# **Bacterial diversity and abundance near and far from drainage pipes** at DuPage County lakes

## **Bacteria and Water Quality**

The abundance and diversity of bacteria can be an indicator of water quality<sup>1</sup>. We hypothesized that water drainage pipes would be a vessel of pollutants from the surrounding roads to the water. This lead us to ask the question: Is there a difference in the abundance and diversity of bacteria near and far from drainage pipes?



Specifically, we wanted to measure the presence of indicator organisms. **Indicator organisms** are easily measured organisms that signal the presence of contamination and pathogens<sup>2</sup>. The two indicator organisms that we chose to measure were Escherichia coli, and Enterococci.

- Escherichia coli are gram-negative, facultative anaerobes, commonly found in the intestines of warm-blooded animals.
- *Enterococci* are gram-positive, facultative anaerobes. They are the leading cause of hospital-acquired secondary infection. They thrive in slightly basic conditions  $(pH > 7)^3$ .



m-Endo agar with *E. Coli* (left); m-Enterococcus agar with *Enterococci* (right)

We also wanted to measure **bacterial diversity**. We were curious to see what species of bacteria were living in the water at different locations.



## Materials and methods

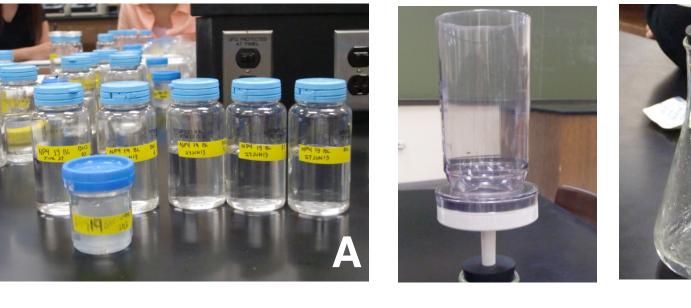
#### Collection

- Two (100 mL) water samples were taken at each location
- All locations were marked with GPS

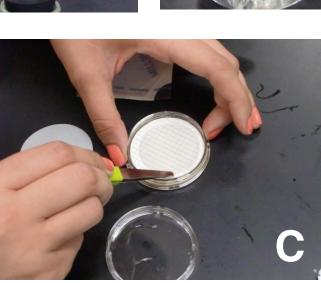


#### 2. Dilution & Filtration

- Water was diluted at ratios of 1:10, 1:100, 1:1,000<sup>A</sup>
- Diluted samples were filtered through a 45 µm filter using a hand pump<sup>B</sup>
- Filters were placed on agar plates and incubated<sup>C</sup>







#### **Total Aerobic Plate Count**

- BHI broth (permissive media)
- m-Endo agar (tests for *E. coli*)
- m-Enterococcus agar (tests for *Enterococci*)

#### To further identify bacteria species:

- EMB agar plate
- SD media
- Gram-negative bacteria
- 5. Slide mounts
- 6. Oxidase test
- Identifies aerobic bacteria
- Enterococcus test
- bile-esculin agar slant
- indole test
- BBL Enterotube II

Flooding slides with crystal violet dve

Self-contained test with 12 media and determines 15 biochemical reactions

### To test for bacterial diversity:

We used the Shannon-Wiener diversity index

$$H = -\sum_{i=1}^{S} p_i \ln p_i$$

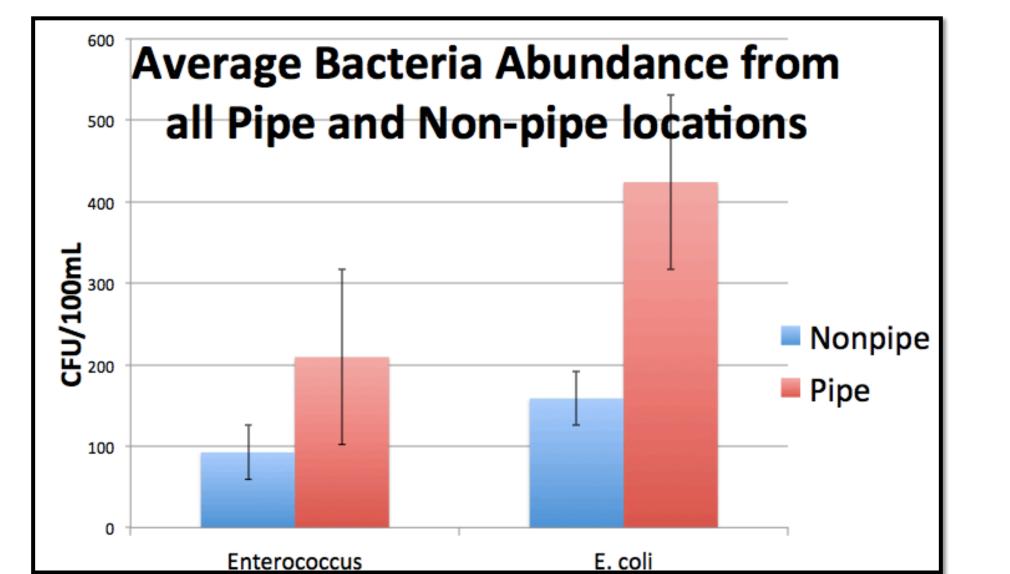
• H = Index P<sub>i</sub> = proportion of species

Ln = natural log (base e)

Eileen Colwell<sup>1</sup>, Vishal Patel<sup>1</sup> and Helena Puche<sup>2</sup> <sup>1</sup>DePaul University; <sup>2</sup> Harry S. Truman College

## Results

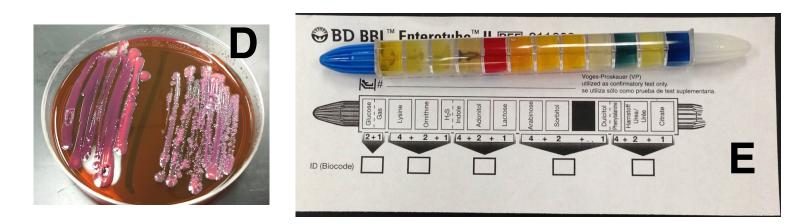
From our total aerobic plate count, we found that the numbers of *E. coli* and *Enterococci* were not statistically different between pipe and non-pipe locations (Kruskal Wallis ANOVA, H = 5.3; df = 4; P > 0.05).



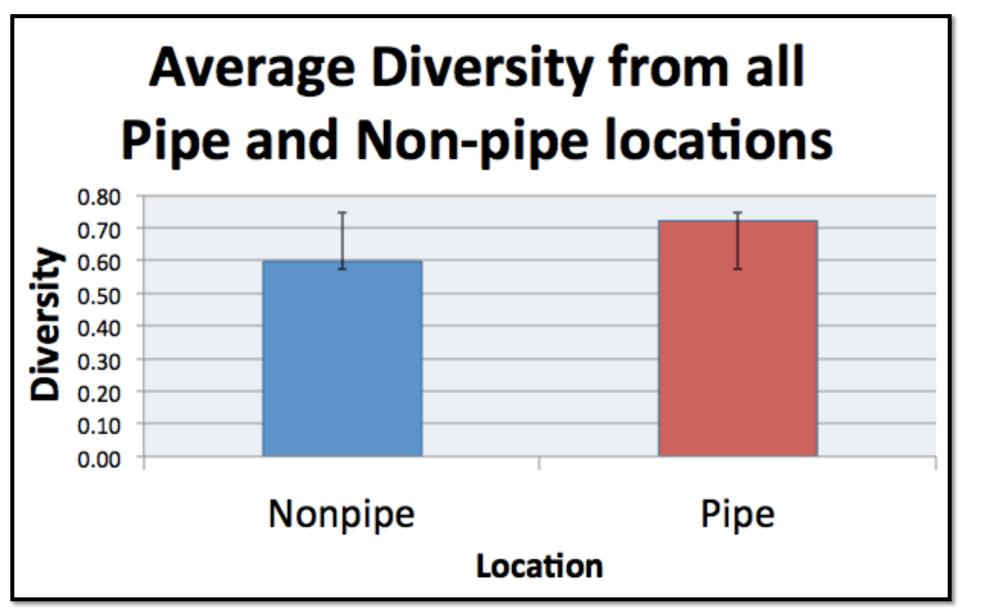
Average count of colony forming units (CFU) per 100 mL of Enterococci and E. coli at pipe and non-pipe locations.

From the results of our identification experiments (i.e. EMB agar plate<sup>D</sup>, slide mounts, oxidase test, *Enterococcus* test, and Enterotube<sup>E),</sup> We were able to confirm the presence of the following bacteria:

- Citrobacter freundii
- Enterococcus faecalis
- Enterobacter clocae



From our diversity calculations, the **diversity index between** pipe and nonpipe locations was not statistically different.



Average diversity of pipe and non-pipe locations, based on the Shannon-Wiener diversity index. A higher diversity index value indicates a more diverse location (Mann Whitney U test: U = 49.5; df = 1; P > 0.05).

## Conclusions

The abundance of E. Coli and Enterococci were not different between pipe and non-pipe locations. This lead us to ask the question: Are there other environmental variables that are shadowing the effects of drainage pipes?



- Several congregates of ducks and geese were observed at our test sites.
- We found *Enterobacter clocae*, a bacteria found in the intestines and fecal matter of birds.
- The presence or absence of bird species may have overshadowed contamination from water drainage pipes.

### Literature cited

- Ibekwe, A. M., Leddy, M. B., Bold, R. M., Graves, A.A. K. (2012). "Bacterial community composition in low-flowing river water with different sources of pollutants."
- FEMS Microbiology Ecology. 79: 155-166.
- 2. Wheeler Alm, E. (2003). "Fecal indicator bacteria are abundant in wet sand at freshwater beaches". Water Research, 37 (16): 3978. Blood, R. M. (1995). "Media for 'total' Enterobacteriaceae, coliforms and Escherichia coli". International Journal of Food Microbiology, 26 (1): 93.

### Acknowledgments

Thanks to Dr. Montgomery (DePaul University) for lending the Environmental Monitoring System, the Dept. of Environmental Sciences (DePaul University) for lending the GPS units, and Mr. Kevin Jankowski (Truman College) for guidance with bacteria. This is DuPage Forest Preserve permit 13-070.

### **Contact Information**

Eileen Colwell Vishal Patel Helena Puche eileenmcolwell@gmail.com vpatel56@mail.depaul.edu hpuche@ccc.edu