



# The Urinary Microbiome

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# Overview

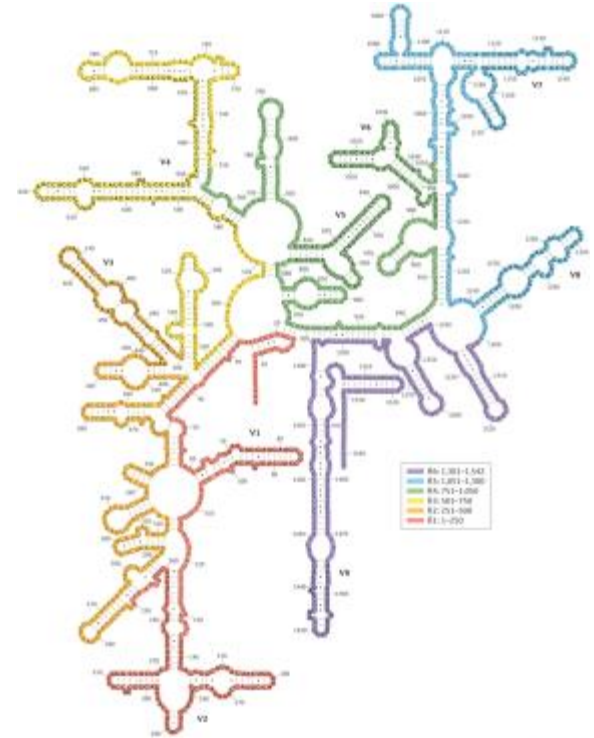
- Existence of a Female Urinary Microbiome & Microbiota
- Implications for Urinary Tract Infections
- Current and Future Projects

# Overview

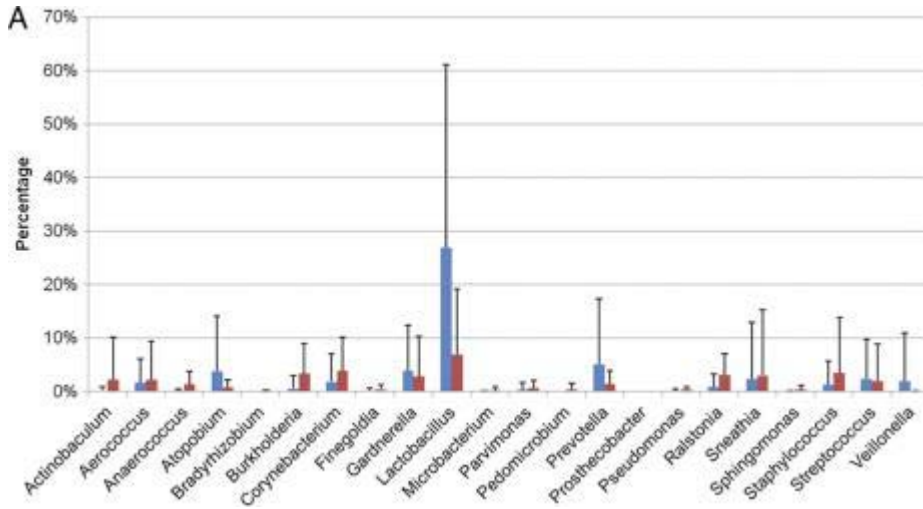
- **Existence of a Female Urinary Microbiome & Microbiota**
- Implications for Urinary Tract Infections
- Current and Future Projects

# A Female Urinary Microbiome exists

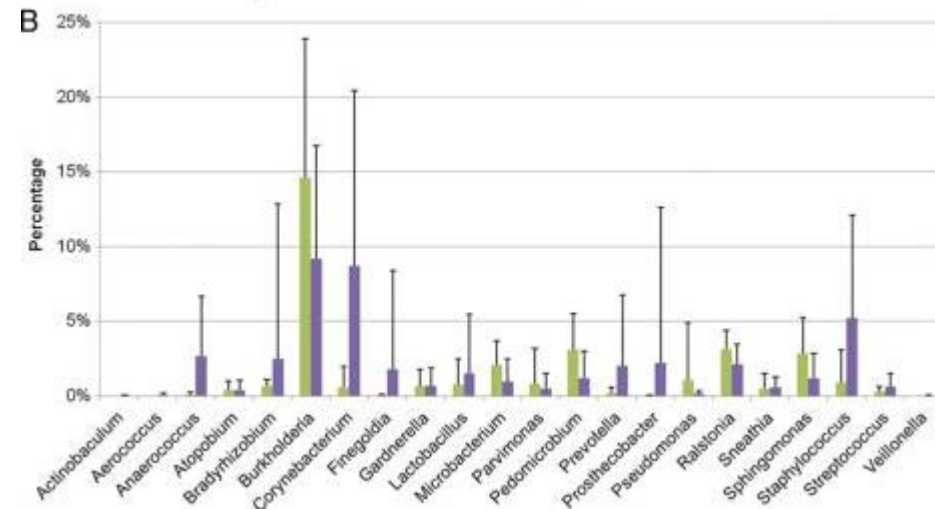
- 16S sequencing – V4 region
- Illumina MiSeq
- Mothur bioinformatics



# A Female Urinary Microbiome exists



■ Transurethral Catheter (TUC)  
 ■ Suprapubic Aspirate (SPA)

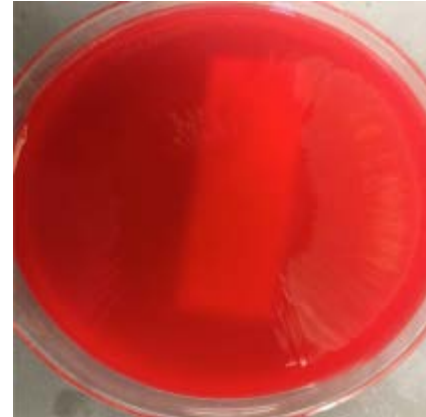


■ Needle Stick  
 ■ Skin Swab

# Urine is not sterile

Protocol (Volume)	Media	Condition	Identification
SUC (1 $\mu$ L)	Blood, MacConkey	Aerobic 35C	24 hr
EQUC (100 $\mu$ L)	Blood, Chocolate, CNA	5% CO <sub>2</sub> 35C	48 hr
	Blood	Aerobic 35C	48 hr
	Anaerobic	Anaerobic 35C	48 hr

## Standard Urine Culture (SUC)



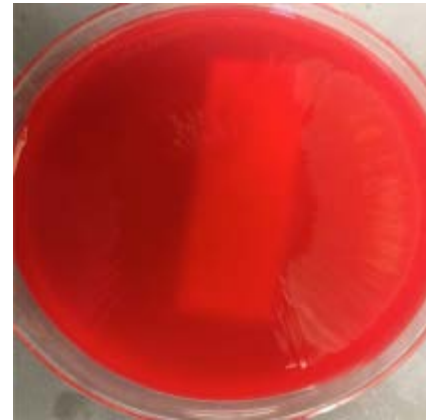
1  $\mu$ L, Aerobic at  
35C, 24hrs

## Enhanced Quantitative Urine Culture (EQUC)

# Urine is not sterile – Female Urinary Microbiota (FUM)

Protocol (Volume)	Media	Condition	Identification
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	Blood	Aerobic 35C	48 hr
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**Standard Urine Culture (SUC)**



1 µL, Aerobic at  
35C, 24hrs

**Enhanced Quantitative Urine Culture (EQUC)**



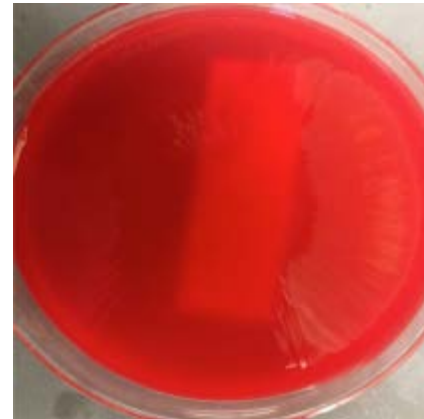
100 µL, Aerobic at  
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# Urine is not sterile – Female Urinary Microbiota (FUM)

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**Standard Urine Culture (SUC)**



1 µL, Aerobic at 35C, 24hrs

**Enhanced Quantitative Urine Culture (EQUC)**

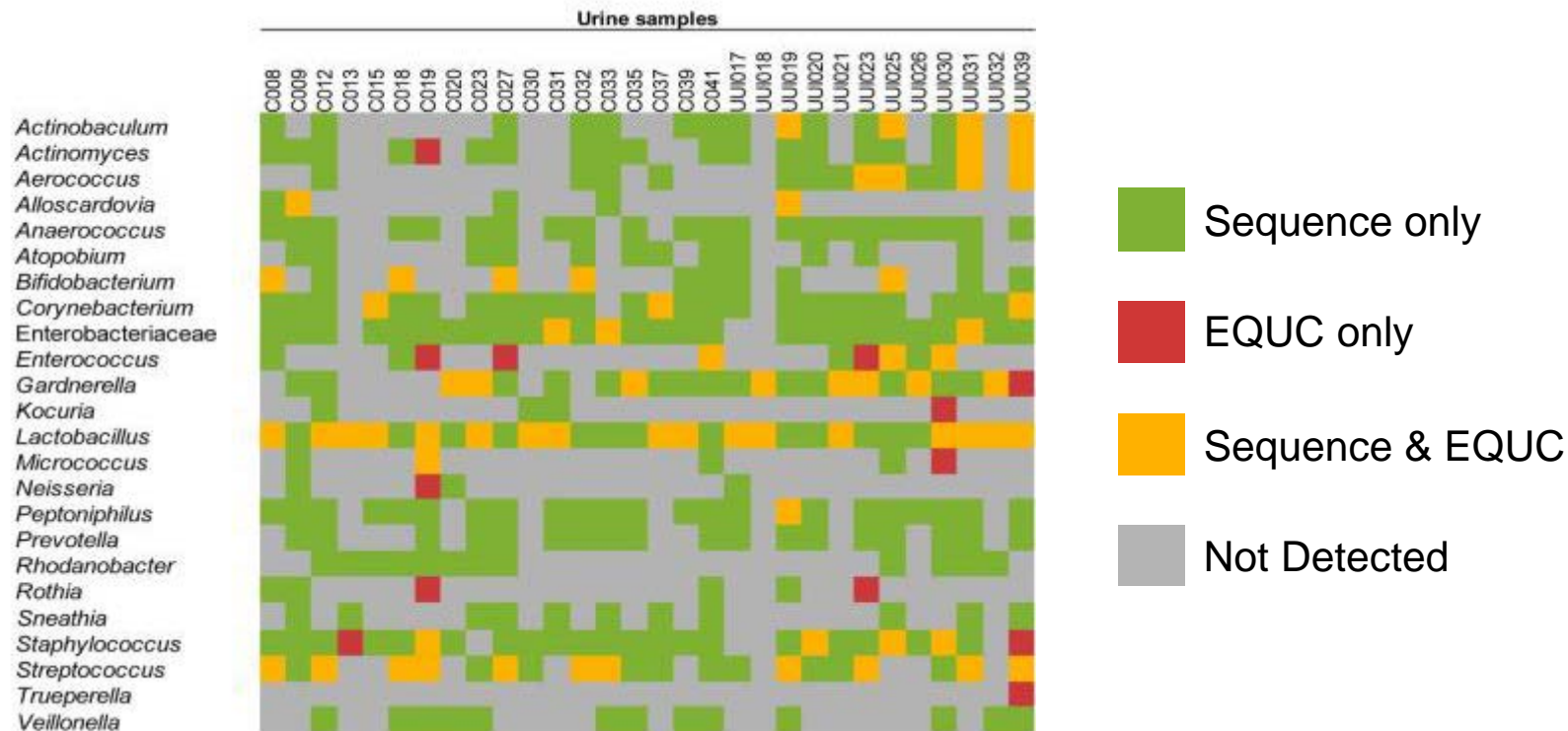


100 µL, Aerobic at 35C, 48hrs

**SUC has a 90% false-negative rate**

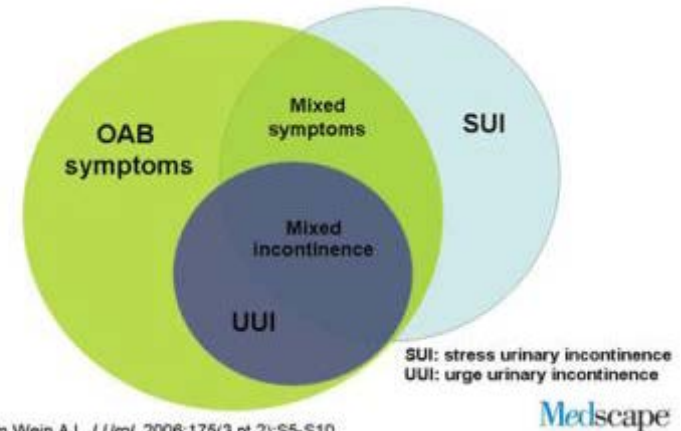
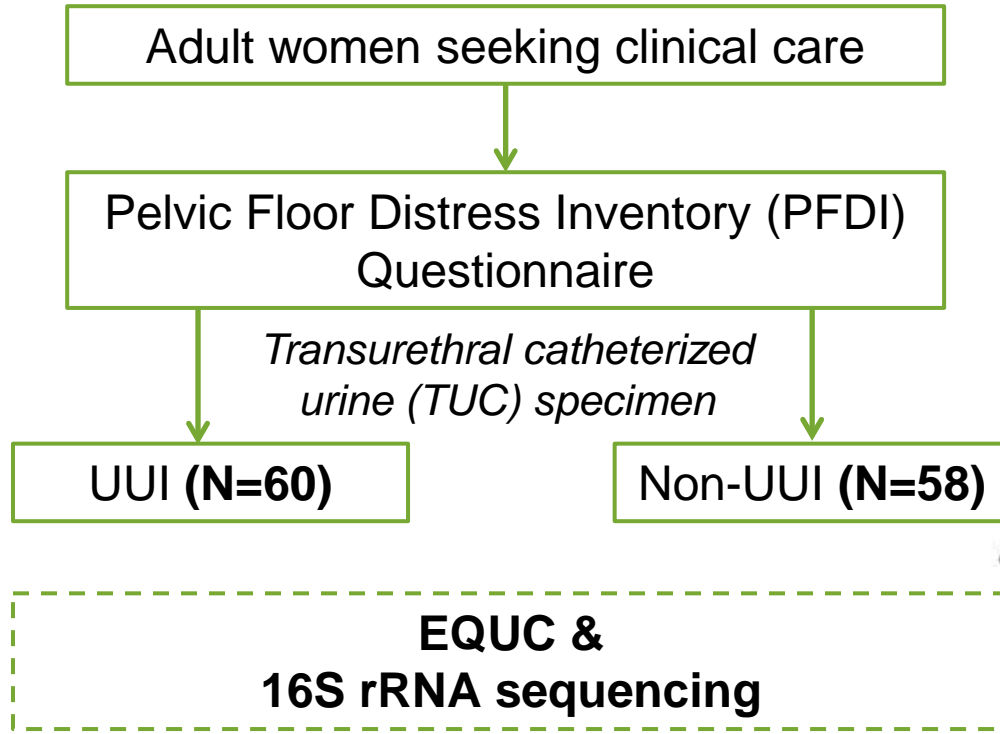
Hilt *et al.*, 2014

# Sequencing & EQUC are complementary

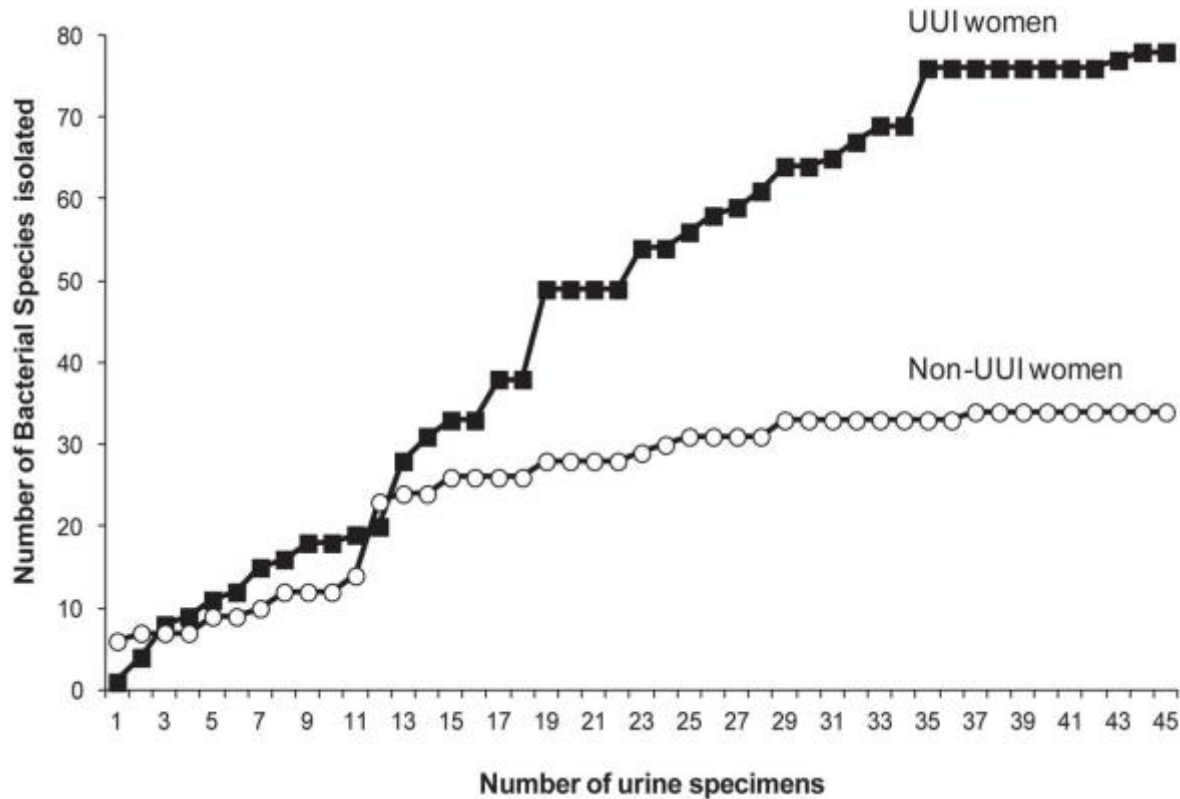


- Sequence only
- EQUC only
- Sequence & EQUC
- Not Detected

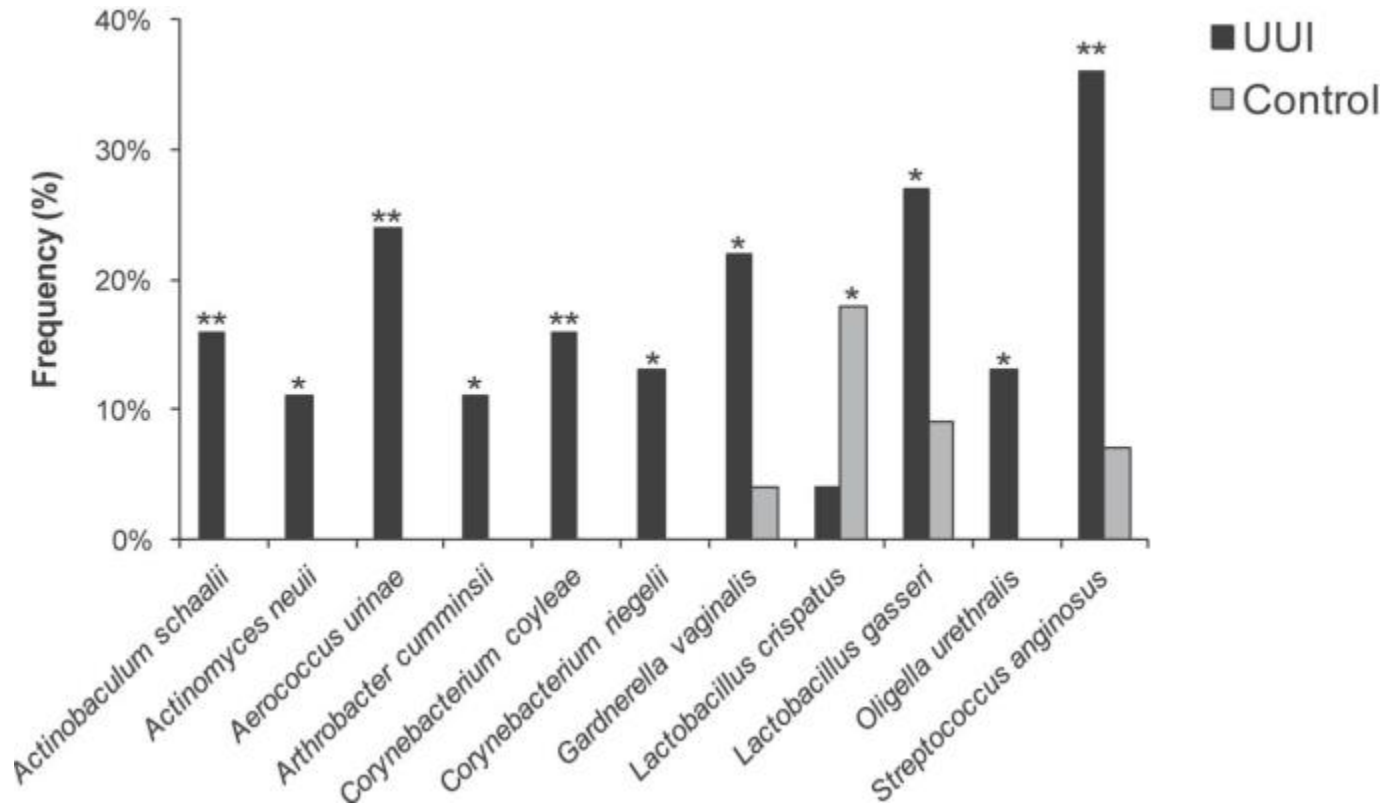
# Is the FUM involved in Lower Urinary Tract Disorders?



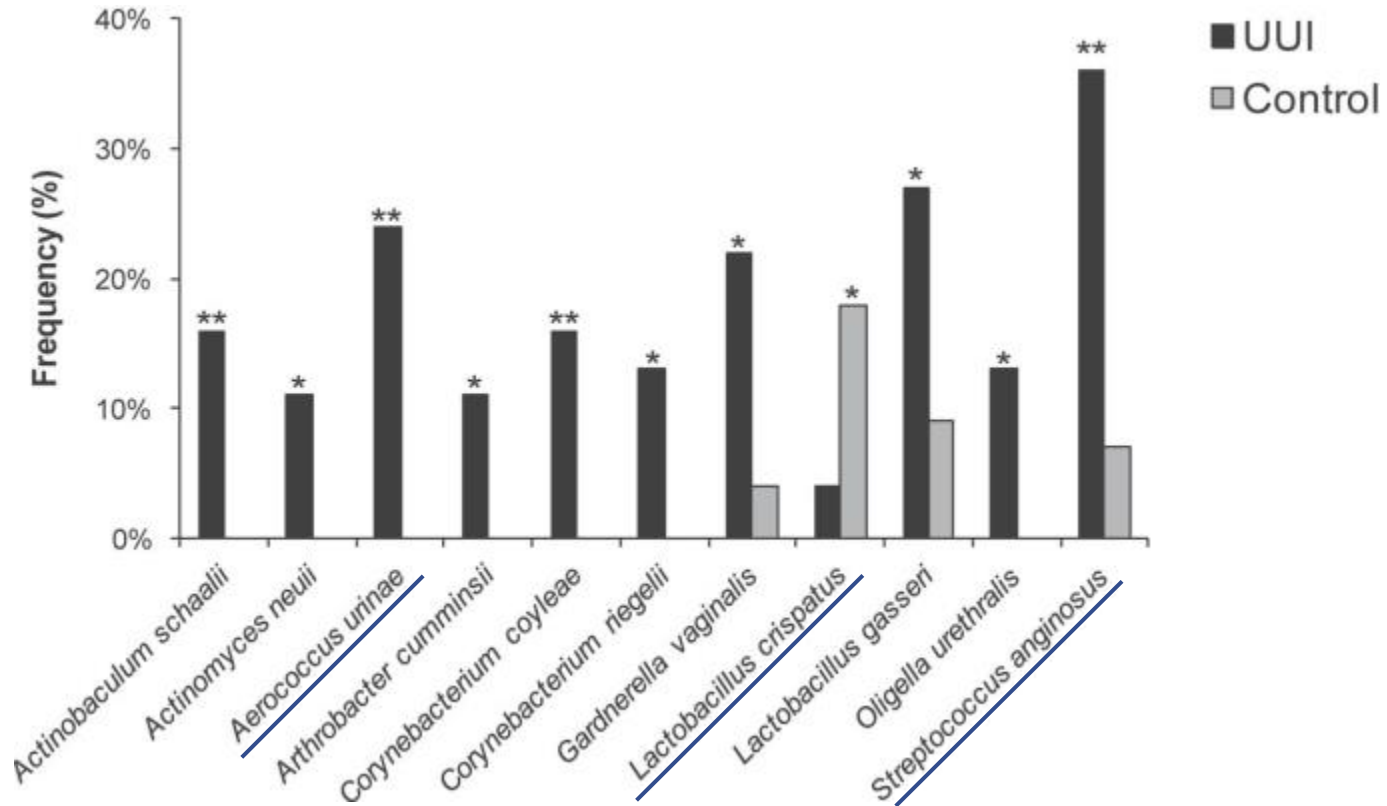
# FUM diversity is higher in women with UUI



# FUM composition differs between UII and non-UII women



# FUM composition differs between UII and non-UII women



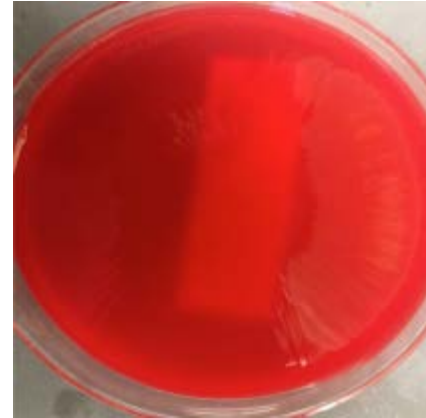
# Overview

- Existence of a Female Urinary Microbiome & Microbiota
- **Implications for Urinary Tract Infections**
- Current and Future Projects

# SUC has a 90% false-negative rate

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**Standard Urine Culture (SUC)**



1  $\mu$ L, Aerobic at  
35C, 24hrs

**Enhanced Quantitative Urine Culture (EQUC)**



100  $\mu$ L, Aerobic at  
35C, 48hrs



# Hypothesis

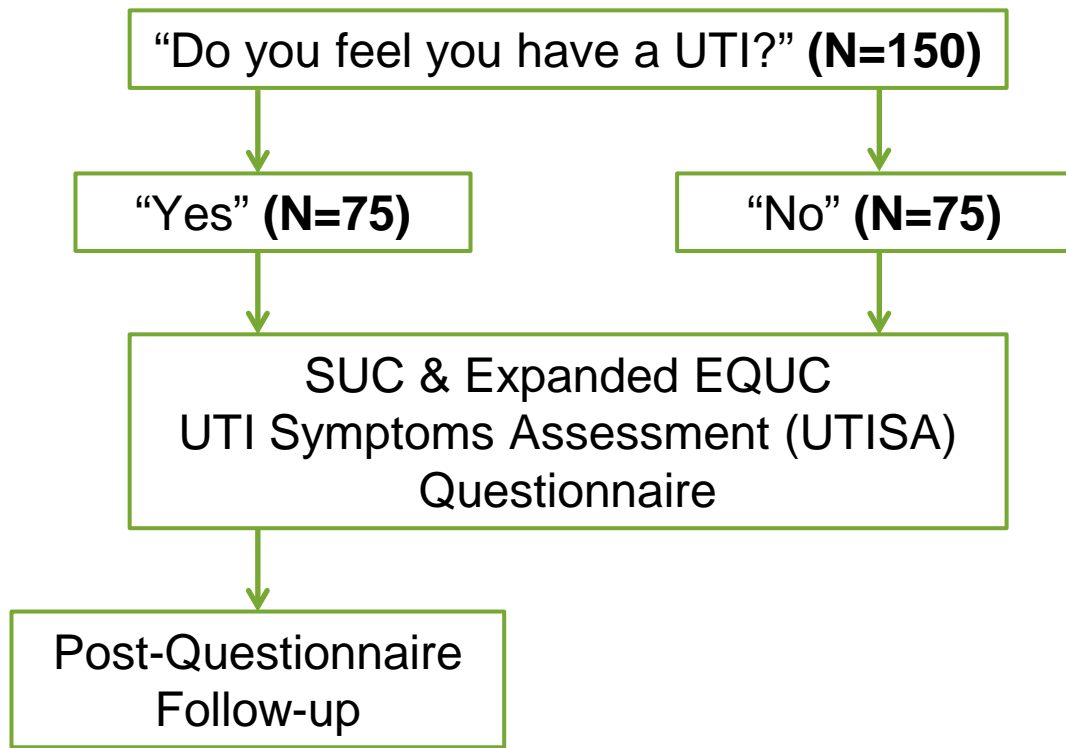
The standard definitions and measures of UTIs  
are inefficient at detecting  
clinically relevant infections.

# Hypothesis

The standard definitions and measures of UTIs are inefficient at detecting clinically relevant infections.

- Standard Urine Culture (SUC) protocol
  - Urinary Symptoms
  - $\geq 10^5$  CFU/mL for diagnosis

# Study design



## UTISA Questionnaire

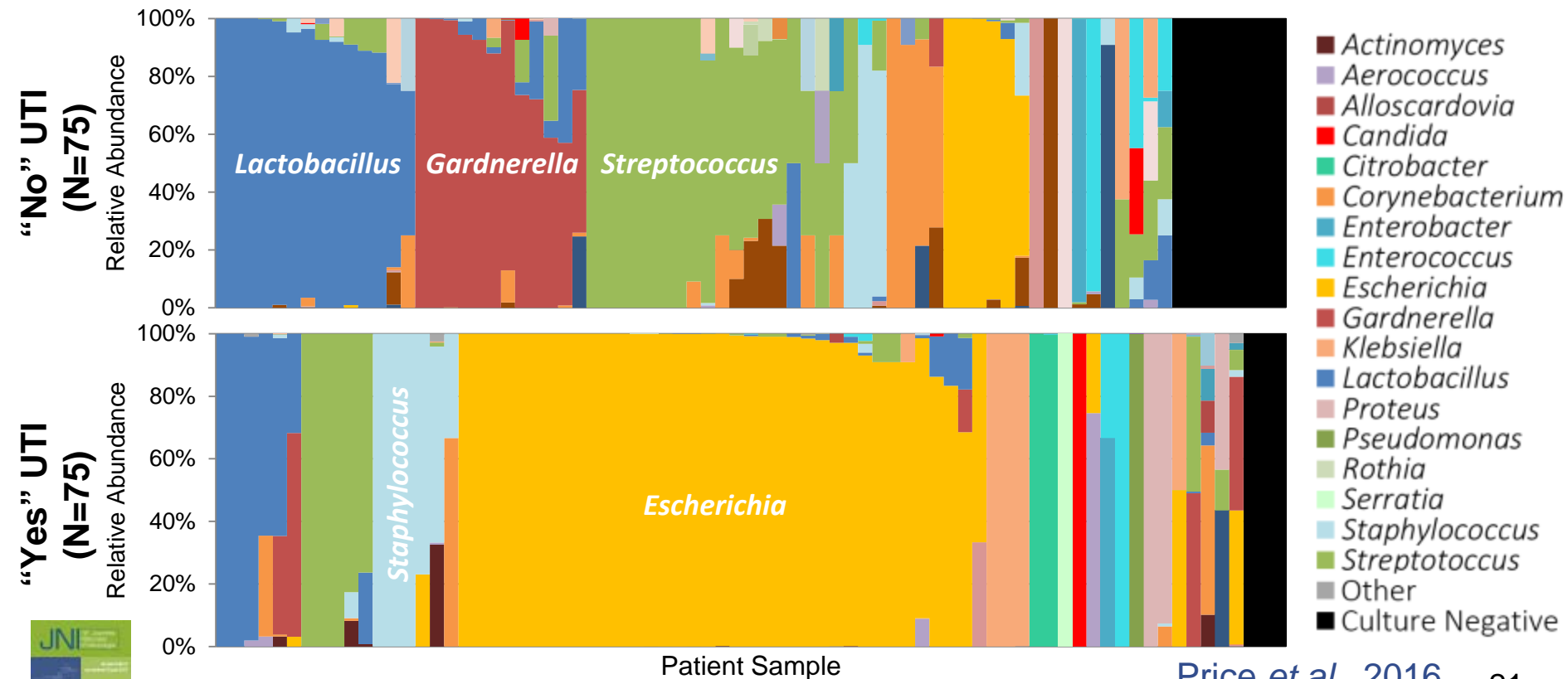
- Urgency
- Frequency
- Dysuria
- Difficulty Urinating
- Abdominal/Pelvic Pain/Pressure
- Low Back Pain
- Blood in Urine

Price *et al.*, 2016  
Clayson *et al.*, 2005

# Study design

Protocol (Volume)	Media	Condition	Identification
SUC (1 $\mu$ L)	Blood, MacConkey	Aerobic 35C	24 hr
Expanded EQUC (1, 10, 100 $\mu$ L)	Blood, Chocolate, CNA	5% CO <sub>2</sub> 35C	24 hr 48 hr
	Blood, MacConkey	Aerobic 35C	24 hr 48 hr
	Anaerobic	Anaerobic 35C	48 hr
	Anaerobic	Microaerophilic (5% O <sub>2</sub> , 10% CO <sub>2</sub> , 85% N) 35C	48 hr

# Composition differs at the genus level



# Hypothesis

The standard definitions and measures of UTIs are inefficient at detecting clinically relevant infections.

- **Standard Urine Culture (SUC) protocol**
  - Urinary Symptoms
  - $\geq 10^5$  CFU/mL for diagnosis

# SUC fails to detect 67% of uropathogens

Culturing Protocol	Uropathogens (N=182)
100ul Expanded EQUC	96% (N=174)
10ul Expanded EQUC	65% (N=118)
1ul Expanded EQUC	52% (N=95)
Standard Urine Culture (SUC)	<b>33% (N=60)</b>

## Uropathogens (SUC/EQUC):

*Actinobaculum schaalii* (0/6), *Aerococcus sanguinicola* (0/1), *Aerococcus urinae* (1/15),  
*Alloscardovia omnicoles* (0/8), *Candida albicans* (0/2), *Candida parapsilosis* (0/4), *Citrobacter freundii* (1/1),  
*Citrobacter koseri* (0/1), *Corynebacterium riegelii* (0/4), *Corynebacterium urealyticum* (0/2),  
*Enterobacter aerogenes* (1/3), *Enterococcus faecalis* (1/16), *Escherichia coli* (44/50),  
*Klebsiella pneumoniae* (4/10), *Morganella morganii* (0/1), *Oligella urethralis* (0/1), *Proteus mirabilis* (2/4),  
*Pseudomonas aeruginosa* (1/1), *Serratia marcescens* (0/1), *Staphylococcus aureus* (3/7),  
*Staphylococcus lugdunensis* (1/2), *Streptococcus agalactiae* (1/10), *Streptococcus anginosus* (0/32)

# SUC fails to detect 88% of non-*E. coli* uropathogens

Culturing Protocol	Detection of <i>E. coli</i> (N=50)	Non- <i>E. coli</i> Uropathogens (N=132)
100ul Expanded EQUC	100% (N=50)	94% (N=124)
10ul Expanded EQUC	94% (N=47)	55% (N=72)
1ul Expanded EQUC	92% (N=46)	36% (N=48)
Standard Urine Culture (SUC)	89%(N=44)	<b>12% (N=16)</b>

## Uropathogens (SUC/EQUC):

*Actinobaculum schaalii* (0/6), *Aerococcus sanguinicola* (0/1), *Aerococcus urinae* (1/15),  
*Alloscardovia omnicoles* (0/8), *Candida albicans* (0/2), *Candida parapsilosis* (0/4), *Citrobacter freundii* (1/1),  
*Citrobacter koseri* (0/1), *Corynebacterium riegelii* (0/4), *Corynebacterium urealyticum* (0/2),  
*Enterobacter aerogenes* (1/3), *Enterococcus faecalis* (1/16), ***Escherichia coli* (44/50)**,  
*Klebsiella pneumoniae* (4/10), *Morganella morganii* (0/1), *Oligella urethralis* (0/1), *Proteus mirabilis* (2/4),  
*Pseudomonas aeruginosa* (1/1), *Serratia marcescens* (0/1), *Staphylococcus aureus* (3/7),  
*Staphylococcus lugdunensis* (1/2), *Streptococcus agalactiae* (1/10), *Streptococcus anginosus* (0/32)



# Hypothesis

The standard definitions and measures of UTIs are inefficient at detecting clinically relevant infections.

- Standard Urine Culture (SUC) protocol
  - **Urinary Symptoms**
  - $\geq 10^5$  CFU/mL for diagnosis

# Urinary symptoms and UTI diagnosis

## UTI Diagnosis:

### Self-Report

UTI: N=75

No UTI: N=75

## UTI Diagnosis:

### SUC ( $\geq 10^5$ CFU/mL)

UTI: N=57

No UTI: N=93

## UTI Diagnosis:

### EQUC ( $\geq 10$ CFU/mL)

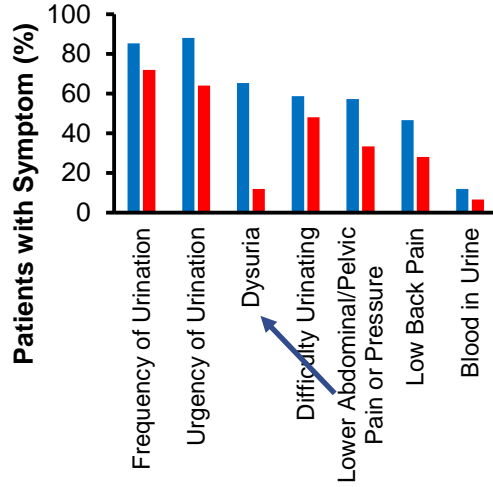
UTI: N=110

No UTI: N=40

# Dysuria is a strong indicator of UTI

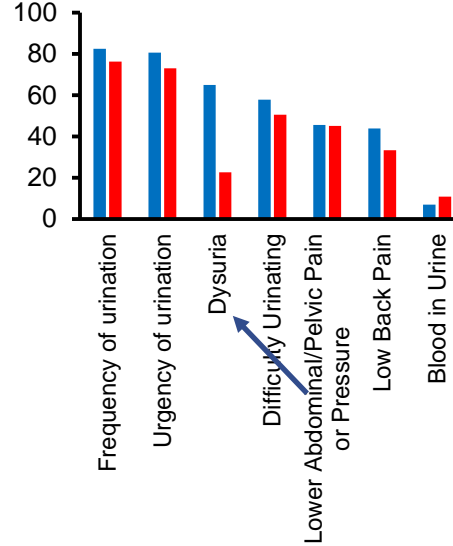
## UTI Diagnosis: Self-Report

UTI: N=75  
No UTI: N=75



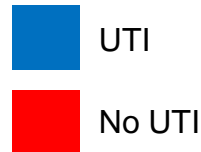
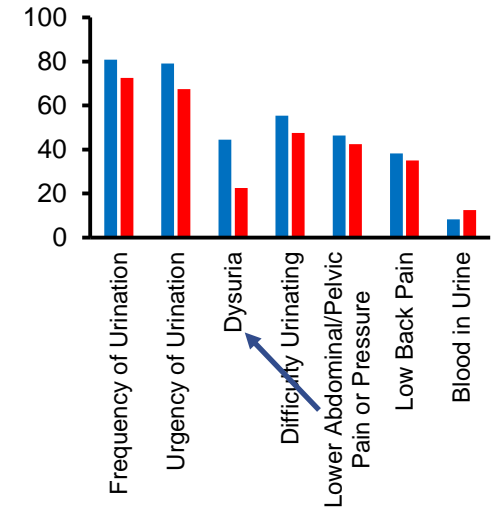
## UTI Diagnosis: SUC ( $\geq 10^5$ CFU/mL)

UTI: N=57  
No UTI: N=93



## UTI Diagnosis: EQUC ( $\geq 10$ CFU/mL)

UTI: N=110  
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Urinary Symptoms



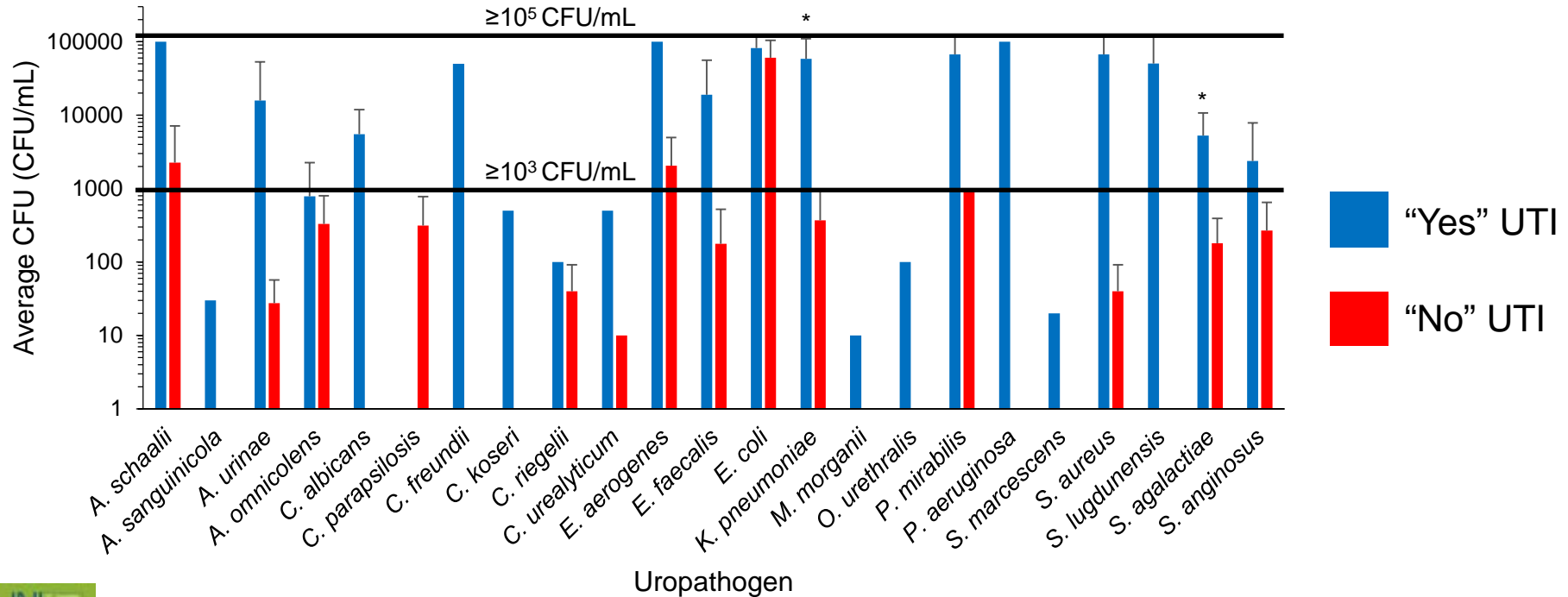
# Hypothesis

The standard definitions and measures of UTIs are inefficient at detecting clinically relevant infections.

- Standard Urine Culture (SUC) protocol
  - Urinary Symptoms
  - **$\geq 10^5$  CFU/mL for diagnosis**

# $\geq 10^5$ CFU/mL fails to detect most uropathogens

Average CFU for Cultivation of Uropathogens



# Conclusions

- **Use of Standard Urine Culture (SUC) is not a good indicator of UTI**
  - **Optimal protocol: 100 $\mu$ L; Blood, MacConkey, CNA; 5% CO<sub>2</sub>; 48 hrs**
- **Symptoms of Frequency and Urgency or urination are not good indicators of UTI**
  - **Presence of Pain and Burning during urination (Dysuria) is associated with multiple UTI definitions**
- **Use of a  $\geq 10^5$  CFU/mL threshold is not a good indicator of UTI**
  - **Lower/No threshold better detects possible Gram-positive and polymicrobial UTIs**
  - **Failure to treat infections  $< 10^5$  CFU/mL results in poor clinical outcome**

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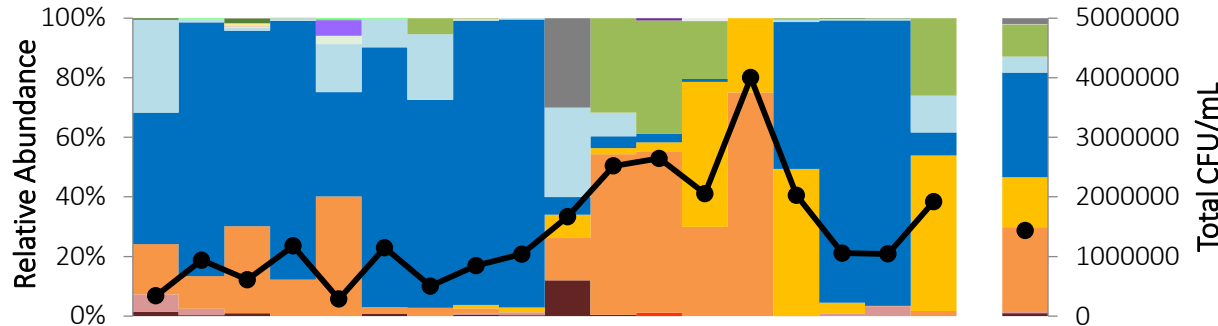
# Current projects

- Clinical trial to determine if treatment based off EQUC leads to improved patient outcome
- Measuring the effects of Estrogen on the FUM
- Determining if the FUM contributes to Interstitial Cystitis (IC/PBS)
- Measuring the stability of the FUM

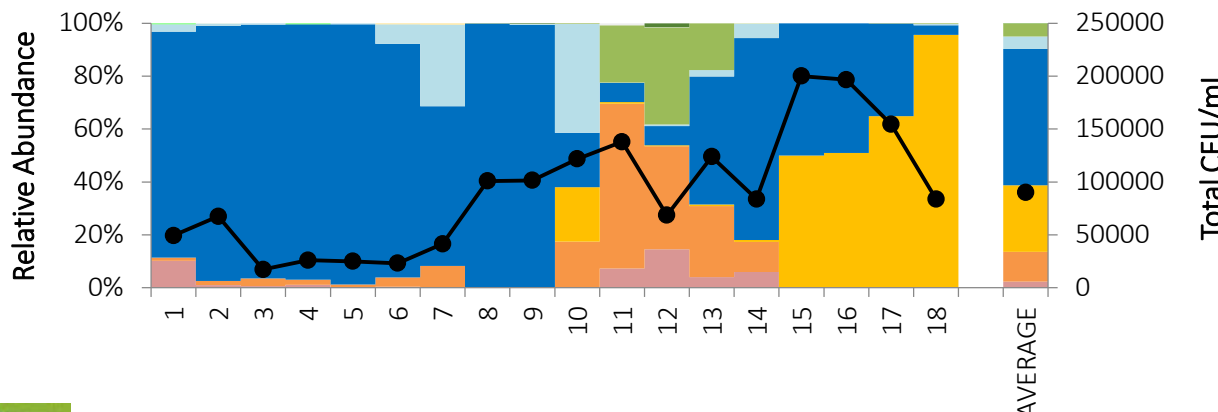


# Natural development of a UTI

Peri-urethral Swab



Midstream Voided Urine



Participant E

- *Actinomyces*
- *Alloscardovia*
- *Bifidobacterium*
- *Corynebacterium*
- *Enterobacter*
- *Enterococcus*
- *Escherichia*
- *Lactobacillus*
- *Micrococcus*
- *Propionibacterium*
- *Pseudomonas*
- *Staphylococcus*
- *Streptococcus*
- Unclassified
- Negative

Time (Days)



# Acknowledgments

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- Linda Brubaker, MD
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