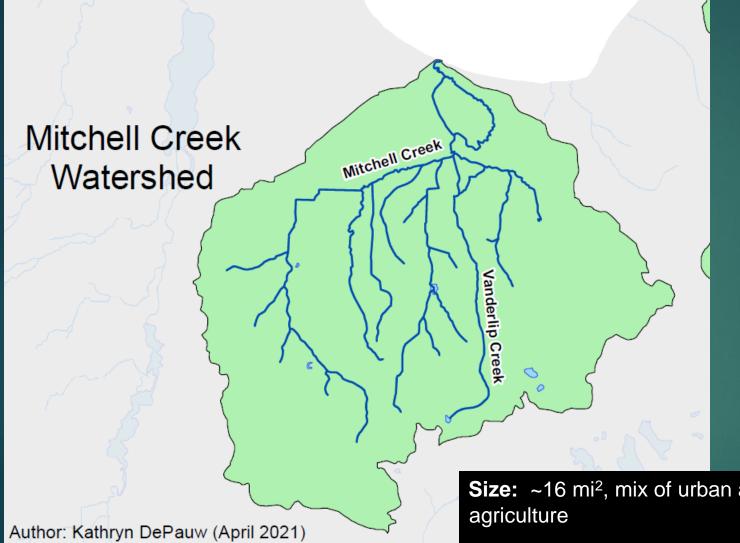
Mitchell Creek Source Tracking Project



Sarah U'Ren Program Director The Watershed Center

Mitchell Creek Impairment



On State Impaired Waters List (303(d) List) for:

Total Body Contact

Results of monitoring efforts by TWC in 2015
11 locations, weekly, 12 weeks

Size: ~16 mi², mix of urban and wetland land cover, with headwaters mostly agriculture

Issues:

- Elevated *E.coli* bacteria levels, not meeting State Water Quality Standards
- Potential public health issue, creek outlets next to State Park Beach

EGLE Grant

September 2020, awarded EGLE NPS grant to conduct microbial source tracking (MST) study to determine causes of elevated bacteria levels

Is it human or animal related? Are septic system contributing to problem? If animal, which one(s)?

Partners:



















Project Tasks

Surface Water Sampling 2021-2023

Run on ALL samples – markers for fecal contamination

- E.coli
- Somatic coliphage indicator virus
- F-specific coliphage indicator virus
- Clostridium perfringens

Microbial Source Tracking Analysis

Completed by MSU lab using ddPCR (digital droplet)

Run on all *E.coli* samples >300 col/100mL

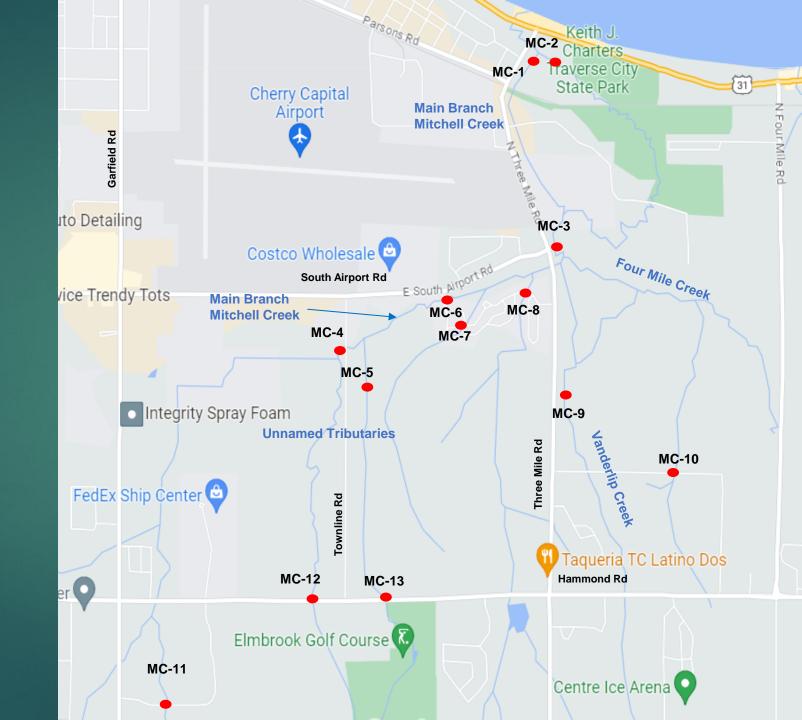
MST Markers* Tested:

- Human
- Cow
- Pig
- Dog
- Gull

*MST Markers:

Molecular genes which are detected primarily in the fecal bacteria group the Bacteroides

Specific to animal hosts

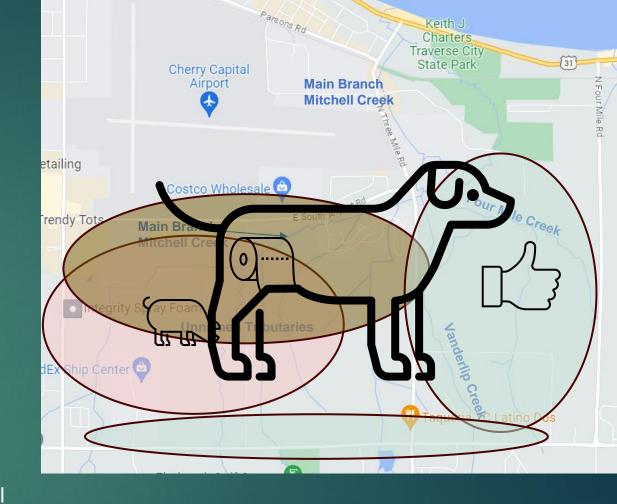


Results and Conclusions -Broad Overview

- Clear temporal pattern, rainfall driven
- Mix of fresh and old/aged sources (comparing Ecoli to other fecal indicators)
- Possible that Ecoli has accumulated in creek sediment and being released during rain events

Microbial Source Tracking:

- Little to no bacteria, fecal indicators, or MST markers found in eastern tribs of Vanderlip or Four Mile creeks or headwaters (Hammond Road and south)
- The western tributaries had more samples containing the pig marker
- Human marker was found in the western tributaries and central main branch
- Canine marker was widespread in all tributaries.



Fecal indicator and microbial source tracking (MST) analysis conducted in this study has not led to any obvious or consistent sources or locations of bacterial contamination, and it is difficult to determine remediation steps to reduce the impairment.

Results and Conclusions Human Markers & Septic Systems

2021-2022

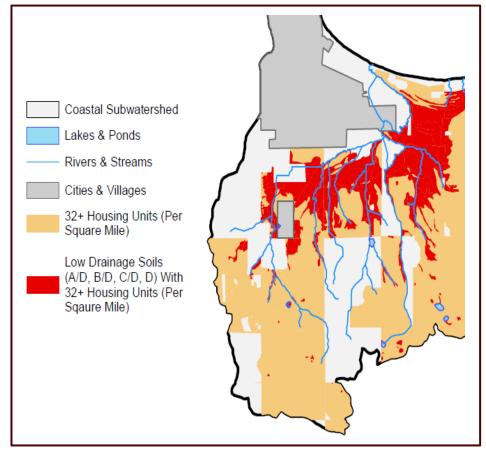
 Human markers were occasionally found in the western and central main branches of Mitchell Creek.

2023

- Expanded testing to include three upstream sites
- Included virus testing to determine potential public health impacts
- All tested sites in 2023 had no viruses or human markers present

Due to the sporadic and inconsistent findings for the human marker we cannot conclusively say at this time that leaking or failing septic systems are a consistent, major problem causing widespread impairment in the watershed.

Excerpt of Figure 25 from Coastal Grand Traverse Bay Watershed Plan – Areas At Risk for Septic Pollution



The Coastal Grand Traverse Bay Watershed Plan states that the Mitchell Creek area has a greater potential for septic system pollution than other areas in the watershed due to the density of systems combined with soil types

Results and Conclusions Pig Markers

- Found throughout, but most often in the unnamed western tributaries.
- Looking at pairings with fecal indicators, most of the time it was from older sources
 - Release from sediments
 - Land application of pig waste that had been properly composted in storage prior to being land applied.

Here's where it gets tricky...

- 2021 Found at nearly all the sites tested (dry and wet weather)
- 2022-2023 Virtually none found
- Met with resource professionals No large pig farms, but some small ones with pigs. However, all are following accepted BMPs for pig waste storage.
- No way to track land application of fresh or aged pig manure (who, where, how old, etc.).

However, if farms are following all BMPs for manure application, they are less likely to be affecting water quality in the creek and the marker may just be showing up through properly aged/composted runoff with a low bacterial and viral risk that is not harmful to human health





Results and Conclusions Other Sources

Cow/Gull

 Both cow and gull markers were virtually non-existent in this study and are not considered a threat at this time.

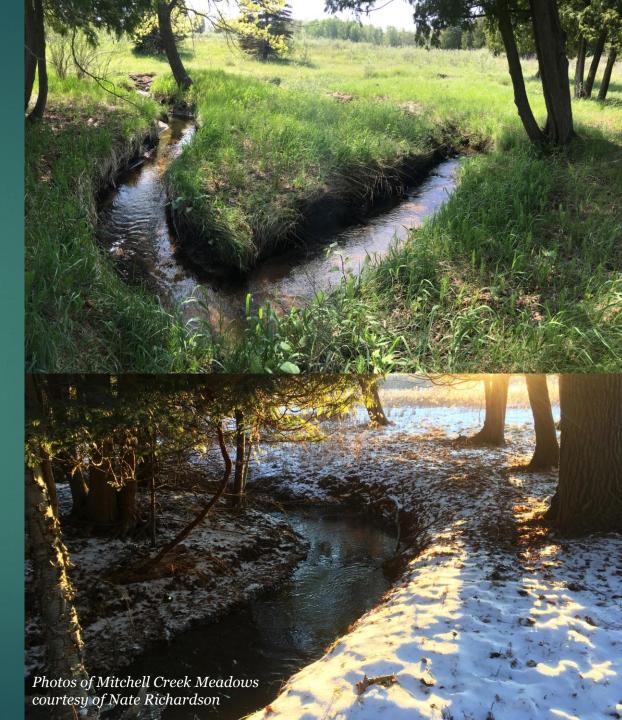


- The canine marker found in almost all samples tested.
- In addition to domesticated dogs, the marker may have also detected wild canines including coyote and fox.
- Therefore, we cannot confirm that fecal material from domestic dogs is a widespread concern for Mitchell Creek. If bacterial impairment is attributed to coyotes and foxes, remedying the issue will be difficult.



Recommendations

- Ensure all farms are following "Generally Accepted Agricultural Management Practices" (GAAMPs) and implement new agricultural BMPs to reduce fecal input to creek where needed.
- Provide septic system education, including maintenance, rightsizing, replacing aged systems, and proper use.
- Conduct continual maintenance of the City of Traverse City's sanitary system and extend service to new homes or developments, when possible.
- Provide education regarding the importance of picking up and properly disposing of dog waste.
- Preserve existing forested and vegetated wetlands that have the ability to reduce bacteria in the watershed and restore forested wetlands when possible.



Thank you!



Sarah U'Ren Program Director The Watershed Center <u>E.coli</u>: A bacteria that is the State of MI standard for recreational and impaired waters associated with fecal contamination. (this is what we use for beach testing)

Other indicators

- <u>Clostridium perfringens:</u> Bacteria, anaerobe, spore former, often used to indicate "older pollution" and materials resuspended from sediments
- F-Specific and Somatic Coliphage: Both are viruses found on *E.coli*. Somatic usually persists longer. Both may be indicative of recent fecal pollution because neither regrows in the environment.



Summary of Sample Collection

•	Added on to original
	sampling program

- Included three upstream sites
- Focused on human/pig sources

Date	Wet/Dry	Total Samples Collected (not including duplicates)	Total Samples Chosen for MST	MST Markers Run
7/12/2021	Dry	9	5	All*
8/4/2021	Dry	9	7	All*
9/20/2021	Dry	9	3	All*
10/4/2021	Wet	9	8	All*
10/18/2021	Dry	9	0	N/A
6/7/2022	Wet	9	9	All*
9/12/2023	Wet	10	8	B. theta and HF183PigViruses
10/25/2023	Wet	10	7	B. theta and HF183PigViruses
Totals	4 Dry 4 Wet	74 (36 dry, 38 wet events)	47 (15 dry, 32 wet events)	

^{*}MST markers include the following: human B. theta and HF193, pig, cow, canine, and gull *Human B. theta* marker indicates more recent pollution, more sensitive and degrades faster than Human HF183

Summary of Sample Collection

Date	Location											
	MC-1	MC-2	MC-3	MC-4	MC-5	MC-6	MC-8	MC-9	MC-10	MC-11	MC-12	MC-13
Phase 1 Sampling												
7/12/2021	326	743	289	403	519	284	449	141	150			
8/4/2021	310	306	305	425	442	414	480	243	91			
	Ź		,									
wet 6/7/2022 1,476 500 1,255 707 64' Phase 3 Sampling						1,000	1,000	7.5 0				
9/12/2023	1,278	577	1,093	611	1,058	880	1,183			126	78	816
10/25/2023				2,179		2,757				151		43
r of exceedances	6/8	6/8	5/8	7/8	7/8	5/8	7/8	2/6	0/6	0/2	0/2	1/2
Exceeds Daily Partial Body Contact Standard (300col/100mL)												
Exceeds Daily Total Body Contact Standard (1,000col/100mL)												
Percentage of Partial/Total Body Contact Exceedances by Type of Sample Event (Dry and Wet)												
ents	75%	75%	63%	88%	88%	63%	88%	25%	0%	0%	0%	50%
% of wet events % of dry events		100%	100% 25%	100% 75%	100% 75%	100% 25%	100% 75%	100%	0%	0% n/a	0% n/a	50% n/a
	7/12/2021 8/4/2021 9/20/2021 10/4/2021 10/18/2021 6/7/2022 8 Sampling 9/12/2023 10/25/2023 r of exceedances Exceeds Daily Parents ents	MC-1 MC-1 326 326 326 310	MC-1 MC-2	MC-1 MC-2 MC-3	MC-1 MC-2 MC-3 MC-4 Sampling	MC-1 MC-2 MC-3 MC-4 MC-5	MC-1 MC-2 MC-3 MC-4 MC-5 MC-6 Sampling	NC-1 NC-2 NC-3 NC-4 NC-5 NC-6 NC-8	MC-1 MC-2 MC-3 MC-4 MC-5 MC-6 MC-8 MC-9	NC-1 NC-2 NC-3 NC-4 NC-5 NC-6 NC-8 NC-9 NC-10 Sampling	NC-1 NC-2 NC-3 NC-4 NC-5 NC-6 NC-8 NC-9 NC-10 NC-11	NG-1 NG-2 NG-3 NG-4 NG-5 NG-6 NG-8 NG-9 NG-10 NG-11 NG-12