

# Human Fecal Score: A standardized method for MST data interpretation

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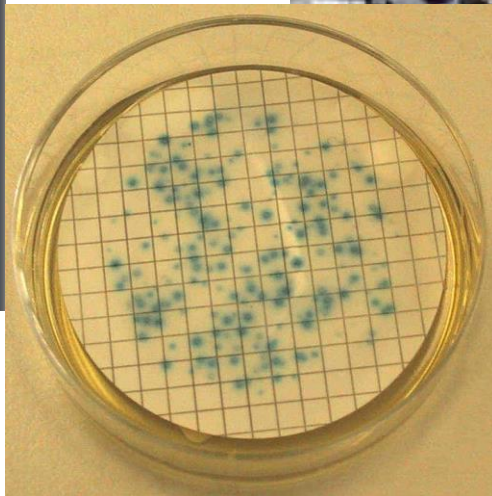
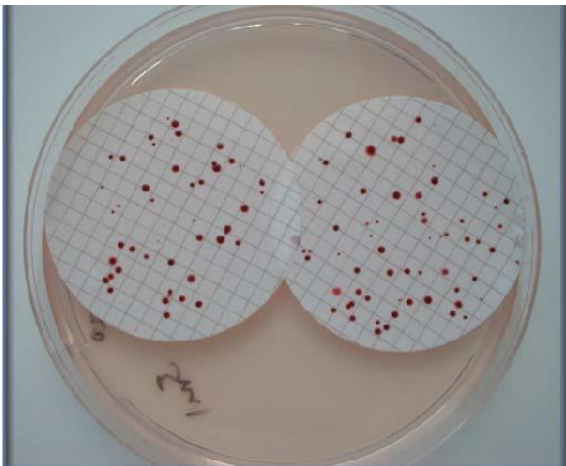
Central/Northern California Ocean and Bay  
Water Quality Monitoring Group

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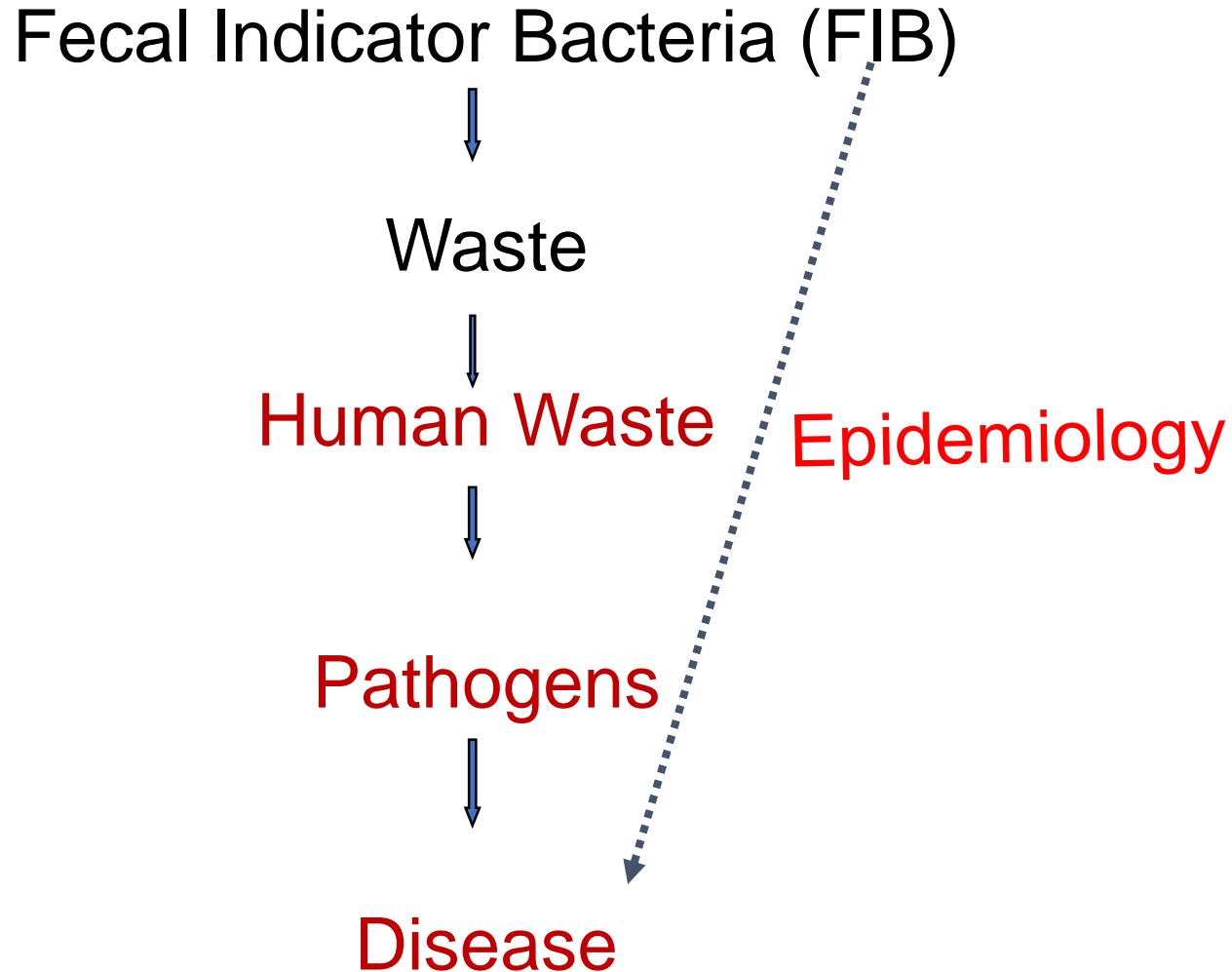
# Water Monitoring

- Fecal Indicator Bacteria (FIB)
  - *Enterococcus* spp.
  - Fecal or total coliforms, *E. coli*



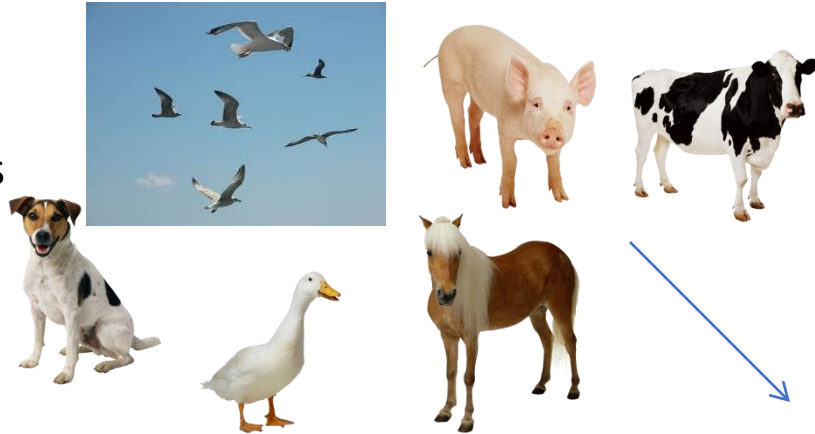
(Image: Donna Ferguson, Google)

# Basis for Monitoring: *the chain of inference*



# Issue: Unspecific

Animals

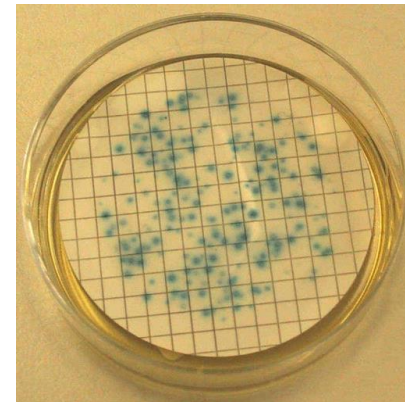


Human

Plants



Soil,  
biofilm



# Consequences

**Mitigation:**

- Hinders source abatement
  - Source identification must precede mitigation

**TMDL:**

- Undermines the basis for water monitoring (i.e. the chain of inference)

**Risk Assessment:**

- Not all sources present the same level of risk

# Microbial Source Tracking

- There are microbes that are only associated with a given fecal source
  - Host and gut microbes co-evolve
    - Physiological difference of the gut
    - Dietary difference between hosts
- MST provides a set of methods to identify sources of contamination
  - Genetic testing of host fecal markers





# MST Applications

- Site assessment: How bad or good is this site?
  - Among all sites within your jurisdiction?
  - Compared to a reference site with little human activities?
  - Compared to a site with measured health risk via epidemiology studies?
  - Before and after implementing BMP remediation actions?
- Answers should be based on data, using “scientifically sound and statistically defensible approaches”
  - Study design
  - Lab analysis
  - Data interpretation

# The Process



Sampling



Lab analysis

sample	Cq	Copy per 100ml
1	36.13	?
1	37.41	?
1	36.05	?
2	Non detect	?
2	Non detect	?
2	Non detect	?
3	Non detect	?
3	Non detect	?
3	Non detect	?
4	30.48	19173
4	30.50	18855
4	30.17	23356
...	...	?
...	...	?
...	...	?
n	Non detect	?
n	Non detect	?
n	Non detect	?

qPCR raw data (marker concentrations) from n samples



Data interpretation  
Site assessment



Action at the site



# The Practice

- Best professional judgement
  - Different experts in different projects
- Worries
  - Unintentional bias: inherent subjectivity and implementation variability by experts?
  - Intentional bias: hired gun by discharger or regulator?



Sampling



Lab analysis



Data Interpretation

# The BPJ Exercise

- Assess variability in MST data interpretation

Create a simulated data set  
(26 site, 20 sample/site)



Ten experts rank the sites 1 to 26 regarding  
relative levels of human fecal contamination

- Experts: research scientists and water quality managers
  - from the federal government
  - a public research agency
  - academic
  - a wastewater treatment agency
- Assessment: Compare ranks among pairs of experts via correlation

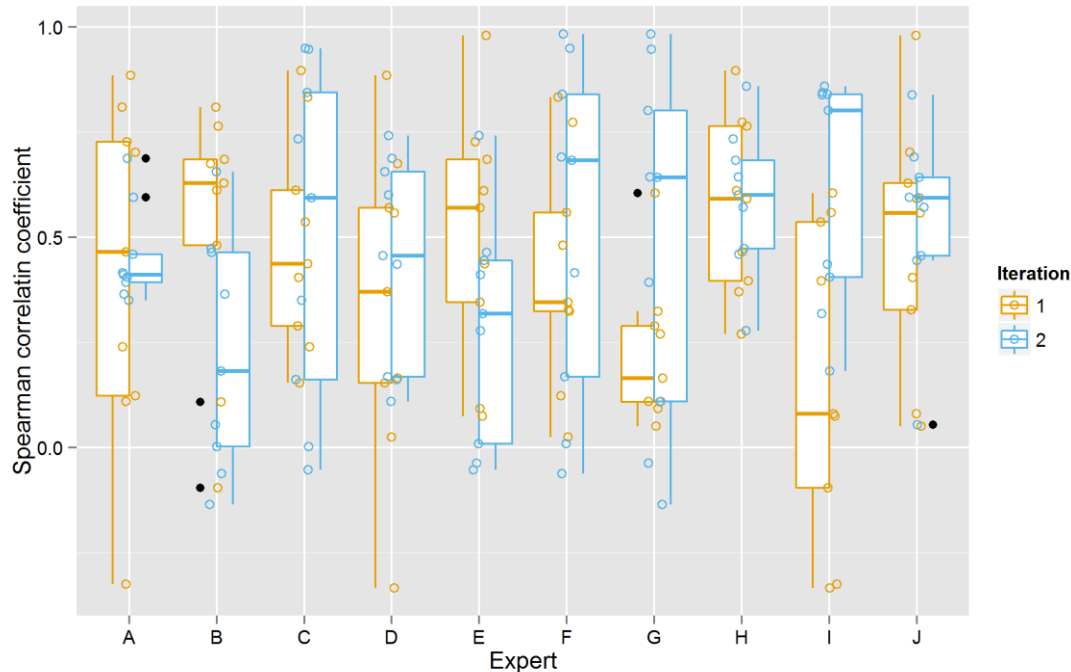
# The BPJ Exercise

- Assess variability in MST data interpretation

Two iterations

- 1<sup>st</sup> iteration: no prior discussion among experts  
r= - 0.33 to 0.98 (avg: 0.41)
- 2<sup>nd</sup> iteration: experts agreed to a set of principles before ranking  
r= - 0.14 to 0.98 (avg: 0.47)

# BPJ Interpretation Highly Inconsistent



- Experts' interpretation of the same data were highly variable
  - 1<sup>st</sup> iteration:  $r = -0.33$  to 0.98 (avg: 0.41)
  - 2<sup>nd</sup> iteration:  $r = -0.14$  to 0.98 (avg: 0.47)

So, how well does BPJ work? – not so well  
Are we right to worry? - yes

# Motivation for Human Fecal Score

- BPJ exercise conclusion: a standardized mathematically defined objective approach is needed!
- Team:
  - SCCWRP: Drs. Yiping Cao, John Griffith, Steve Weisberg
  - USEPA: Drs. Orin Shanks, Mano Sivaganesan, Catherine Kelty
  - Stanford: Drs. Ali Boehm, Dan Wang



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Water Research

journal homepage: [www.elsevier.com/locate/watres](http://www.elsevier.com/locate/watres)



A human fecal contamination score for ranking recreational sites using the HF183/BacR287 quantitative real-time PCR method



(Cao et al 2018)

# Human Fecal Score (HFS): Simple

- Simple
  - Site average concentration of HF183 marker
  - One number to characterize the extent of human fecal pollution at a site

(n samples, 3n data points, for some we don't even have a number for )

sample	Cq	Copy per 100ml
1	36.13	?
1	37.41	?
1	36.05	?
2	Non detect	?
2	Non detect	?
2	Non detect	?
3	Non detect	?
3	Non detect	?
3	Non detect	?
4	30.48	19173
4	30.50	18855
4	30.17	23356
...	...	?
...	...	?
...	...	?
n	Non detect	?
n	Non detect	?
n	Non detect	?

## Human Fecal Score

=  
55





# HFS: Complete

- Uses all data
  - non-detect
  - detected but not quantifiable
  - Quantifiable

( Can't average non-number, e.g. ? )

sample	Cq	Copy per 100ml
1	36.13	?
1	37.41	?
1	36.05	?
2	Non detect	?
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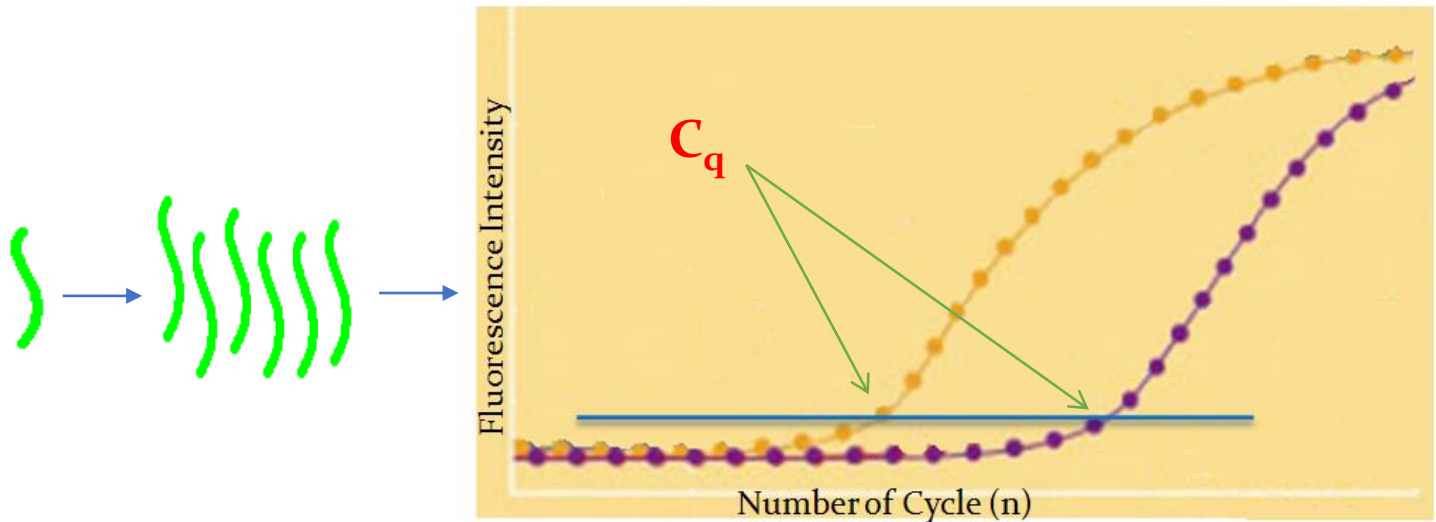
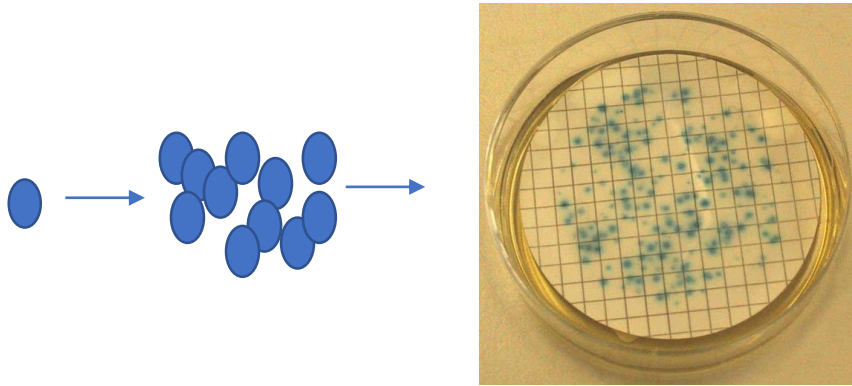
## Human Fecal Score

=  
55



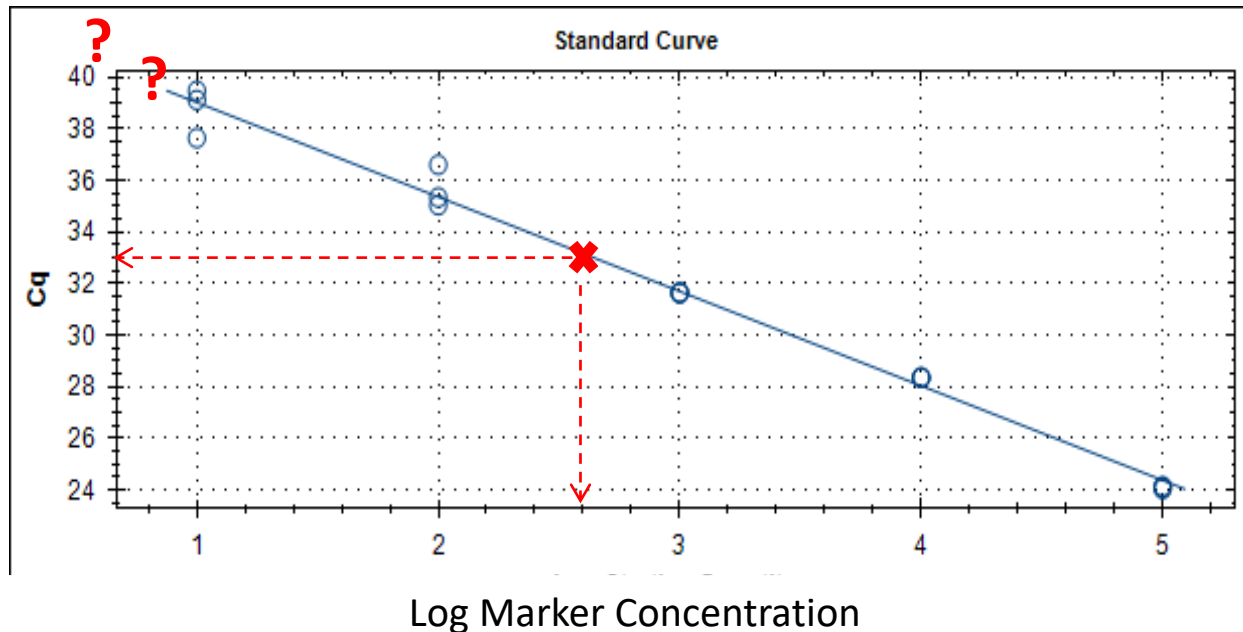
# How do qPCR quantify?

Grow cells vs. “grow” DNA



# qPCR Range of Quantification (ROQ)

- Within Range:  $C_q$  linearly inversely relates to marker concentration
- At low concentration: no more linear relationship
  - Can't quantify using the standard curve
- Non-detect: no quantification



- Previous “solutions”

- Ignore non-detect and detected but not quantifiable

- Arbitrarily assign a number
    - 0, DL/2, DL, LLOQ ...

- Force standard curve outside ROQ

- Statistics for censored data
    - Not applicable in most cases

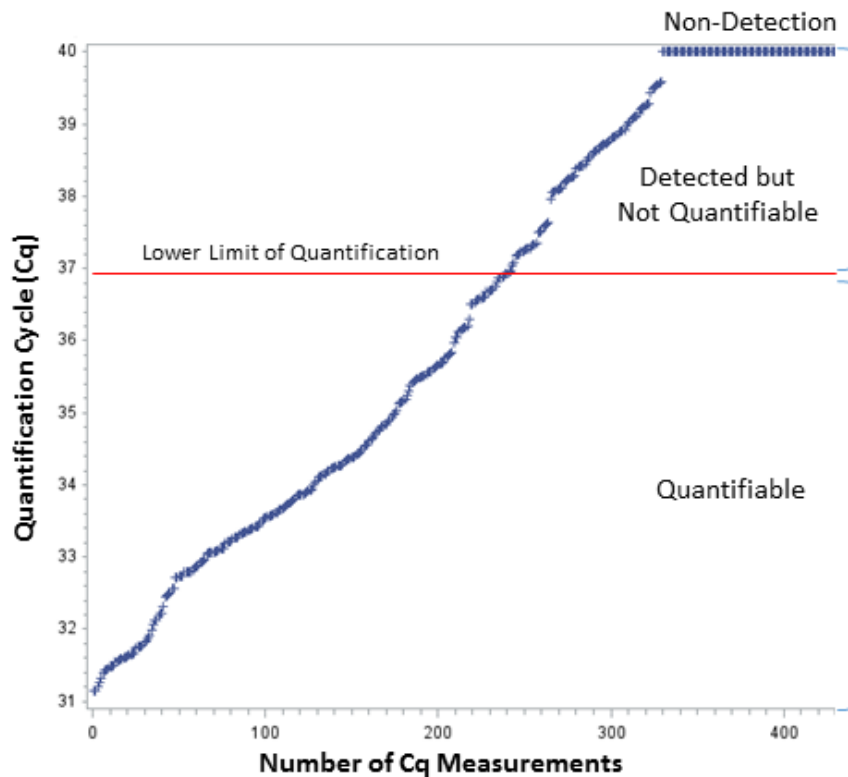
- HFS: use underlying Poisson distribution to estimate ?’s outside ROQ

( Can’t average non-number, e.g. ? )

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# HFS: Based on statistics

- Two different quantification mechanisms
  - Executed by Bayesian models, integrating data uncertainty



**MPN Range**  
Quantification by  
MPN approach

**Human Fecal Score**  
(weighted average)

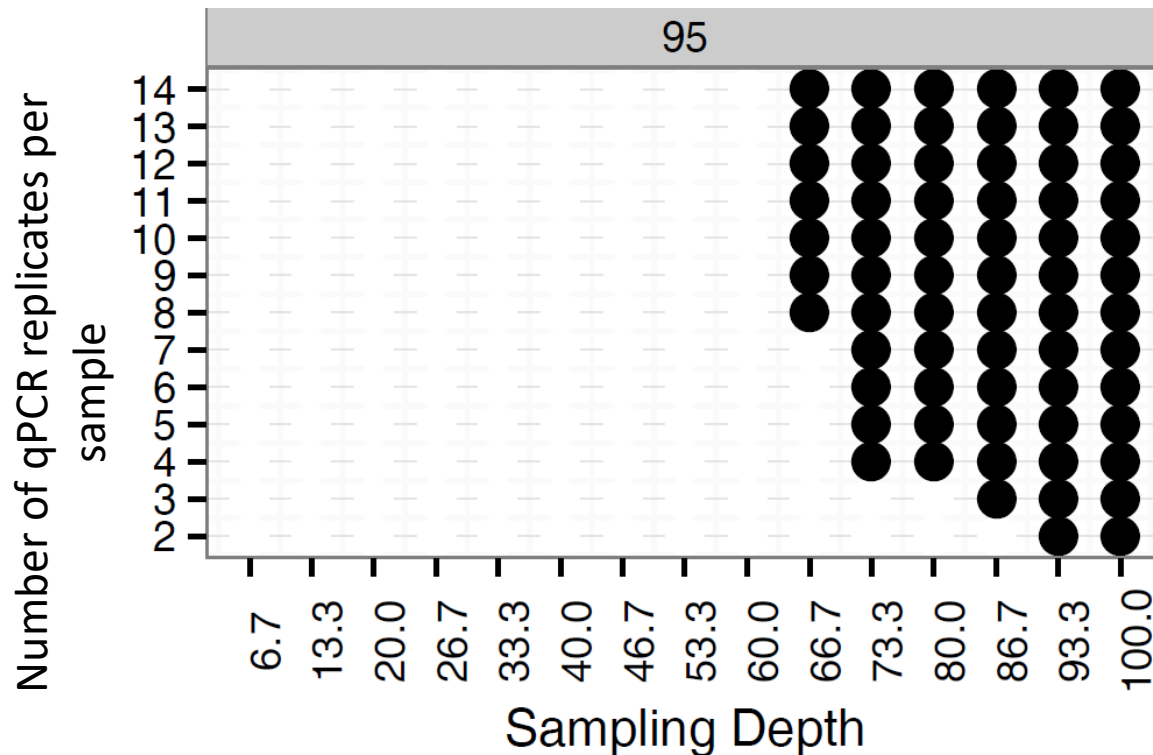
**ROQ Range**  
Quantification by  
calibration model

$$\text{Log}_{10} \text{HFS} = W \cdot \text{Log}_{10} C_1 + (1 - W) \cdot \text{Log}_{10} C_2$$

(Cao et al 2018)

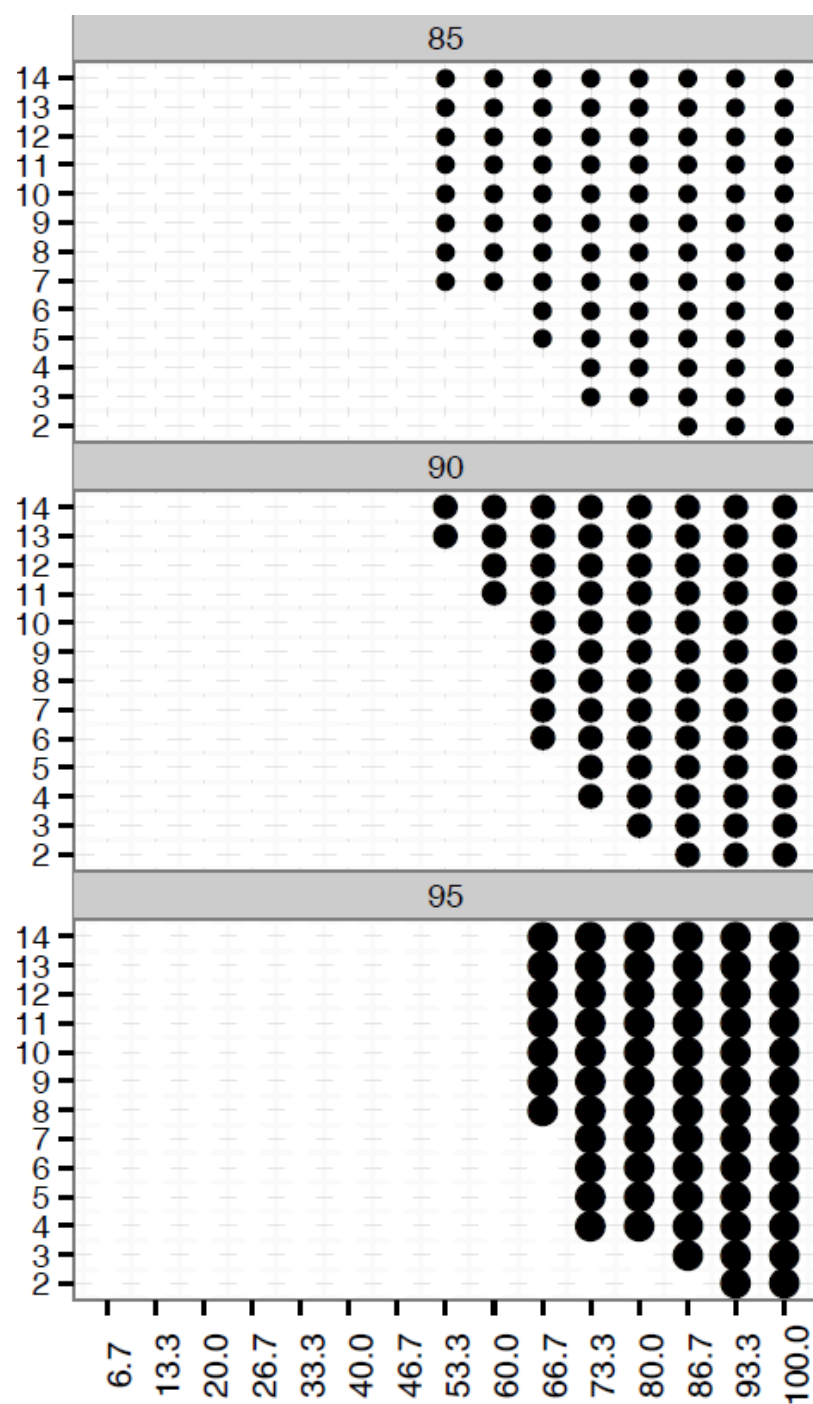
# HFS allows sampling design optimization

- Certainty accepted by managers and/or regulators
- Trade-off between sample size and qPCR replication



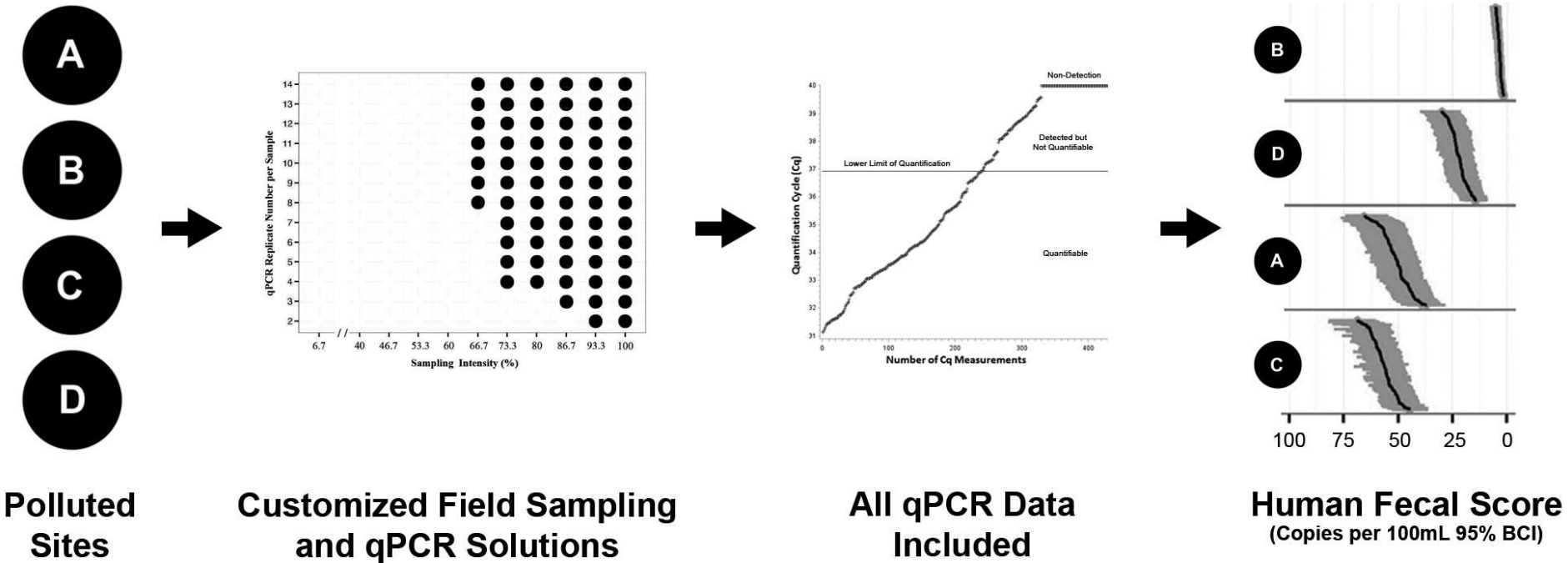


Willing to accept different chances of getting the right answer?



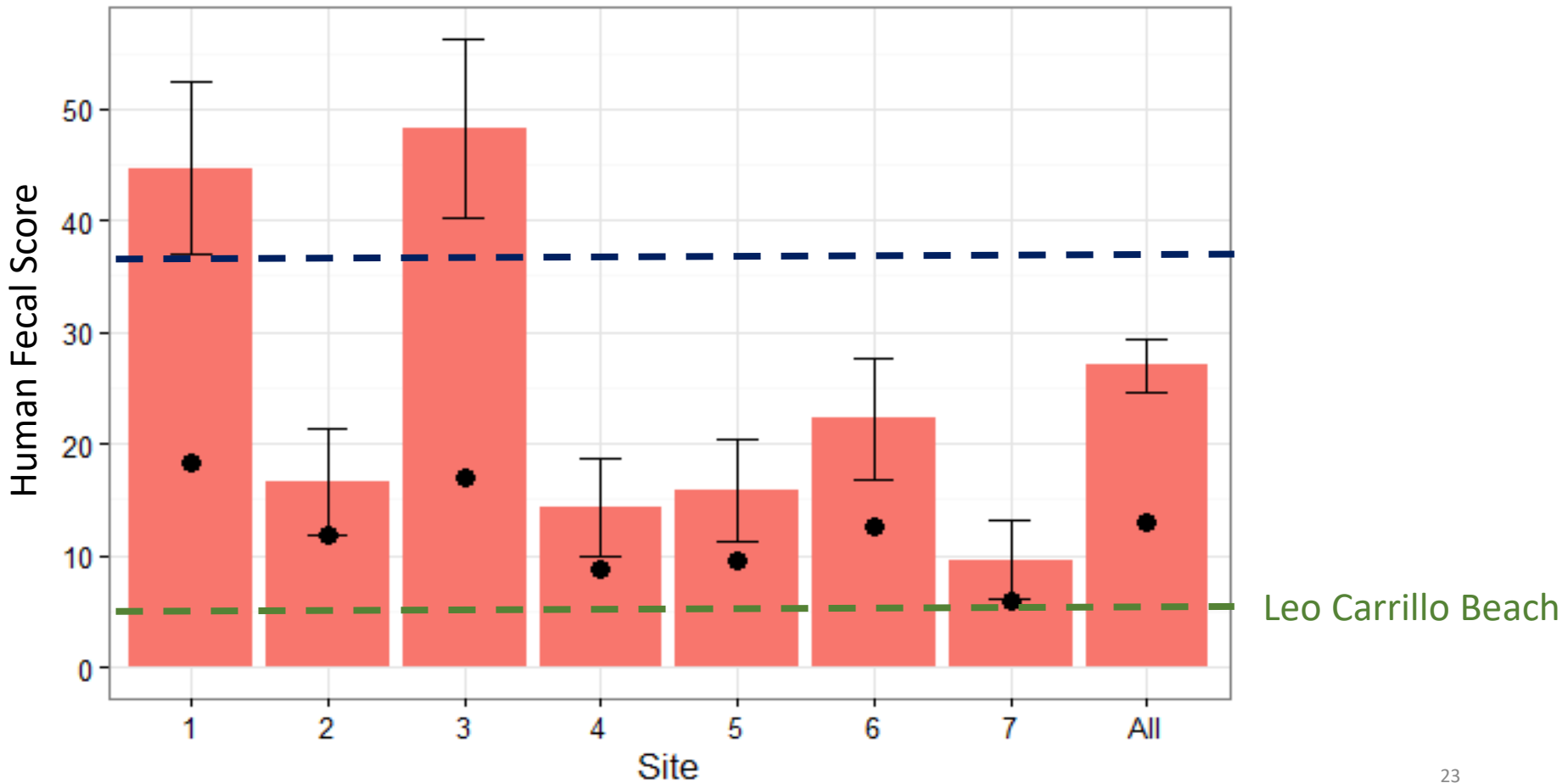
# HFS Application: Prioritizing Remediation

## HUMAN FECAL SCORE FOR SITE RANKING



## STANDARDIZED PROCEDURE

# HFS Application: Real Site Scores



# HFS Summary

- Simple
- Respect data
  - Use everything
  - Add nothing
  - Respect underlying data distribution
  - Integrate uncertainty in data
- Objective
  - Mathematically defined
  - Build on formulas instead of narratives
- Standardization
  - Use the U.S. EPA standard HF183 qPCR method
  - Sampling design



Sampling



Lab analysis



Data Interpretation

# Implications for water quality management

- HFS describes a standardized method for characterizing human fecal pollution level at a site
- General:
  - Other markers: Cow Fecal Score, Gull Fecal Score
  - Other technology: digital PCR
- Potential applications
  - BMP effectiveness
  - Rank sites
  - CSO consent decree compliance
  - QMRA site eligibility

# Thank you!

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