Version of Record:<https://www.sciencedirect.com/science/article/pii/S0048969721033039> Manuscript\_b0fbf94a4d6a97f0036cedf6ba9e8da9

# Integrating Culture and Molecular Quantification of Microbial Contaminants into a Predictive Modeling Framework in a Low-Lying, Tidally-Influenced Coastal Watershed

Matthew T. Price<sup>a</sup>, Angelia D. Blackwood<sup>a</sup>, Rachel T. Noble<sup>a</sup>

<sup>a</sup> UNC Institute of Marine Sciences, 3431 Arendell St., Morehead City, NC 28557, USA

Corresponding author: Rachel T. Noble Email: rtnoble@email.unc.edu Phone: (252) 726-6841; ext. 150

December 2020

**ABSTRACT** 

Examinations of stormwater delivery in the context of tidal inundation are lacking. Along the coastal plains of the southeast, tidal inundation is increasing in frequency and severity, often with dramatic adverse impacts on stormwater discharge and "sunny day flooding". Therefore, a comprehensive study was conducted to examine tidally-influenced stormwater outfalls discharging to Taylor's Creek, an estuary off the coast of Beaufort, NC used regularly for recreation and tourism. Over a wide range of meteorological conditions, water samples were collected and analyzed for fecal indicator bacteria (FIB, used for regulatory decision-making) and published quantitative microbial source tracking (qMST) markers. Nineteen sampling events were conducted from July 2017 – June 2018 with samples classified as inundated, receding or transition depending on collection during tidal stage. A first-of-its-kind multiple linear regression model was developed to predict concentrations of *Enterococcus* sp. by tidal cycle, salinity and antecedent rainfall. We demonstrated that the majority of variability associated with the concentration of *Enterococcus* sp. could be predicted by *E. coli* concentration and tidal phase. 16 FIB concentrations were significantly  $(50.05)$  influenced by tide with higher concentrations observed in samples collected during receding (low) tides (EC: log 3.12 MPN/100 mL; ENT: 2.67 MPN/100 mL) compared to those collected during inundated (high) (EC: log 2.62 MPN/100 mL; ENT: 2.11 MPN/100 mL) or transition (EC: log 2.74 MPN/100 mL; ENT: 2.53 MPN/100 mL) tidal periods. Salinity, was also found to significantly (<0.05) correlate with *Enterococcus* sp. concentrations during inundated (high) tidal conditions (sal: 17 ppt; ENT: 2.04 MPN/100 mL). Tide, not precipitation, was shown to be a significant driver in explaining the variability in *Enterococcus* sp. concentrations. Precipitation has previously been shown to be a driver of



#### **1. INTRODUCTION**

Stormwater runoff is one of the most important hydrological factors affecting surface water quality (Ahn et al., 2005; Mallin et al., 2009). Flowing directly overland, stormwater picks up pollutants including potentially pathogenic bacteria and viruses from animal and human waste (Griffin et al., 2003; Haile et al., 1999; Mallin et al., 2000; Prüss, 1998). Often times, this runoff enters stormwater conveyance systems that then carry the untreated runoff into downstream waterbodies, adversely impacting water quality and health for primary contact recreators. The United States (US) Environmental Protection Agency (US EPA) has recommended the use of enterococci (ENT) and *Escherichia* coli (EC) as fecal indicator bacteria (FIB) to monitor both marine and fresh surface waters (US EPA, 2012). FIB serve as a proxy for the presence of microbial pathogens associated with feces. Ingesting water with high concentrations of FIB through recreation can lead to gastrointestinal and other illnesses (Colford et al., 2007; Haile et al., 1999; Soller et al., 2017). Additionally, FIB have been selected due to their low pathogenic potential and high concentrations in sewage and feces (Ahmed et al., 2008; Ahmed et al., 2019; Harwood et al., 2014; Sidhu et al., 2012). As such, FIB have been widely used by states as a mitigation tool to meet US EPA water quality requirements. States have the discretion, however, to implement either or both FIB in monitoring programs. The State of North Carolina (NC) utilizes enterococci solely to monitor recreational surface waters (NC DEQ, 2020). While studied significantly across coastal waters, one major drawback towards the use of FIB, however, is their lack of source-specificity (Ex. human vs. non-human) regarding fecal contamination. As such, quantitative microbial source tracking tools (qMST) have been proposed. Quantitative microbial source tracking methods aim to discriminate between human and

non-human fecal sources in contaminated waterbodies (Lee et al., 2020; Nguyen et al., 2018;



much of the coastal zone in NC has a low topographic slope increasing at less than 0.09 m elevation for every horizontal mile (Corbett et al., 2008). As such, coastal NC remains susceptible to the effects of global climate change, including sea level rise, intensifying extreme storm events and increasing tidal ranges and sunny-day flooding (Hino et al., 2019). Sea level off the NC coast has increased 0.28 m as compared to 1950. The rate of rise accelerating over the last decade to now increasing by over 0.03 m every 2 years (NOAA, 2020; NC Coastal Resources Commission, 2015). This coupled with increased nuisance flooding frequency events suggest coastal surface waters along the coast of NC are at risk for continual impairment (King Tides Project, 2020; Sweet et al., 2014).





**Figure 1: Digital Elevation Model (DEM) depicting elevation in coastal, eastern NC and sampling area.** 



#### **2. MATERIALS AND METHODS**

## **2.1 Study Sites and Sample Collection**

Water samples were collected at three sampling locations throughout Beaufort (Figure 2): two at stormwater outfall locations (Orange St. and Marsh/Pollock) proximal to downstream receiving waters (Taylor's Creek) and a third site (Ann St.) one block inland. The two downstream locations were selected to underline the performance of the stormwater conveyance system, while the inland site was selected to characterize upstream watershed conditions. Following a land survey campaign, the Orange St. and Marsh/Pollock outfall sites were found to be the only two with an above-ground end-of-pipe access point and, as such, were selected as sampling locations. Nineteen sampling events were conducted seasonally over the course of 11 months from July 2017 – June 2018, with samples collected during both storm and ambient conditions. Storm sampling was initiated after a sustained period of moderate to heavy rainfall which produced accumulation of at least ∼0.25 in until ∼1 h after the storm ended. Dry weather samples were collected following three days without rainfall accumulation.

 Raleigh **Cape Hatteras**  Atlantic Ocean Wilmington



- **Figure 2: Three sampling locations: Orange St. (OS) and Marsh/Pollock (MP) are located adjacent to Taylor's Creek while Ann St. (AS) is one block inland.**
- Samples were collected using both an automatic and grab sampling approach. Automatic grab sampling was conducted using an ISCO 6712 Portable Sampler where composite samples
- were collected every 3 hours and stored for up to 6 hours before processing. Following
- collection, samples were stored on ice and transported to the laboratory where they were
- analyzed within 2 hours of collection.

# **2.2 Environmental Parameters**

Water temperature, total suspended solids (TSS) and salinity were measured in situ using a YSI probe (YSI 6600 multiparameter probe, USA). Additionally, meteorological observations (Ex. 24-h antecedent rainfall, tidal height and air temperature) were collected from publicly available data provided by NOAA: Station (ID: 8656483). We were able to determine the relative meteorological conditions by rounding sample collection time to the nearest NOAA sampling point (6-minute increments). 

#### **2.3 Tidal Characterization**

163 Similar to methods conducted in Boehm & Weisberg (2005), samples were classified into three tidal categories (Ex. receding, inundated and transition) classified by collection time as it related to the nearest recorded high tide. Given the semi-diurnal nature of tides within our system, samples were separated into three tidal categories: inundated (high tide), receding (low tide) or transition. Inundated samples were classified so if they had been collected within 2 hours of the previous high tide, while receding samples were collected >4 hours from the previous high tide. Transition samples were those collected in between the two groups (2-4 hours from nearest high tide). In addition, GPS locations and elevations were collected (Table 1) using a Trimble R8 171 RTK GPS relative to NAVD88 where average vertical error was  $\pm$  1.2 in. Outfall elevations were then used to verify coverage given NOAA verified tidal recordings.

**Table 1: Latitude, longitude, elevation and pipe size for OS, MP and AS sampling locations**

<b>Site</b>	Latitude	Longitude	Elevation (m, <b>NAVD88)</b>	<b>Pipe Radius</b> (m)
<b>Orange Street</b>	34.71751	-76.66740	0.105	0.3
<b>Marsh/Pollock</b>	34.71454	$-76.66190$	$-0.515$	0.43
<b>Ann Street</b>	34.71613	$-76.66070$	0.446	0.46

#### **2.4 Sample Preparation**

177 FIB *E. coli* and enterococci were enumerated using Colilert-18<sup>®</sup> and Enterolert™ at a 1:10 dilution (sample: DI water) per manufacturer instructions (IDEXX Laboratories, Westbrook, ME). For downstream molecular analysis, triplicate 100-150 mL samples were vacuum filtered through 0.45 μm pore size, 47 mm polycarbonate (PC) filters (HTTP, Millipore, Bedford, MA)

using a six-place filtration manifold and vacuum pump assembly. The filters were placed into sterile, DNase/RNase-free microcentrifuge tubes and stored at -80 °C. DNA extractions were 183 performed using the NUCLISENS<sup>®</sup> MINIMAG<sup>®</sup> extraction kit per manufacturer instructions, 184 with extracts then stored at -20 °C. Assays were performed in a CFX96 Touch<sup>TM</sup> Real-Time PCR Detection System (Bio-Rad Laboratories., Hercules, CA) with the following cycling conditions: 186 10 min at 95 °C, followed by 40 cycles of 15 s at 95 °C and 1 min at 60 °C. Extracted samples 187 were processed using TaqMan<sup>®</sup> Environmental Master Mix 2.0 (Applied Biosystems, Waltham, 188 Massachusetts). Primers (100  $\mu$ M) and probes (10  $\mu$ M) were synthesized by LGC Biosearch 189 Technologies (Petaluma, CA). Each reaction had a total volume of 25 µL, 20 µL including nuclease-free water,  $TaqMan^{\circledR}$  Environmental Master Mix 2.0, as well as appropriate primers and probes, and 5 µL of unknown sample, standard, or control. No template controls (NTCs) were processed with every plate.

### **2.5 Assessment of qPCR Specimen Processing Control and Inhibition Control**

Performance of the qPCR assays through evaluation of recovery efficiency and qPCR inhibition was measured using β actin (*ACTB*) cDNA as a specimen processing control (SPC) as previously conducted by Conn et al. (2012). 5 µL of *ACTB* solution (4000 copies/µL) was pipetted into each of the samples, calibrators, and negative controls prior to processing. Following this, samples were extracted. Inhibition was determined by calculating the difference between the cycle threshold (Ct) of the SPC in samples with (experimental) and without (control, only SPC) target DNA. Extracts were analyzed without dilution with samples having more than 0.5 log units (2.32 Ct) difference from control samples deemed inhibited (Lambertini et al., 203 2008). Since the total number of inhibited samples (11 out of 167 samples) constituted only 6.6% of total samples inhibited, no adjustment for inhibition was made. For all qPCR runs, appropriate controls were employed and showed no contamination: no template control (omission of DNA template from the qPCR reaction), and negative extractions control (inclusion of filter blank during DNA extraction). Plasmid standards were used for HF183 and Entero1-qPCR assays. Standards were synthesized by GenScript (Piscataway, NJ). Gene sequences were synthesized and inserted into a linearized pUC57 vector which was cloned into DH5α competent cells. Plasmids were extracted using Wizard® Plus SV 10 Minipreps DNA Purification System (Promega Corp., Madison, WI) and linearized using Eco R1 digestion. They were then confirmed via a 1% agarose gel in Tris-Acetate-EDTA buffer. The weight of purified plasmids was then calculated spectrophotometrically (Nanodrop 2000c, Thermo Scientific, Waltham, MA). Nanograms of plasmids were transformed to copy number by using a copy number calculator (SciencePrimer.com). Linearized plasmids were diluted and stored at a concentration 216 of  $1 \times 10^8$  copies per uL at -20 $^{\circ}$ C.

## **2.6 Standard Curves**

Standard curves for HF183 and Entero1-qPCR consisted of the calibration standard and five 10-fold serial dilutions that were run in triplicate. For each of the molecular markers, standard dilution curves were aggregated to form a singular curve. The theoretical limit of detection (LOD) was the lowest concentration where the standard could be detected reliably in at least 50% of qPCR replicates. The limit of quantification (LOQ) for qPCR assays was defined as the lowest concentration above the lowest point on the standard curve where amplification was observed in at least 50% of qPCR replicates.

## **2.7 Multiple Linear Regression Models**

Predictive modeling was also incorporated in the form of MLR models, which serve as a statistical technique that uses several explanatory variables to predict the outcome of a response variable. For the purposes of our study, enterococci consistently served as our response variable, given its regulatory importance in surface water quality monitoring in NC. Additionally, FIB *E. coli* and 24-h antecedent rainfall were incorporated with three tidal variables: tidal height (TH), tidal phase (TP) and tidal cycle (TC). Tidal height was incorporated using verified tidal height data recorded by NOAA, while the tidal phase variable incorporated distance the sample was taken from the nearest high tide. An additional variable accounting for tidal cycle was also included in regression analysis. This was done using the sine and cosine functions to characterize the cyclical nature of tides:  $\sin(2 \times \pi \times \frac{Minutes from high tide}{T_{\text{total}} + T_{\text{initial}} + T_{\text{total}} + T_{\text{initial}} + T_{\text{total}} + T_{\text{initial}}}$  $\sin(2 \times \pi \times \frac{\text{Minutes from high tide}}{\text{Total minutes between high tide}}))$  Cos(2 x π x ( 
240  $\cos(2 \times \pi \times (\frac{Minutes from high tide}{Total minutes between high tides}))$  Tidal Cycle =  $\sin(2 \times \pi \times \frac{\text{Minutes from high tide}}{\text{Total minutes between high-tark}})$ Minutes from high tide<br>Total minutes between high tides<sup>)</sup>) + Cos(2 x  $\pi$  x ( $\frac{Minutes \ from \ high \ tide}{Total \ minutes \ between \ high}$ 242 Tidal Cycle =  $\sin(2 \times \pi \times (\frac{Minutes from high tide}{Total minutes between high tides})) + \cos(2 \times \pi \times (\frac{Minutes from high tide}{Total minutes between high tides}))$  Using the regression model formula: 246  $Y_i = \beta_0 + \beta_1 + \beta_2 x_1 + \beta_2 x_2$  248 where Y<sub>i</sub> is the log-transformed outcome ENT concentrations,  $\beta_k$  is the estimated coefficient (EC 249 concentration, 24-h antecedent rainfall and tidal height) for variables  $X_1$  (tidal phase) and  $X_2$ 

(tidal cycle). Including the aforementioned terms, the final regression model was as follows:

 $YENT = \beta_{EC} + \beta_{Rain} + (\beta_{Tidal Height} \times \beta_{Tidal Phase}) + (\beta_{Tidal Height} \times \beta_{Tidal Cycle})$  **2.8 Statistical Analysis**  Log10 concentrations between FIB and qMST markers and environmental parameters were compared using matched paired t-tests for lognormally distributed samples or the nonparametric Wilcoxon Ranks-Sum Test for samples that did not fit a lognormal distribution. Non-detect samples were assigned a value of 5 copies/100 mL (log 0.7) with significance level set at 0.05 for all analyses. Analyses were conducted in OriginPro 8.5 (OriginLab, Northampton, MA). **3. RESULTS 3.1 Summary Statistics**  In total, 137 samples were collected and analyzed using culture-based FIB enumeration, qPCR-based *Enterococcus* sp. enumeration and qMST marker enumeration using vetted, 276 published qPCR-based approaches. Concentrations of EC ( $log 0.7 - 4.94$  MPN/100 mL) and ENT (log 0.7 – 4.78 MPN/100 mL) were comparable to those of the molecular markers, HF183 (log 0.7 – 4.07 copies/100 mL) and *Enterococcus* sp. quantification via qPCR (log 0.7 – 5.03 copies/100 mL). Significant correlations were observed across combinations of FIB and qMST 280 markers with significant positive correlations found between ENT and EC (r: 0.65;  $p \le 0.01$ ), 281 Entero1-qPCR (r: 0.71; p < 0.01) and HF183 (r: 0.45; p < 0.01). In an attempt to understand stormwater conveyance as it relates to tidal cycle, samples were collected over a wide range of precipitation and tidal conditions (Figure 3). On average, log EC and ENT concentrations in samples collected during storm events were 2.90 and 2.39 MPN/100 mL respectively, compared to average concentrations of 2.41 and 2.14 MPN/100 mL respectively during dry conditions. This was also true for qMST markers as HF183 and *Enterococcus* sp. quantified via qPCR were also found at mean higher concentrations in samples







300	Samples were collected across a diverse range of environmental conditions (Table 2), with
301	salinity measurements indicating an array of samples were collected across both storm and tidal
302	variations, as these values ranged from 0-35 parts per thousand (ppt). This suggests periods of
303	both fresh, stormwater inundation and marine, creek water inundation were included in overall
304	analysis. Additionally, a wide range of water temperatures that ranged from $9.0^{\circ}$ C during the
305	winter months, to 28 <sup>o</sup> C during the summer months, indicate seasonality was also considered in
306	sample collection.

**Table 2: Summarized data for environmental parameters: salinity (ppt), TSS (mg/L), water temp. (<sup>o</sup>C), 24h antecedent rainfall (in), tidal height (m) and air temp. ( <sup>o</sup>C) across the three sampling sites (OS, MP and AS).** 



#### **3.2 Inter-Site Variability**

On average, mean FIB and qMST marker concentrations where consistently higher at AS compared to those at the OS and MP locations (Table 3). Concentrations of EC*,* ENT and Entero1-qPCR concentrations at the upstream, inland AS location averaged 3.62 MPN/100 mL, 3.10 MPN/100 mL and 3.96 copies/100 mL respectively, compared to average values of 2.15 MPN/100 mL, 1.76 MPN/100 mL and 2.19 copies/100 mL at OS and 2.69 MPN/100 mL, 2.39 MPN/100 mL and 3.08 copies/100 mL at MP. The distributions of qMST marker and FIB marker concentrations measured across the sample sites were skewed, with relatively low average EC and ENT concentrations observed for the two downstream locations (OS and MP),



**Table 3: Summarized data for EC, ENT, HF183 and Entero1-qPCR concentrations at sampling sites (Orange St., Marsh/Pollock and Ann St.) including the distribution and prevalence of samples that exceeded recreational contact standards.** 

330



Samples collected at the AS location consistently exceeded recommended concentrations for both culture- and qPCR-based quantification of FIB concentration. For ENT, 79% of samples collected during all environmental conditions exceeded the NC Department of Environmental Quality (DEQ) state threshold of 104 MPN/100 mL. This was also true when samples were analyzed for concentration of Entero1-qPCR, which exceeded US EPA recommended criteria in approximately 83% of samples. When we compare these exceedances to the two downstream locations, which are influenced more greatly by tidal inundation, exceedance of FIB concentrations decreases. FIB exceedances were lowest at the OS outfall with approximately 32% and 27% of samples exceeding recommended EC and ENT concentrations respectively. This compares to an exceedance rate of 14% for samples analyzed for ENT concentrations via qPCR. HF183 concentrations, which are specifically associated with human fecal sources, only exceeded suggested thresholds (4200 copies per 100 mL, (Boehm et al., 2015)) in approximately one-third of samples at AS and MP with fewer samples (15%) exceeding suggested thresholds at OS.

## **3.3 Tidal Characterization**

Descriptive statistics were calculated across sample sites as characterized by collection time within the tidal cycle (Table 4). Across the three tidal categories (inundated, transition and receding), FIB and qMST marker concentrations were consistently higher at the AS location when compared to the two downstream sites: OS and MP. FIB and qMST marker concentrations were compared across tidal classifications using one-way ANOVA calculations with only EC 352 concentrations significantly  $(p<0.05)$  differing between inundation and receding tidal periods. The same analyses were performed between FIB characterized by sites across the different tidal phases. At OS, significant (p<0.05) differences were found between ENT and HF183





**Table 4: Descriptive statistics of FIB characterized by tidal cycle (inundated, receding or transition) sampling location.** 

360

359

A representative number of samples were collected across the tidal cycle in order to better represent FIB and qMST marker concentrations in the context of storm events and ambient (dry) conditions. Across the three tidal classifications, correlation coefficients were determined between ENT concentrations and EC, *Enterococcus* sp. concentrations determined via qPCR and HF183. A similar analysis was conducted with environmental parameters such as water temperature, salinity and TSS. Regardless of tidal cycle, ENT concentrations were found to significantly (p<0.05) correlate with other FIB concentration and qMST marker concentration,

368 regardless of enumeration approach (culture vs. molecular). Only salinity measurements  $(r = -$ 369 0.448, p-value =  $0.042$ ) revealed a significant relationship, with regards to the environmental parameters measured, indicating negative correlation with ENT concentrations only during periods of tidal inundation.

**3.4 Multiple Linear Regression Models** 

Three models in total were created to predict concentrations of ENT in a tidally-influenced estuarine system. The models were created using data from all sampling locations, however only 375 the two downstream location (OS and MP) were significant ( $p \le 0.05$ ) in their prediction of variation in ENT concentrations; therefore, the models are appropriate for locations regularly influenced by tidal inundation. For all three models, a combination of biological (EC concentrations) and environmental parameters (24-h antecedent rainfall, tidal height, tidal cycle and tidal phase) were found to maximize the ability to predict the observed variation in ENT concentrations explained. FIB and qMST markers, such as HF183 and *Enterococcus* sp. determined via qPCR, as well as environmental parameters, such as water temperature, salinity, TSS, 24h antecedent rainfall and water temperature, were considered when making a data training set. However, the five variables used in our models that consistently performed the best across the three sites, when compared to other data training sets. Models were evaluated by 385 comparing the p-value and adjusted  $R^2$  values. Table 5 summarizes the model performances for the pooled data from the three sites. The OS model demonstrated that 55% of its variation could be explained by five variables, with EC concentration and tidal phase and cycle exhibiting significant influences on ENT concentrations. Similar results were observed for the MP model with 63% of the variation in *Enterococcus* sp. concentration explained by the same variables. In this model, however, only EC concentration and tidal cycle were found to significantly

391 contribute to ENT concentrations. Interestingly enough, 24-h antecedent rainfall was not a

- 392 significant contributor to the variation observed in *Enterococcus* sp. concentrations for any of the
- 393 models.

394

395

**Table 5: Multiple regression model for the association of log10 Enterococci with biological and environmental characteristics by sampling location (Orange St., Marsh/Pollock and Ann St.). The regression model looks to better characterize the effect of tidal cycle on bacterial concentrations delivered with the system.**







#### **4. DISCUSSION**

 Historically, rainfall has long been associated with elevated FIB concentrations in receiving waters (Coulliette & Noble, 2008; Hart et al., 2020; Silva et al., 2014). However, the influence of tide on contaminant delivery during storms is poorly understood, particularly in low-lying coastal plain systems. This study evaluated the relationships of both culture- and qPCR-based FIB and qMST markers in the context of tidal cycle in an estuarine system exposed to stormwater delivery across a wide range of weather conditions. To further evaluate relationships observed for ENT, EC and qMST marker concentrations according to tide, we developed a MLR tool to better understand stormwater contamination dynamics in a complex, tidally-influenced estuarine system. MLR has been recommended as part of the US EPA 2012 "Update to the Recreational Water Quality Criteria," but to our knowledge there are no published models that incorporate tide. Predictive modeling tools have previously shown their utility in NC estuaries such as the one we studied (Gonzalez et al., 2012; Gonzalez & Noble, 2014) and therefore may be used to better serve coastal water quality managers by better explaining microbial contaminants in the context of tide and other environmental parameters. We hope to provide a framework for stormwater researchers needing to incorporate a tidal parameter in their monitoring regimes for the future, while also highlighting some of the major limitations associated with using such an approach.

# **4.1 Summary Statistics**

Samples were collected over a broad range of rainfall and dry weather conditions and across tidal cycles. While concentrations of FIB and qMST markers increased slightly during wet weather conditions, the concentrations were not significantly greater as compared to dry weather. Previous studies did find significant increases in FIB concentrations following rain events (Converse et al., 2011; Gonzalez et al., 2012; Parker et al., 2010; Stumpf et al., 2010), indicating the potential for a different driver of FIB and qMST marker concentrations. To analyze this further, inter-site variability was studied with regards to FIB and qMST marker concentrations. On average, the upstream, non-tidally impacted sampling location (AS) consistently had higher FIB and qMST marker concentrations compared to the downstream locations. We speculated that tidal inundation was impacting the downstream locations, but not the upstream location and was the factor dictating the observed differences in concentrations. Lewis et al., (2013) observed a decrease in FIB concentrations with increases in tide stage dependent on the extent of the tidal height. They concluded that tidal shifts exceeding 1.5 m within the tidal range resulted in decreased FIB concentrations as the system is inundated and diluted with seawater. Conversely, decreased tidal inundation was characterized by maximum inflows of freshwater which promote bacterial replication in systems with high concentrations of fecal contamination. This could explain why higher concentrations of FIB were observed at the AS location as compared to OS and MP. Findings from this study support the idea of a dilution effect on FIB and qMST marker concentrations related to tidal mixing causing both dilution and bacterial cell rupture during high tide events that ultimately reduces measured FIB concentrations (Chen et al., 2019; De Brauwere et al., 2011; Kirchman et al., 1984; Pednekar et al., 2005).

Environmental parameters validated the observed, shifting dynamics across the various tidal classifications. Salinity measurements were found to be the highest during periods of tidal inundation (17 ppt) compared to transition (10 ppt) and receding (16 ppt) tidal periods. While not significantly different than average values during low tide events, significant correlations to ENT concentrations during high tide suggest the potential utility of such a parameter as has been reported in previous research (Byappanahalli et al., 2012; Dorsey et al., 2010; Sinton et al., 2002). Neither TSS nor water temperature exhibited strong relationships with either FIB or qMST indicators. This could be attributed to fewer measurements collected over the course of the study, which was the result of evolving research goals that emerged as the complexity of the system became apparent.

# **4.2 Multiple Linear Regression Models**

To our knowledge, this was the first application of a MLR that incorporated both qualitative and quantitative tidal variables to examine drivers of microbial contaminant concentrations. This approach, when compared to other statistical methods, may serve as more appropriate tool to routinely evaluate stormwater-impacted water quality. As evidenced by the recent USEPA buy-in, predictive modeling tools (Cyterski et al., 2013) can offer an opportunity to identify drivers of contamination, especially as related to stormwater inputs, and environmental parameters. In the end, these tools conserve valuable resources by allowing predictions rather than routine sample collection and monitoring to manage recreational exposure and risk. Previous modeling done by Gonzalez et al., (2014) was conducted in a neighboring system and demonstrated successful application of MLR. In this study, however, no tidal variable was incorporated to explain variation in either EC or ENT concentrations. Furthermore, rainfall was found to be a significant driver of FIB concentrations. The utility of our study is the incorporation of both wellestablished biological parameters (Hamilton et al., 2017; Jin et al., 2004; Parker et al., 2010) with less-understood environmental influences, such as tidal condition.

ENT and EC have long shown co-occurrence within fecal waste natural environments, with some proposing that EC is a superior metric of fecal contamination given its specificity and relationship to human health (Cabelli et al., 1982; Soller et al., 2010). Therefore, the relevance of EC concentration within the model makes sense due to its known, previously published, positive correlation with ENT (Boehm & Sassoubre, 2014; Steele et al., 2018; Stumpf et al., 2010). Tidal cycle, however, which has been studied much less frequently, also appeared to exhibit great influence on ENT concentration variation. We believe this implies that contaminant transport is more dependent on the timing of storm events as they relate to the state of the tide, compared to simply the extent, intensity of the storm event itself. If this is true, downstream waters could be susceptible to impairment long after a storm event ceases and related to the release of the system as the tide retreats. Thus, contaminated waterways remain open during contamination events increasing the likelihood of deleterious public health effects (Leecaster & Weisberg, 2001; Noble, Blackwood, Griffith, McGee, & Weisberg, 2010). Furthermore, in this framework, antecedent rainfall patterns would carry increased weight and value to future predictive model development. This is because long periods of increased rainfall will begin to favor higher surficial groundwater levels, as well as decreased infiltration capacity, potentially driving a compounded issue of stormwater delivery hampered by localized increased tidal elevation due to increased localized runoff (Yau et al., 2014).

# **4.3 Application**

In low-lying, rural systems, such as Beaufort, NC, it is not uncommon to find some degree of spatial autocorrelation in water quality studies (Partyka et al., 2017; Tu & Xia, 2008) suggesting

that the qualities under investigation are determined somewhat by unmeasured, and possibly external factors. If these influences are not taken into consideration, bias can be introduced into microbial water quality monitoring programs and the subsequent management decisions. In this particular study, we considered tidal variation, which is surprisingly understudied. Coastal communities across the entire NC coast sit at elevations around or below those found in Beaufort (e.g. Currituck (7 ft), Hatteras (3 ft), and Ocracoke (3 ft)) and, as such, experience similar degrees of tidal inundation. By addressing this issue in more depth, stormwater researchers may have greater success in developing a more inclusive framework for stormwater management that may be applied in susceptible coastal communities throughout the US (Poulter et al., 2009; Pricope, Halls, & Rosul, 2019). We recognize the limitations of this study and the possible influence this may have on the reliability of model predictions. For instance, laboratory-based measures (e.g. salinity and TSS) not comprehensively conducted across all sample types throughout the study. Furthermore, it would have been of great interest to understand the elevation and pipe dimension and flow and discharge across the entire system, but these parameters were difficult to measure in practice and resulted in intermittent data collection. Additionally, sampling regimes varied between automatic and grab sampling, introducing bias related to sample collection frequency and type. Previous studies applying a tidal description in their sampling methods have primarily occurred during one tidal phase (Ex. low or high) which limits one's understanding of shifting FIB and qMST concentrations that change with the tide. Much of the previous literature shows geographic or socio-economic biases as many were conducted in the western US or in highly developed watersheds with lower tidal intrusion and greater financial resources to combat coastal flooding. With the greatest risks falling on low-

lying, rural populations, accurate classifications of tidal inundation and its impact on microbial contaminant delivery in stormwater is necessary for future consideration.

We understand there is no "one-size-fits-all" model for the prediction of *Enterococcus* sp. concentration in discharge to coastal, surface waters. However, once baseline interactions between environmental parameters and microbial dynamics have been established through routine monitoring, data can then be interpreted in the context of tide. Without reliable spatial and temporal knowledge of tidal cycle, we cannot fully rely on the results of published models to answer today's questions of acceptable water quality.

## **5. CONCLUSIONS**

• Concentrations of culture FIB (*E. coli* and enterococci), Entero1-qPCR and qMST (HF183) markers were significantly influenced by tide with higher concentrations found during receding (low) tides compared to those from inundated (high) or transition tidal periods.

526 • Environmental parameters, such as salinity, were found to significantly  $(p<0.05)$  correlate with ENT concentrations during periods of tidal inundation. Salinity is likely a valuable conservative marker for future dispersion studies.

• Study successfully showed the application of MLR using qualitative and quantitative tidal

variables as driver of variation in both EC and ENT concentrations. However, 24-h

antecedent rainfall was not determined to have a major influence on FIB concentration as has

been previously reported.



#### **7. REFERENCES**

- Ahmed, W., Stewart, J., Powell, D., & Gardner, T. (2008). Evaluation of Bacteroides markers for the detection of human faecal pollution. *Letters in Applied Microbiology*, *46*(2), 237–242. https://doi.org/10.1111/j.1472-765X.2007.02287.x
- Ahmed, W., Hamilton, K., Toze, S., Cook, S., & Page, D. (2019). A review on microbial contaminants in stormwater runoff and outfalls: Potential health risks and mitigation strategies. In *Science of the Total Environment* (Vol. 692, pp. 1304–1321). https://doi.org/10.1016/j.scitotenv.2019.07.055
- Ahmed, W., Payyappat, S., Cassidy, M., Harrison, N., & Besley, C. (2020). Sewage-associated marker genes illustrate the impact of wet weather overflows and dry weather leakage in urban estuarine waters of Sydney, Australia. *Science of the Total Environment*, *705*, 135390. https://doi.org/10.1016/j.scitotenv.2019.135390
- Ahn, J. H., Grant, S. B., Surbeck, C. Q., Digiacomo, P. M., Nezlin, N. P., & Jiang, S. (2005). Coastal water quality impact of stormwater runoff from an urban watershed in Southern California. *Environmental Science and Technology*, *39*(16), 5940–5953. https://doi.org/10.1021/es0501464
- Al Aukidy, M., & Verlicchi, P. (2017). Contributions of combined sewer overflows and treated effluents to the bacterial load released into a coastal area. *Science of the Total Environment*, *607*–*608*, 483–496. https://doi.org/10.1016/j.scitotenv.2017.07.050
- Aragonés, L., López, I., Palazón, A., López-Ubeda, R., & García, C. (2016). Evaluation of the quality of coastal bathing waters in Spain through fecal bacteria Escherichia coli and Enterococcus. *Science of the Total Environment*, *566*–*567*, 288–297. https://doi.org/10.1016/j.scitotenv.2016.05.106
- Arnold, B. F., Wade, T. J., Benjamin-Chung, J., Schiff, K. C., Griffith, J. F., Dufour, A. P., Weisberg, S. B., & Colford, J. M. (2016). Acute gastroenteritis and recreational water: Highest burden among young US children. *American Journal of Public Health*, *106*(9), 1690–1697. https://doi.org/10.2105/AJPH.2016.303279
- Badgley, B. D., Steele, M. K., Cappellin, C., Burger, J., Jian, J., Neher, T. P., Orentas, M., & Wagner, R. (2019). Fecal indicator dynamics at the watershed scale: Variable relationships with land use, season, and water chemistry. *Science of the Total Environment*, *697*, 134113. https://doi.org/10.1016/j.scitotenv.2019.134113
- Bichai, F., & Ashbolt, N. (2017). Public health and water quality management in low-exposure stormwater schemes: A critical review of regulatory frameworks and path forward. *Sustainable Cities and Society*, *28*, 453–465. https://doi.org/10.1016/j.scs.2016.09.003

Boehm, A. B., & Weisberg, S. B. (2005). Tidal forcing of enterococci at marine recreational beaches at fortnightly and semidiurnal frequencies. *Environmental Science and Technology*, *39*(15), 5575–5583. https://doi.org/10.1021/es048175m Boehm, A. B., & Sassoubre, L. M. (2014). Enterococci as Indicators of Environmental Fecal Contamination. In *Enterococci: From Commensals to Leading Causes of Drug Resistant Infection*. Retrieved from http://www.ncbi.nlm.nih.gov/pubmed/24649503 Boehm, A. B., Soller, J. A., & Shanks, O. C. (2015). Human-Associated Fecal Quantitative Polymerase Chain Reaction Measurements and Simulated Risk of Gastrointestinal Illness in Recreational Waters Contaminated with Raw Sewage. *Environmental Science and Technology Letters*, *2*(10), 270–275. https://doi.org/10.1021/acs.estlett.5b00219 Byappanahalli, M. N., Nevers, M. B., Korajkic, A., Staley, Z. R., & Harwood, V. J. (2012). Enterococci in the Environment. *Microbiology and Molecular Biology Reviews*, *76*(4), 685– 706. https://doi.org/10.1128/mmbr.00023-12 Cabelli, V. J., Dufour, A. P., Mccabe, L. J., & Levin, M. A. (1982). Swimming-associated gastroenteritis and water quality. *American Journal of Epidemiology*, *115*(4), 606–616. https://doi.org/10.1093/oxfordjournals.aje.a113342 Cabelli, V. J. (1983). Microbial indicator systems for assessing water quality. *Antonie van Leeuwenhoek*, *48*(6), 613–618. https://doi.org/10.1007/BF00399546 Chen, X., Wei, W., Wang, J., Li, H., Sun, J., Ma, R., Jiao, N., & Zhang, R. (2019). Tide driven microbial dynamics through virus-host interactions in the estuarine ecosystem. *Water Research*, *160*, 118–129. https://doi.org/10.1016/j.watres.2019.05.051 Climate Signals. (2016). *King Tides Flooding*. Retrieved from https://www.climatesignals.org/events/king-tides-flooding-november-2016 Coelho, M. P. P., Marques, M. E., & Roseiro, J. C. (1999). Dynamics of microbiological contamination at a marine recreational site. *Marine Pollution Bulletin*, *38*(12), 1242–1246. https://doi.org/10.1016/S0025-326X(99)00169-1 Colford, J. M., Wade, T. J., Schiff, K. C., Wright, C. C., Griffith, J. F., Sandhu, S. K., Burns, S., Sobsey, M., Lovelace, G., & Weisberg, S. B. (2007). Water quality indicators and the risk of illness at beaches with nonpoint sources of fecal contamination. *Epidemiology*, *18*(1), 27–35. https://doi.org/10.1097/01.ede.0000249425.32990.b9 Conn, K. E., Habteselassie, M. Y., Denene Blackwood, A., & Noble, R. T. (2012). Microbial water quality before and after the repair of a failing onsite wastewater treatment system adjacent to coastal waters. *Journal of Applied Microbiology*, *112*(1), 214–224. https://doi.org/10.1111/j.1365-2672.2011.05183.x

- Converse, R. R., Blackwood, A. D., Kirs, M., Griffith, J. F., & Noble, R. T. (2009). Rapid QPCR-based assay for fecal Bacteroides spp. as a tool for assessing fecal contamination in recreational waters. *Water Research*, *43*(19), 4828–4837. https://doi.org/10.1016/j.watres.2009.06.036
- 
- Converse, R. R., Piehler, M. F., & Noble, R. T. (2011). Contrasts in concentrations and loads of conventional and alternative indicators of fecal contamination in coastal stormwater. *Water Research*, *45*(16), 5229–5240. https://doi.org/10.1016/j.watres.2011.07.029
- Corbett, D., Walsh, J. P., Cowart, L., Riggs, S. R., Ames, D. V, & Culver, S. J. (2008). Shoreline Change Within the Albemarle-Pamlico Estuarine System North Carolina. Retrieved from http://hdl.handle.net/10342/2862
- Coulliette, A. D., & Noble, R. T. (2008). Impacts of rainfall on the water quality of the Newport River Estuary (Eastern North Carolina, USA). *Journal of Water and Health*, *6*(4), 473–482. https://doi.org/10.2166/wh.2008.136
- Cyterski, M., Brooks, W., Galvin, M., Wolfe, K., Carvin, R., Roddick, T., & Corsi, S. (2013). Virtual beach 3: User's guide. United States Environmental Protection Agency. EPA/600/R-13/311
- De Brauwere, A., De Brye, B., Servais, P., Passerat, J., & Deleersnijder, E. (2011). Modelling Escherichia coli concentrations in the tidal Scheldt river and estuary. *Water Research*, 45(9), 2724-2738. https://doi.org/10.1016/j.watres.2011.02.003
- Dorsey, J. H., Carter, P. M., Bergquist, S., & Sagarin, R. (2010). Reduction of fecal indicator bacteria (FIB) in the Ballona Wetlands saltwater marsh (Los Angeles County, California, USA) with implications for restoration actions. *Water Research*, *44*(15), 4630–4642. https://doi.org/10.1016/j.watres.2010.06.012
- Eregno, F. E., Tryland, I., Myrmel, M., Wennberg, A., Oliinyk, A., Khatri, M., & Heistad, A. (2018). Decay rate of virus and faecal indicator bacteria (FIB) in seawater and the concentration of FIBs in different wastewater systems. *Microbial Risk Analysis*, 8, 14-21. https://doi.org/10.1016/j.mran.2018.01.001
- Gonzalez, R. A., Conn, K. E., Crosswell, J. R., & Noble, R. T. (2012). Application of empirical predictive modeling using conventional and alternative fecal indicator bacteria in eastern North Carolina waters. *Water Research*, *46*(18), 5871–5882.
- https://doi.org/10.1016/j.watres.2012.07.050
- Gonzalez, R. A., & Noble, R. T. (2014). Comparisons of statistical models to predict fecal indicator bacteria concentrations enumerated by qPCR- and culture-based methods. *Water Research*, *48*(1), 296–305. https://doi.org/10.1016/j.watres.2013.09.038
- Green, H. C., & Field, K. G. (2012). Sensitive detection of sample interference in environmental qPCR. *Water Research*, *46*(10), 3251–3260. https://doi.org/10.1016/j.watres.2012.03.041
- Green, H. C., Haugland, R. A., Varma, M., Millen, H. T., Borchardt, M. A., Field, K. G., Walters, W. A., Knight, R., Sivaganesan, M., Kelty, C. A., & Shanks, O. C. (2014). Improved HF183 quantitative real-time PCR assay for characterization of human fecal pollution in ambient surface water samples. *Applied and Environmental Microbiology*, *80*(10), 3086–3094. https://doi.org/10.1128/AEM.04137-13
- Griffin, D. W., Donaldson, K. A., Paul, J. H., & Rose, J. B. (2003). Pathogenic human viruses in coastal waters. *Clinical Microbiology Reviews*, *16*(1), 129–143. https://doi.org/10.1128/CMR.16.1.129-143.2003
- Haile, R. W., Witte, J. S., Gold, M., Cressey, R., McGee, C., Millikan, R. C., Glasser, A., Harawa, N., Ervin, C., Harmon, P., Harper, J., Dermand, J., Alamillo, J., Barrett, K., Nides, M., & Wang, G. Y. (1999). The health effects of swimming in ocean water contaminated by storm drain runoff. *Epidemiology*, *10*(4), 355–363. https://doi.org/10.1097/00001648- 199907000-00004
- Hamilton, K. A., Ahmed, W., Palmer, A., Smith, K., Toze, S., & Haas, C. N. (2017). Seasonal Assessment of Opportunistic Premise Plumbing Pathogens in Roof-Harvested Rainwater Tanks. *Environmental Science and Technology*, *51*(3), 1742–1753. https://doi.org/10.1021/acs.est.6b04814
- Hart, J. D., Blackwood, A. D., & Noble, R. T. (2020). Examining coastal dynamics and recreational water quality by quantifying multiple sewage specific markers in a North Carolina estuary. *Science of the Total Environment*, *747*, 141124. https://doi.org/10.1016/j.scitotenv.2020.141124
- Harwood, V. J., Staley, C., Badgley, B. D., Borges, K., & Korajkic, A. (2014). Microbial source tracking markers for detection of fecal contamination in environmental waters:
- Relationships between pathogens and human health outcomes. In *FEMS Microbiology Reviews 38*(1), 1–40. https://doi.org/10.1111/1574-6976.12031
- Hathaway, J. M., & Hunt, W. F. (2011). Evaluation of first flush for indicator bacteria and total suspended solids in urban stormwater runoff. *Water, Air, and Soil Pollution*, *217*(1–4), 135– 147. https://doi.org/10.1007/s11270-010-0574-y
- Haugland, R. A., Siefring, S. C., Wymer, L. J., Brenner, K. P., & Dufour, A. P. (2005). Comparison of Enterococcus measurements in freshwater at two recreational beaches by quantitative polymerase chain reaction and membrane filter culture analysis. *Water Research*, *39*(4), 559–568. https://doi.org/10.1016/j.watres.2004.11.011
- Haugland, R. A., Varma, M., Sivaganesan, M., Kelty, C., Peed, L., & Shanks, O. C. (2010). Evaluation of genetic markers from the 16S rRNA gene V2 region for use in quantitative detection of selected Bacteroidales species and human fecal waste by qPCR. *Systematic and Applied Microbiology*, *33*(6), 348–357. https://doi.org/10.1016/J.SYAPM.2010.06.001
- Hino, M., Belanger, S. T., Field, C. B., Davies, A. R., & Mach, K. J. (2019). High-tide flooding disrupts local economic activity. *Science Advances*, *5*(2), eaau2736. https://doi.org/10.1126/sciadv.aau2736
- Jeong, Y., Grant, S. B., Ritter, S., Pednekar, A., Candelaria, L., & Winant, C. (2005). Identifying pollutant sources in tidally mixed systems: Case study of fecal indicator bacteria from marinas in Newport Bay, Southern California. *Environmental Science and Technology*, *39*(23), 9083–9093. https://doi.org/10.1021/es0482684
- Jin, G., Englande, A. J., Bradford, H., & Jeng, H.-W. (2004). Comparison of *E. coli*, Enterococci, and Fecal Coliform as Indicators for Brackish Water Quality Assessment. *Water Environment Research*, *76*(3), 245–255. https://doi.org/10.2175/106143004X141807
- Jothikumar, N., Cromeans, T. L., Hill, V. R., Lu, X., Sobsey, M. D., & Erdman, D. D. (2005). Quantitative real-time PCR assays for detection of human adenoviruses and identification of serotypes 40 and 41. *Applied and Environmental Microbiology*, *71*(6), 3131–3136. https://doi.org/10.1128/AEM.71.6.3131-3136.2005
- Jovanovic, D., Coleman, R., Deletic, A., & McCarthy, D. T. (2017). Tidal fluctuations influence *E. coli* concentrations in urban estuaries. *Marine Pollution Bulletin*, *119*(1), 226–230. https://doi.org/10.1016/j.marpolbul.2017.04.004
- Kemp, A. C., Bernhardt, C. E., Horton, B. P., Kopp, R. E., Vane, C. H., Peltier, W. R., Hawkes, A. D., Donnelly, J. P., Parnell, A. C., & Cahill, N. (2014). Late Holocene sea- and land-level change on the U.S. southeastern Atlantic coast. *Marine Geology*, *357*, 90–100. https://doi.org/10.1016/j.margeo.2014.07.010
- Kildare, B. J., Leutenegger, C. M., McSwain, B. S., Bambic, D. G., Rajal, V. B., & Wuertz, S. (2007). 16S rRNA-based assays for quantitative detection of universal, human-, cow-, and dog-specific fecal Bacteroidales: A Bayesian approach. *Water Research*, *41*(16), 3701– 3715. https://doi.org/10.1016/j.watres.2007.06.037
- King Tides Project (2020). *North Carolina Tides*. Retrieved from http://nckingtides.web.unc.edu/astronomical-tides/
- Kirchman, D., Peterson, B., & Juers, D. (1984). Bacterial growth and tidal variation in bacterial abundance in the Great Sippewissett Salt Marsh. *Marine Ecology Progress Series*, 19, 247- 259. https://doi.org/10.3354/meps019247
- Kopp, R. E., Horton, B. P., Kemp, A. C., & Tebaldi, C. (2015). Past and future sea-level rise along the coast of North Carolina, USA. *Climatic Change*, *132*(4), 693–707. https://doi.org/10.1007/s10584-015-1451-x
- Krometis, L. A. H., Characklis, G. W., Simmons, O. D., Dilts, M. J., Likirdopulos, C. A., & Sobsey, M. D. (2007). Intra-storm variability in microbial partitioning and microbial
- loading rates. *Water Research*, *41*(2), 506–516. https://doi.org/10.1016/J.WATRES.2006.09.029
- Lambertini, E., Spencer, S. K., Bertz, P. D., Loge, F. J., Kieke, B. A., & Borchardt, M. A. (2008). Concentration of enteroviruses, adenoviruses, and noroviruses from drinking water by use of glass wool filters. *Applied and Environmental Microbiology*, *74*(10), 2990–2996. https://doi.org/10.1128/AEM.02246-07
- Lee, S., Suits, M., Wituszynski, D., Winston, R., Martin, J., & Lee, J. (2020). Residential urban stormwater runoff: A comprehensive profile of microbiome and antibiotic resistance. *Science of the Total Environment*, *723*, 138033. https://doi.org/10.1016/j.scitotenv.2020.138033
- Leecaster, M. K., & Weisberg, S. B. (2001). Effect of sampling frequency on shoreline microbiology assessments. *Marine Pollution Bulletin*, *42*(11), 1150–1154. https://doi.org/10.1016/S0025-326X(01)00130-8
- Lewis, D. J., Atwill, E. R., Pereira, M. das G. C., & Bond, R. (2013). Spatial and Temporal Dynamics of Fecal Coliform and Escherichia coli Associated with Suspended Solids and Water within Five Northern California Estuaries. *Journal of Environmental Quality*, *42*(1), 229–238. https://doi.org/10.2134/jeq2011.0479
- Lipp, E. K., Kurz, R., Vincent, R., Rodriguez-Palacios, C., Farrah, S. R., & Rose, J. B. (2001). The effects of seasonal variability and weather on microbial fecal pollution and enteric pathogens in a subtropical estuary. *Estuaries*, *24*(2), 266–276. https://doi.org/10.2307/1352950
- Mallin, M. A., Esham, E. C., Williams, K. E., & Nearhoof, J. E. (1999). Tidal stage variability of fecal coliform and chlorophyll-a concentrations in coastal creeks. In *Marine Pollution Bulletin*, *38*(5). https://doi.org/10.1016/S0025-326X(99)00024-7
- Mallin, M. A., Williams, K. E., Esham, E. C., & Lowe, R. P. (2000). Effect of human development on bacteriological water quality in coastal watersheds. *Ecological Applications*, *10*(4), 1047–1056. https://doi.org/10.1890/1051- 0761(2000)010[1047:EOHDOB]2.0.CO;2

Mallin, M. A., Johnson, V. L., & Ensign, S. H. (2009). Comparative impacts of stormwater runoff on water quality of an urban, a suburban, and a rural stream. *Environmental Monitoring and Assessment*, *159*(1–4), 475–491. https://doi.org/10.1007/s10661-008-0644- 

Mill, A., Schlacher, T., & Katouli, M. (2006). Tidal and longitudinal variation of faecal indicator bacteria in an estuarine creek in south-east Queensland, Australia. *Marine Pollution Bulletin*, *52*(8), 881–891. https://doi.org/10.1016/j.marpolbul.2005.11.018



Parker, J. K., McIntyre, D., & Noble, R. T. (2010). Characterizing fecal contamination in stormwater runoff in coastal North Carolina, USA. *Water Research*, *44*(14), 4186–4194. https://doi.org/10.1016/j.watres.2010.05.018 Partyka, M. L., Bond, R. F., Chase, J. A., & Atwill, E. R. (2017). Monitoring bacterial indicators of water quality in a tidally influenced delta: A Sisyphean pursuit. *Science of the Total Environment*, *578*, 346–356. https://doi.org/10.1016/j.scitotenv.2016.10.179 Paule-Mercado, M. A., Ventura, J. S., Memon, S. A., Jahng, D., Kang, J. H., & Lee, C. H. (2016). Monitoring and predicting the fecal indicator bacteria concentrations from agricultural, mixed land use and urban stormwater runoff. *Science of the Total Environment*, *550*, 1171–1181. https://doi.org/10.1016/J.SCITOTENV.2016.01.026 Pednekar, A.M., S.B. Grant, Y. Jeong, Y. Poon, and C. Oancea (2005). Influence of climate change, tidal mixing, and watershed urbanization on historical water quality in Newport Bay, a saltwater wetland and tidal embayment in Southern California. *Environmental Science and Technology, 39*(23):9071-9082. https://doi.org/10.1021/es0504789 Poulter, B., Feldman, R. L., Brinson, M. M., Horton, B. P., Orbach, M. K., Pearsall, S. H., Reyes, E., Riggs, S. R., & Whitehead, J. C. (2009). Sea-level rise research and dialogue in North Carolina: Creating windows for policy change. *Ocean and Coastal Management*, *52*(3–4), 147–153. https://doi.org/10.1016/j.ocecoaman.2008.09.010 Pricope, N. G., Halls, J. N., & Rosul, L. M. (2019). Modeling residential coastal flood vulnerability using finished-floor elevations and socio-economic characteristics. *Journal of Environmental Management*, *237*, 387–398. https://doi.org/10.1016/j.jenvman.2019.02.078 Prüss, A. (1998). Review of epidemiological studies on health effects from exposure to recreational water. In *International Journal of Epidemiology*, *27*(1). https://doi.org/10.1093/ije/27.1.1 827 Rajal, V. B., McSwain, B. S., Thompson, D. E., Leutenegger, C. M., & Wuertz, S. (2007). Molecular quantitative analysis of human viruses in California stormwater. *Water Research*, *41*(19), 4287–4298. https://doi.org/10.1016/j.watres.2007.06.002 Rippy, M. A., Stein, R., Sanders, B. F., Davis, K., McLaughlin, K., Skinner, J. F., Kappeler, J., & Grant, S. B. (2014). Small drains, big problems: The impact of dry weather runoff on shoreline water quality at enclosed beaches. *Environmental Science and Technology*, *48*(24), 14168–14177. https://doi.org/10.1021/es503139h Seruge, J., Wong, M., Noble, R. T., Blackwood, A. D., Moravcik, P. S., & Kirs, M. (2019). Application of a rapid qPCR method for enterococci for beach water quality monitoring purposes in Hawaii: Loss of DNA during the extraction protocol due to coral sands. *Marine Pollution Bulletin*, *149*. https://doi.org/10.1016/j.marpolbul.2019.110631

- Shanks, O. C., Green, H., Korajkic, A., & Field, K. G. (2015). Overview of Microbial Source Tracking Methods Targeting Human Fecal Pollution Sources. In *Manual of Environmental Microbiology* (3.4.3-1-3.4.3-8). https://doi.org/10.1128/9781555818821.ch3.4.3
- Shehane, S. D., Harwood, V. J., Whitlock, J. E., & Rose, J. B. (2005). The influence of rainfall on the incidence of microbial faecal indicators and the dominant sources of faecal pollution in a Florida river. *Journal of Applied Microbiology*, *98*(5), 1127–1136. https://doi.org/10.1111/j.1365-2672.2005.02554.x
- Sidhu, J. P. S., Hodgers, L., Ahmed, W., Chong, M. N., & Toze, S. (2012). Prevalence of human pathogens and indicators in stormwater runoff in Brisbane, Australia. *Water Research*, *46*(20), 6652–6660. https://doi.org/10.1016/j.watres.2012.03.012
- Silva, M. R., Bravo, H. R., Cherkauer, D., Klump, J. V., Kean, W., & McLellan, S. L. (2014). Effect of hydrological and geophysical factors on formation of standing water and FIB reservoirs at a Lake Michigan beach*. Journal of Great Lakes Research, 40*(3), 778-789. https://doi.org/10.1016/j.jglr.2014.06.003
- Sinclair, R. G., Jones, E. L., & Gerba, C. P. (2009). Viruses in recreational water-borne disease outbreaks: A review. *Journal of Applied Microbiology*, *107*(6), 1769–1780. https://doi.org/10.1111/j.1365-2672.2009.04367.x
- Sinton, L. W., Hall, C. H., Lynch, P. A., & Davies-Colley, R. J. (2002). Sunlight inactivation of fecal indicator bacteria and bacteriophages from waste stabilization pond effluent in fresh and saline waters. *Applied and Environmental Microbiology*, *68*(3), 1122–1131. https://doi.org/10.1128/AEM.68.3.1122-1131.2002
- Soller, J. A., Schoen, M. E., Bartrand, T., Ravenscroft, J. E., & Ashbolt, N. J. (2010). Estimated human health risks from exposure to recreational waters impacted by human and non-human sources of faecal contamination. *Water Research*, *44*(16), 4674–4691. https://doi.org/10.1016/j.watres.2010.06.049
- Soller, J. A., Schoen, M. E., Varghese, A., Ichida, A. M., Boehm, A. B., Eftim, S., Ashbolt, N. J., & Ravenscroft, J. E. (2014). Human health risk implications of multiple sources of faecal indicator bacteria in a recreational waterbody. *Water Research*, *66*, 254–264. https://doi.org/10.1016/j.watres.2014.08.026
- Soller, J., Bartrand, T., Ravenscroft, J., Molina, M., Whelan, G., Schoen, M., & Ashbolt, N. (2015). Estimated human health risks from recreational exposures to stormwater runoff containing animal faecal material. *Environmental Modelling and Software*, *72*, 21–32. https://doi.org/10.1016/J.ENVSOFT.2015.05.018
- Soller, J. A., Schoen, M., Steele, J. A., Griffith, J. F., & Schiff, K. C. (2017). Incidence of gastrointestinal illness following wet weather recreational exposures: Harmonization of quantitative microbial risk assessment with an epidemiologic investigation of surfers. *Water Research*, *121*, 280–289. https://doi.org/10.1016/J.WATRES.2017.05.017
- Steele, J. A., Blackwood, A. D., Griffith, J. F., Noble, R. T., & Schiff, K. C. (2018). Quantification of pathogens and markers of fecal contamination during storm events along popular surfing beaches in San Diego, California. *Water Research*, *136*, 137–149. https://doi.org/10.1016/J.WATRES.2018.01.056 Stumpf, C. H., Piehler, M. F., Thompson, S., & Noble, R. T. (2010). Loading of fecal indicator bacteria in North Carolina tidal creek headwaters: Hydrographic patterns and terrestrial runoff relationships. *Water Research*, *44*(16), 4704–4715. https://doi.org/10.1016/j.watres.2010.07.004 Sweet, W., Park, J., Marra, J., Zervas, C., & Gill, S. (2014). Sea Level Rise and Nuisance Flood Frequency Changes around the United States. *NOAA Technical Report NOS CO-OPS 073* (Issue June). Retrieved from http://tidesandcurrents.noaa.gov/publications/NOAA\_Technical\_Report\_NOS\_COOPS\_07 3.pdf Tiefenthaler, L., Stein, E. D., & Schiff, K. C. (2011). Levels and patterns of fecal indicator bacteria in stormwater runoff from homogenous land use sites and urban watersheds. *Journal of Water and Health*, *9*(2), 279–290. https://doi.org/10.2166/wh.2010.056 Titus, J. G., & Richman, C. (2001). Maps of lands vulnerable to sea level rise: Modeled elevations along the US Atlantic and Gulf coasts. *Climate Research*, *18*(3), 205–228. https://doi.org/10.3354/cr018205 Tolouei, S., Dewey, R., Snodgrass, W. J., Edge, T. A., Andrews, R. C., Taghipour, M., Prévost, M., & Dorner, S. (2019). Assessing microbial risk through event-based pathogen loading and hydrodynamic modelling. *Science of the Total Environment*, *693*. https://doi.org/10.1016/j.scitotenv.2019.07.373
- Tu, J., & Xia, Z. G. (2008). Examining spatially varying relationships between land use and water quality using geographically weighted regression I: Model design and evaluation. *Science of the Total Environment*, *407*(1), 358–378. https://doi.org/10.1016/j.scitotenv.2008.09.031
- US Census Bureau. (2020). Population and Housing, Population and Housing Unit Counts. *U.S. Department of Commerce*, *July*, 1–141. Retrieved from http://www.census.gov/prod/cen2010/cph-2-53.pdf
- US Climate (2020). *Climate Wilmington North Carolina and Weather averages Wilmington*. Retrieved from https://www.usclimatedata.com/climate/wilmington/north-carolina/united-states/usnc0760/2018/12
- US EPA (1986). Ambient Water Quality Criteria for Bacteria 1986. *U. S. Environmental Protection Agency*, *Washington, DC*. Retrieved from https://www.epa.gov/sites/production/files/2019-03/documents/ambient-wqc-bacteria-
- 1986.pdf
- US EPA (2012). Recreational Water Quality Criteria. *U. S. Environmental Protection Agency*, *Washington, DC*. Retrieved from https://www.epa.gov/sites/production/files/2015- 10/documents/rwqc2012.pdf
- US EPA Method 1609 (2012). Enterococci in Water by TaqMan® Quantitative Polymerase Chain Reaction (qPCR) with Internal Amplification Control (IAC) Assay. Retrieved from https://www.epa.gov/sites/production/files/2015-08/documents/method\_1609\_2013.pdf
- US EPA Method 1611 (2012). Enterococci in Water by TaqMan ® Quantitative Polymerase Chain Reaction (qPCR) Assay. Washington DC. Retrieved from https://www.epa.gov/sites/production/files/2015-08/documents/method\_1611\_2012.pdf
- USGS. (2020). Runoff: Surface and Overland water runoff. *USGS Official Website*. Retrieved from https://www.usgs.gov/special-topic/water-science-school/science/runoff-surface-and-923 overland-water-runoff?qt-science center objects=0#qt-science center objects
- Van De Plassche, O., Wright, A. J., Horton, B. P., Engelhart, S. E., Kemp, A. C., Mallinson, D., & Kopp, R. E. (2014). Estimating tectonic uplift of the Cape Fear Arch (south-eastern United States) using reconstructions of Holocene relative sea level. *Journal of Quaternary Science*, *29*(8), 749–759. https://doi.org/10.1002/jqs.2746
- Visit North Carolina, (2019). 2018 North Carolina Regional Visitor Profile. *Economic Development Partnership of North Carolina, Cary, NC*. Retrieved from https://partners.visitnc.com/contents/sdownload/71007/file/2018-North-Carolina-Regional-Visitor-Profile.pdf
- Wade, T. J., Calderon, R. L., Sams, E., Beach, M., Brenner, K. P., Williams, A. H., & Dufour, A. P. (2006). Rapidly measured indicators of recreational water quality are predictive of swimming-associated gastrointestinal illness. *Environmental Health Perspectives*, *114*(1), 24–28. https://doi.org/10.1289/ehp.8273
- Wade, T. J., Calderon, R. L., Brenner, K. P., Sams, E., Beach, M., Haugland, R., Wymer, L., & Dufour, A. P. (2008). High sensitivity of children to swimming-associated gastrointestinal illness: Results using a rapid assay of recreational water quality. *Epidemiology*, *19*(3), 375– 383. https://doi.org/10.1097/EDE.0b013e318169cc87
- Wilhelm, S. W., Brigden, S. M., & Suttle, C. A. (2002). A dilution technique for the direct measurement of viral production: A comparison in stratified and tidally mixed coastal waters. *Microbial Ecology*, *43*(1), 168–173. https://doi.org/10.1007/s00248-001-1021-9
- Yau, V. M., Schiff, K. C., Arnold, B. F., Griffith, J. F., Gruber, J. S., Wright, C. C., & Gold, M. (2014). Effect of submarine groundwater discharge on bacterial indicators and swimmer health at Avalon Beach, CA, USA. *Water Research*, *59*, 23-36.
- https://doi.org/10.1016/j.watres.2014.03.050

Zhu, X., Wang, J. D., Solo-Gabriele, H. M., & Fleming, L. E. (2011). A water quality modeling study of non-point sources at recreational marine beaches. *Water Research*, *45*(9), 2985– 2995. https://doi.org/10.1016/j.watres.2011.03.015

