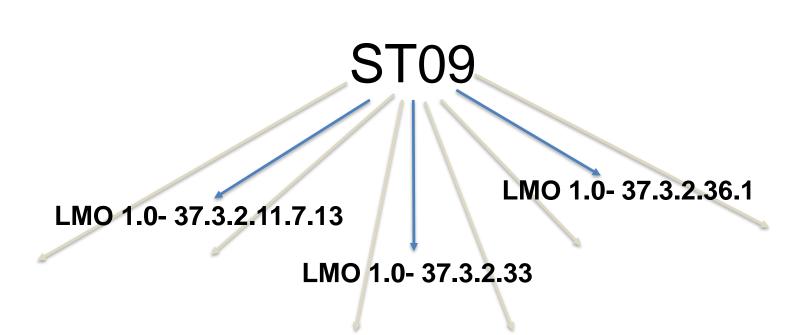
- compares over 1,800 genes
- named by using CDC PulseNet numerical code (e.g., LMO 1.0-5.1.1.2.5.1)

Food Safety and Inspection Service: Whole Genome Sequencing - SNP

Single Nucleotide Polymorphism (SNP)

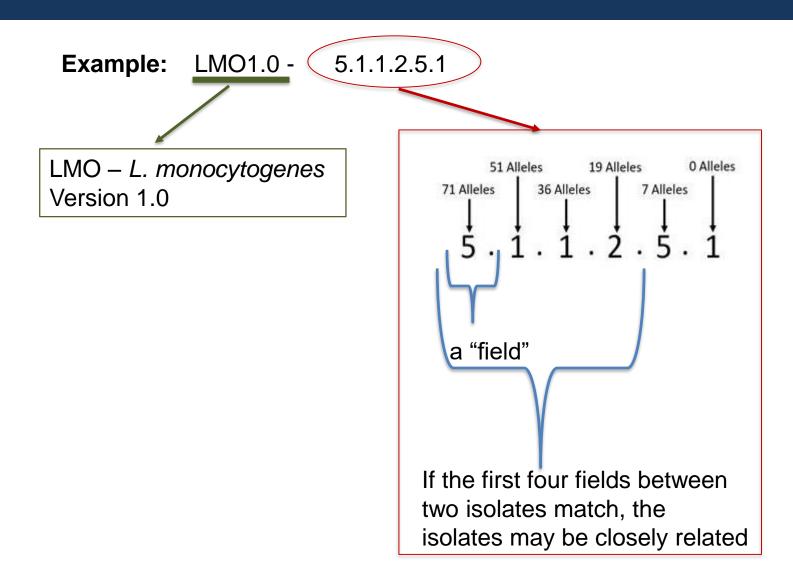
ATGTTCCTC isolate A ATGTTCCTC isolate B This is a single SNP difference!

# Whole Genome Sequencing – Sequence Typing



Allele Code is more specific than public Sequence Type; one public Sequence Type can be inclusive of many Allele Codes.

## Whole Genome Sequencing – Reading Allele Codes



Whole Genome Sequencing



# Allele Code

hqSNP

# Whole Genome Sequencing – Reporting from MCB

Email on Lm-positive from Microbial Characterization Branch (MCB; Eastern Lab, Athens GA)

| Establishment  | Field                                     | 853334755                    | 853344746                    | 853344744                    |                 |
|----------------|---|------------------------------|------------------------------|------------------------------|-----------------|
|                | FormID                                    | 102015479                    | 102051750                    | 102051751                    | Form ID         |
|                | Collect Date                              | 2018-07-30                   | 2018-10-09                   | 2018-10-09                   |                 |
| [              | Project                                   | RTEPROD_RISK                 | INTENV LM W                  | INTENV_LM_W                  | 1               |
|                | FSIS Identifier                           | FSIS31800872                 | FSIS31801180                 | FSIS31801179                 | FSIS Identifier |
|                | NCBI Accession<br>Number                  | SAMN09830008                 | SAMN10269641                 | SAMN10269640                 | ]               |
|                | MLST ST                                   | publicST288                  | publicST288                  | publicST288                  |                 |
| M12345         | Allele Code                               | LMO1.0 - 73.1.1.2.14         | LMO1.0 - 73.1.1.2.14         | LMO1.0 - 73.1.1.2.14         | Allele Code     |
| (LocID: 11981) | NCBI SNP Cluster<br>(Retrieve Date)       | PDS000032940.4<br>2018-10-26 | PDS000032940.4<br>2018-10-26 | PDS000032940.4<br>2018-10-26 | ]               |
|                | Min Food Env (SNP)*                       | 1                            | 1                            | 2                            |                 |
|                | Indicative of Potential<br>Harborage**    |                              | Yes                          | Yes                          |                 |
|                | Indicative of cross-<br>contamination***  |                              | Yes                          | Yes                          |                 |
|                | Min Clinical (SNP)*                       | None                         | None                         | None                         | 1               |
|                | Potentially related to a clinical isolate |                              | No                           | No                           | ]               |

<sup>&</sup>quot;A value of "None" indicates greater than 50 SNPs for this isolate source

<sup>\*\*</sup>Harborage, or repeated introduction is suggested if WGS analysis indicates closely related Lm isolates are found in product, food contact, or nonfood contact environmental samples collected over multiple days, weeks, months, or years. <u>FSIS Notice 48-18</u>

<sup>\*\*\*</sup>Cross-contamination is suggested when closely related Lm isolates are found in product, food contact, and environmental (nonfood contact) samples collected during the same sampling event. <u>FSIS Notice 48-18</u>

## Whole Genome Sequencing – EIAOs needing more info

- When performing a PHRE in establishments with more than one positive RTE sample, EIAOs are to:
  - Request WGS report through <u>Outbreaks WGS@fsis.usda.gov</u>
  - Use WGS to assess if there is a history of harborage or crosscontamination in the establishment.

<u>Note</u>: OPARM is working on updating the "Public Health Risk Evaluation for Establishment" report to include the MLST designations for any historical samples included in the report.

• After an IVT/RLm positive, EIAOs are to make a request through the <u>Outbreaks WGS@fsis.usda.gov</u> Outlook mailbox for WGS analyses.

# Food Safety and Inspection Service: Whole Genome Sequencing – The Future

- FSIS continues to work with FDA, CDC PulseNet, local & state health departments.
- WGS will be the primary subtyping tool for *Campylobacter*, Shiga toxin-producing *Escherichia coli* (STEC), and *Salmonella*.

Stay tuned ....

# **Existing Agency Guidance - askFSIS**

- askFSIS Q&A sometimes contains additional information on testing methods and WGS
- If you cannot answer your question there, please submit to askFSIS

**Questions?** 

Contact me at udit.minocha@fsis.usda.gov

OR:

- Enter question into askFSIS
- Provide documentation for review
- Request "Sampling Queue"

# Food Safety and Inspection Service: Can't find what you are looking for?

| United States Departme<br>Food Safety and Inspection                             |  | FSIS District Offices | Careers Contact Us | Ask Karen askFSIS En Español  |
|--|--|-----------------------|--------------------|---|
| Topics Programs & Servi  | ces Newsroom Forms   |                       |                    | Site Map   Help   |
| Home Answers Submit a G  | uestion My Questions Hel   |                       |                    | Log In   Sign Up  |
| Limit by product All Find the answer to your question                            | Limit by category<br>◄ All   | •                     | Search Tips        | Contact Us<br>Ask a Question<br>Submit a question to our                                      |
| Sort by Default<br>Direction Ascending   |  |                       |                    | Support team.<br>Contact Us<br>If you can't find what you're<br>looking for on our site, give |
| Most Popular Answers Pepperoni Pizza with Whole Wh Use of Celery Powder and Othe | us a call.<br>Give Feedback<br>How can we make this site<br>more useful for you? |                       |                    |   |