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[Simon Jackson](#)*, Christian Good, Alistair White, [Christopher Seymour](#), [Joao Brandao](#)

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Article

Water Quality Assessment: Endotoxin Brings Real-time Measurements and Non-Faecally Transmitted Bacteria to the Table

Christian Good ¹, Alistair White ¹, Joao Brandao ^{2,3}, Christopher Seymour ⁴
and Simon K. Jackson ^{1,5}

¹ Molendotech Limited, Brixham Laboratory, Blackball Lane, Freshwater Quarry, Brixham TQ5 8BA, UK

² National Institute of Health Dr. Ricardo Jorge, Department of Environmental Health, Av. Padre Cruz 1649-016 Lisboa, Portugal

³ cE3c – Centre for Ecology, Evolution and Environmental Changes, Faculdade de Ciências da Universidade de Lisboa, Campo Grande 016, 1749-016 Lisboa, Portugal.

⁴ Anglian Water, Kingfisher Way, Hinchingsbrooke Business Park, Huntingdon, Cambridgeshire, PE29 6FL, UK

⁵ School of Biomedical Science, Faculty of Health, University of Plymouth, Drake Circus, Plymouth PL4 8AA, UK

* Correspondence: simon.jackson@molendotech.com

Abstract: We have used a rapid, portable assay (Bacterisk) to determine bacterial water quality along several inland waters in SW England. Water samples were compared by conventional membrane filter and culture methods for faecal indicator bacteria (FIB; *E. coli* and enterococci) and endotoxin measurement by Bacterisk. The Bacterisk data, measured in near real-time, correlates well with both *E. coli* and enterococci, but also allowed the presence of potential pathogens of non-faecal origin to be detected. The sensitivity was calculated to be 92.96% with a specificity of 46.3% for *E. coli* with an expanded uncertainty of 22.07% and an Endotoxin Risk detection limit of 25 units. The presence of Bacterisk detectable non-faecal pathogenic bacteria in the water samples was successfully confirmed by Illumina MiSeq sequencing followed by target species-specific qPCR. Sequencing showed the presence of pathogens including *Pseudomonas aeruginosa*, *Salmonella typhi*, *Acinetobacter baumannii*, *Shigella* spp, *Legionella* spp as well as antimicrobial resistance genes. Furthermore, the portable Bacterisk assay was able to acquire data on water quality from different locations and at different time points providing a comprehensive surveillance tool that challenges the time to results by conventional methods (minutes instead of days), yielding compatible results.

Keywords: water quality; endotoxin; faecal indicator bacteria; public health

Introduction

Faecal indicator bacteria have been used for one and a half centuries to indicate faecal contamination of water and associated health risks (Holcomb & Stewart, 2020). While useful as indicators of potential faecally transmitted pathogens, reliance on *E. coli* and Enterococci as proxies can miss out the presence of other pathogens and importantly for pathogens of non-faecal origin. The WHO guidelines (WHO 2021) recommend a risk-based approach to consider pathogens that are not necessarily and almost exclusively associated with faecal contamination. Non-faecally transmitted bacteria in recreational waters pose significant health risks to users, underscoring the limitations of relying solely on faecal indicator bacteria (FIB) to assess water quality. These bacteria, which originate from sources other than faecal contamination, can include environmental pathogens such as *Vibrio* spp., *Pseudomonas aeruginosa*, and *Legionella* spp., as well as opportunistic bacteria that thrive in aquatic ecosystems (Boehm & Soller. 2013, Fewtrell & Kay. 2015, Rodrigues et al. 2017). In fact, a recent report

from the Centers for Disease Control and Prevention (CDC) (Hlavsa et al. 2021) shows that these organisms were the main waterborne illness provoking agents in treated recreational water between 2015-2019. All these pathogens can enter recreational waters through various sources and pathways. For instance, *Vibrio* species thrive in warm, brackish, or marine environments, with their populations often peaking during the summer months (Sampaio et al., 2022). Climate change with increased water temperatures is also an emerging factor for increases in *Vibrio* spp. abundance. Similarly, bacteria such as *Pseudomonas aeruginosa* can persist in sediments or biofilms, where they are shielded from environmental stressors (Brandão et al. 2022). Human activity also plays a role; the introduction of skin flora and the use of improperly treated recreational water features, such as pools or water parks, can amplify the presence of these bacteria. (Ayi B. 2015)

Exposure to non-fecally transmitted bacteria can lead to a range of health problems for recreational water users. These bacteria are often associated with skin and soft tissue infections, such as cellulitis, dermatitis, or, in severe cases, necrotising infections caused by pathogens like *Vibrio vulnificus* or *Pseudomonas aeruginosa*. (Guida et al., 2016). Contact with contaminated water can also result in ear, eye, and respiratory issues, including swimmer's ear (*otitis externa*) (Pantazidou et al. 2022) and respiratory illnesses linked to *Legionella* exposure (National Academy of Sciences et al. 2019). For immunocompromised individuals, these pathogens may lead to systemic infections, including sepsis, particularly if the skin barrier is compromised through cuts or abrasions.

The presence of these pathogens presents a challenge for water quality monitoring, which relies on FIB such as *E. coli* or enterococci. (WHO, 2021; European Parliament and Council, 2006). As a result, a water body considered safe based on FIB levels may still harbour significant health risks from environmental bacteria. Additionally, the standard culture methods for determining FIB for water quality assessment take over 24h for results, meaning water quality is only known after exposure (ISO 7899-1 or ISO 7899-2 for enterococci and ISO 9308-3 or ISO 9308-1 for *E. coli*). Moreover, the last revision of these standards recommends their use in either very clean or very dirty waters for enterococci. In addition, sampling is not performed continuously so discrete points are all that regulators can use (Wade et al., 2010). Moreover, by focussing solely on FIB it is possible that other pathogens that are not necessarily of faecal nature, may go undetected (Topić et al. 2021). These drawbacks to the current culture-based approaches highlight the need for not only rapid, but also more complete solutions with the ability to monitor water quality at many locations and at different times and to better indicate a broader set of pathogens.

A rapid and simple test of bacterial water quality would therefore be a very useful tool. Previous studies have suggested that measuring endotoxin (lipopolysaccharide (LPS)), present in the outer membrane of Gram-negative bacteria and some cyanobacteria, may be a useful technique for rapidly determining bacterial biomass and quality of water (Evans et al., 1978; Jorgensen et al., 1979; Haas et al., 1983). Previous work by our group has shown the applicability of using endotoxin as a marker of faecal contamination of seawater (Sattar et al., 2014, Good et al., 2024) and that measuring endotoxin correlates with inflammatory effects of contaminated water samples (Sattar et al., 2019). Researchers at Molendotech have developed a near real-time assay (Bacterisk®) to assess bacterial water quality based on endotoxin, which can be conducted by non-specialist staff *in situ*. The advantage of measuring endotoxin as an indicator for contaminated water is that the test is specific for LPS, a compound which only naturally occurs in the cell walls of Gram-negative bacteria. LPS comprises a relatively constant proportion of a Gram-negative bacterial cell and Gram-negative bacteria account for 80 to 95% of the prokaryotes found in waters (Anwar and Choi, 2014). Moreover, endotoxin could indicate the presence of Gram-negative pathogens not detected by current culture of total coliforms or *E. coli*. Therefore, this novel assay would allow near real-time assessment of water quality and the flexibility to sample at several locations and at different time points. It would thus provide a more complete assessment of water quality and also allow rapid testing in remote and disaster areas where access to laboratories and water testing facilities is challenging.

The presence of AntiMicrobial Resistance (AMR) genes is also not covered by FIB monitoring. Recreational waters, including coastal bathing areas, urban lakes, and estuarine environments, act as

reservoirs and transmission pathways for antimicrobial resistance genes (ARGs). Leonard et al. (2015) emphasise the risk of human exposure to antibiotic-resistant bacteria during recreational activities in coastal waters, where the dissemination of ARGs is influenced by anthropogenic pollution. In this perspective, Farrell et al. (2023) link recreational water use to an increased carriage of antimicrobial-resistant organisms and Singh et al. (2022) describe how wastewater and natural water systems act as vectors for the spread of ARGs into recreational waters. Variations in ARG profiles across different recreational water sources, shaped by local microbial communities and water management practices have been demonstrated by Han et al. (2022) and Li et al. (2022). These findings underscore the critical need to monitor and mitigate ARG contamination to protect both environmental and public health.

Previous reports have highlighted the usefulness of the rapid Bacterisk technology in determining coastal water quality. The present study was undertaken to assess the use of Bacterisk as a rapid method to determine bacterial contamination in fresh (inland) waters and to provide evidence for the detection on non-faecal pathogens by this method. This study therefore challenges the sole use of FIB detection for the assessment of water quality.

Materials and Methods

Water Sampling

A total of 36 inland water samples were collected from various river locations in the southwest of England. Briefly, 500 mL samples were taken using sterile bottles 30 cm below the water's surface in water at least one meter deep. The samples were then transported in the dark and tested within 4 hours or stored in a fridge (2-8 °C) and tested no later than 24 hours after collection.

Bacterial Culture Identification (ISO Methods)

Appropriate volumes of each water sample (1, 10, and 100 mL) were aseptically filtered through a 0.45 µm membrane (Whatman, UK) using a 6-branch vacuum manifold (Sartorius, UK). Following ISO 9308-1:2014 and ISO 7899-2:2000, Membranes were placed on membrane TBX agar (Oxoid, UK) and incubated at 30°C for 4 hours, then at 37°C for 14 hours for the detection of presumptive *E. coli* or on Slanetz and Bartley medium (Oxoid, UK) and incubated at 36°C for 44 hours for the detection of presumptive intestinal *enterococci*. The numbers of colony-forming units (CFU) were then calculated and expressed as CFU/100mL.

Bacterisk Assay

The Bacterisk assay (Molendotech Ltd., Brixham, UK; www.molendotech.com) was performed following the manufacturer's instructions. Briefly, the samples were diluted 1 in 40 in dilution buffer and then 200µL was transferred to a reaction tube containing the lyophilised detection reagent. The samples were then incubated at 37°C for 13 minutes using the integrated Bacterisk incubator and reader. An Endotoxin Risk Unit (ERU) score was then calculated by the device based on the absorbance of the sample at 405 nm. Both a negative control (endotoxin-free water) and a positive control (Endotoxin from *E. coli* 055:B5) was run with every assay.

Sequencing Analysis

Bacterial gene sequencing was performed by Eurofins using the INVIEW Microbiome Profiling package with amplification and Illumina MiSeq sequencing of the hypervariable regions in the 16S rRNA gene. This method amplifies and sequences three targets from all DNA samples (16S V1-V3, 16S V3-V4 or 16S V3-V5).

qPCR

Species-specific quantitative real-time PCR and subsequent amplicon detections were performed on inland water samples from the river Dart by Friends of the Dart and Surfers Against Sewage UK.

Determination of Uncertainty of Measurement (UoM)

Data on samples using the Bacterisk methodology were analysed in duplicate (A and B). Analysis for Measurement of Uncertainty was performed following the guidelines for expanded uncertainty (Magnusson et al., 2017), often referred to as the square root of the sum of the squares multiplied by a coverage factor (*k*) to the desired confidence.

Sum ((log₁₀B-log₁₀A)²) n = total variance (T)

Standard Deviation (SD) = $\sqrt{T/n}$

To calculate the expanded uncertainty, the relative SD is calculated by dividing the standard deviation by the mean of the Log10 observed values, and represented as a percentage. To calculate the expanded uncertainty this relative standard deviation is multiplied by a coverage factor (*k*) at the required confidence limit. Various values have been suggested for confidence limits both fixed and variable. A standard approach is to use a fixed *k* value of 2 for a 95% confidence.

Statistical Analysis

Statistical analyses on the distribution of samples and the correlation between Bacterisk data and culture-derived water quality were performed with Fisher’s exact test and Chi-square test using GraphPad prism version 9 for Windows, GraphPad Software, Boston, Massachusetts USA, www.graphpad.com. Receiver Operating Characteristics (ROC) curve, was used to determine the optimal threshold ERU value used to discriminate the water quality groups. The ROC curve uses 1 – specificity on the x-axis, as calculated:

$$Specificity = \frac{True\ negatives}{True\ negatives + False\ positives}$$

and sensitivity (true positive) on the y-axis, as calculated:

$$Sensitivity = \frac{True\ positives}{True\ positives + False\ negatives}$$

The ROC curve also provides an area under the curve (AUC) value between 0 and 1. The closer the AUC value is to 1 the better the model is at predicting a correct classification, whereas a value of 0.5 represents a model with no ability to predict a correct classification. A model with an AUC of greater than 0.8 is considered acceptable (Nahm 2022).

Results

Use of Expanded Uncertainty was used to provide a limit of quantification for the Bacterisk method, i.e. a value where we could provide confidence in the observed result. To aid this, the observed values were plotted as pairs in a low-high format. This data appeared to demonstrate that values begin to show greater significance between 20 and 30 Endotoxin Risk Units (ERU). The expanded uncertainty was calculated for values of 20 ERU, 25ERU, and 30ERU; to be included in assessment only one of the pairs of results required to satisfy this limit. Results obtained at the 3 different *k* values are shown in Table 1.

Table 1. Expanded Uncertainty of Measurements at 20ERU, 25ERU, and 30ERU.

ERU limit	Degrees of Freedom (<i>df</i>)	Fixed <i>k</i> =2	Fixed <i>k</i> =1.96	Varying <i>k</i> based on <i>df</i>	Expanded UoM
20	287	31.94%	31.30%	1.968264	31.43%
25	249	22.41%	21.97%	1.969537	22.07%

30	199	15.87%	15.56%	1.971957	15.65%
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From the results obtained in Table 1, the Expanded Uncertainty of Measurement decreases as the lower limit of inclusion increases. Comparing this to results obtained from culture-based microbiology methods (in house testing) values range from 12% to around 25%. Therefore, there is a limit of accurate quantification of 25 ERU.

A total of 36 inland water samples were analysed in parallel by the Bacterisk assay to calculate Endotoxin Risk Units (ERU) and by membrane filtration to enumerate the levels of *E. coli* and intestinal *enterococci* (CFU/100 mL), according to ISO 9308-1:2014 and ISO 7899-2:2000, respectively. The results were compared with data obtained for coastal water samples collected and analysed by the same methods as published previously (Good et al., 2024).

Figure 1 shows the water quality of inland water samples determined by Bacterisk and compared with culture of *E. coli* and enterococci by membrane filtration method. As can be seen, many of the samples of inland waters are seen to be of poor quality as assessed by the EU bathing water directive. All the data are presented in Table S1 in supplementary data.

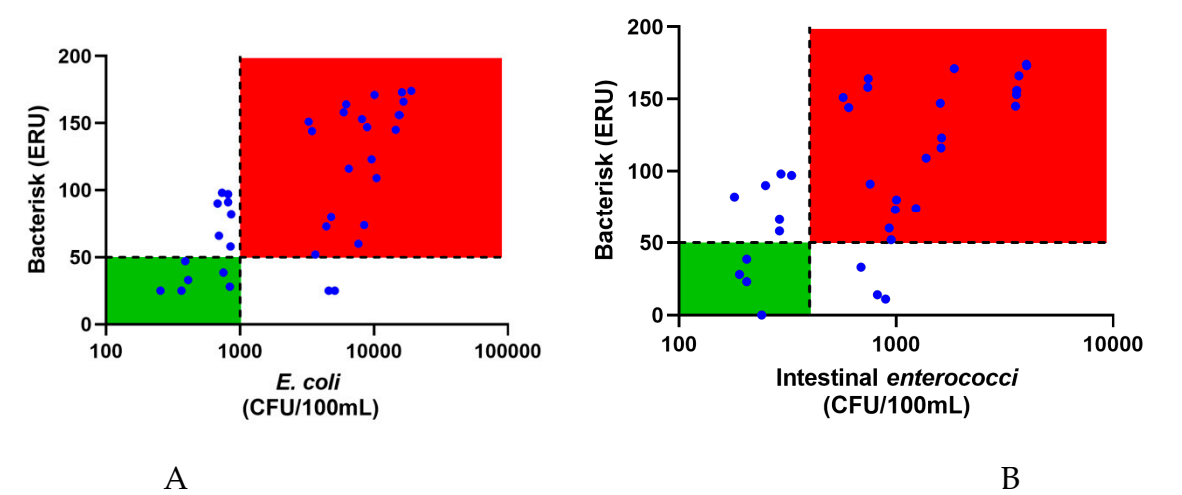


Figure 1. Scatter plot displaying the Bacterisk Endotoxin Risk Units against (A) *E. coli* and (B) Enterococci enumerated using the membrane filtration method on TBX agar for inland water samples. The chart is split into quadrants based on the ERU threshold (Y axis 50) and (A) *E. coli* cut-off (X axis 1000CFU/100mL), (B) Enterococci cut-off (X axis 400CFU/100mL). n=36. TN: true negative, TP: true positive, FN: false negative, OP: off-target positive.

The ROC analysis of the data from Figure 1A gave an area under the curve of 0.826 (p=0.0013) and sensitivity of 91.3% and specificity of 46.2%. The low specificity is due to the Bacterisk assay detecting all Gram-negative bacteria not just *E. coli* and hence alerting to the presence of potential pathogens.

Though intestinal *enterococci* are also used as a FIB it is a Gram-positive bacterium, and in fact the only current parameter recommended by the most recent WHO guidelines for recreational water quality; there was a correlation between intestinal *enterococci* and ERU. As can be seen from Figure 1, Bacterisk ERU values could track enterococci and ERU data were a good proxy for enterococci levels.

Due to Bacterisk determining the levels of a Gram-negative molecule (endotoxin) as a marker of water contamination, it not only provides good correlation with *E. coli* but will respond to other Gram-negative bacteria including non-FIB. These may be pathogenic and of concern for human health. These have been included in our data as ‘other’ or ‘off-target’ positives. It is important to understand what these other bacteria are and how they may contribute to the Bacterisk data. To accomplish this, 10 water samples from known ‘off-target’ positive results together with water samples giving low or high *E. coli* results, were DNA-sequenced to determine the bacterial flora composition. Representative results are shown in Figure 2. In addition, we obtained qPCR data from

samples from one of the river locations. The qPCR data (Figure 3) reveals the presence of several pathogenic species including *S. typhi*.

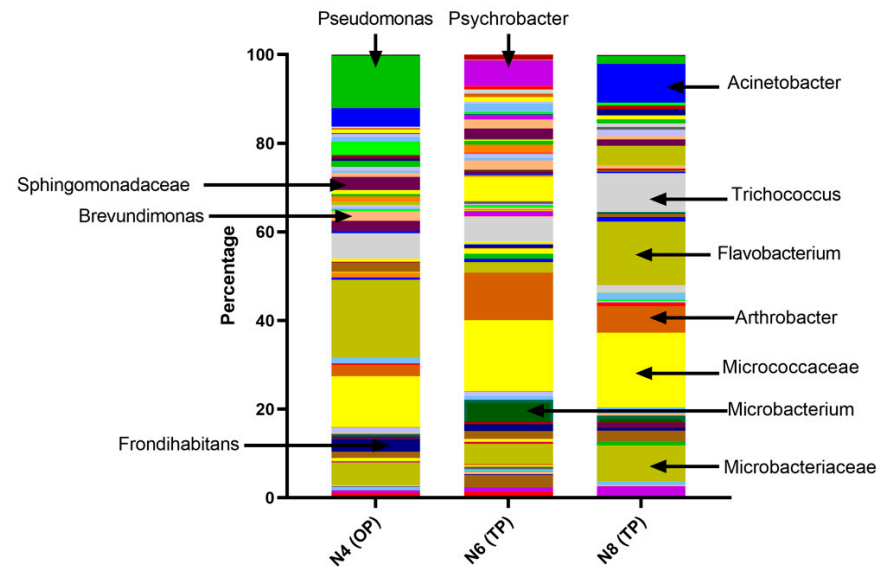


Figure 2. 16S RNA sequencing data from inland water samples showing the different genera present in typical samples. OP = off target positives; TP = True positives based on *E. coli* >1000CFU/100mL.

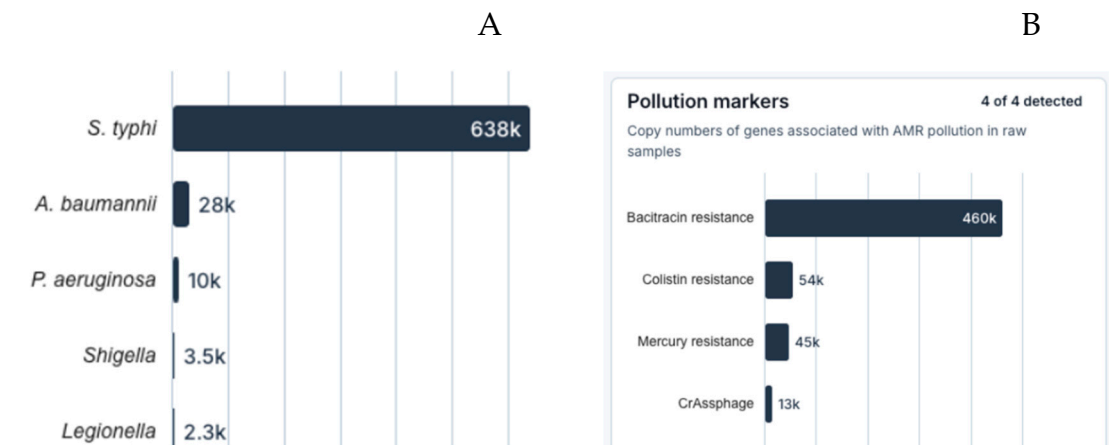


Figure 3. (A) Copy number of independently detected pathogen marker genes, in river water samples by qPCR. (B) AMR genes detected from these samples.

A disadvantage of current methods of water quality monitoring, in addition to the time to result delays, is the restriction to single sites and infrequent sampling. We took samples at different times from the river locations we had sampled for the data in Figures 1-3 to determine how water quality might vary with time and location. Results are shown in Figure 4. It should be noted that water quality assessed by Bacterisk correlates well with *E. coli* culture and that the water quality varies greatly on different dates of sampling highlighting the flux in water contamination within rivers.

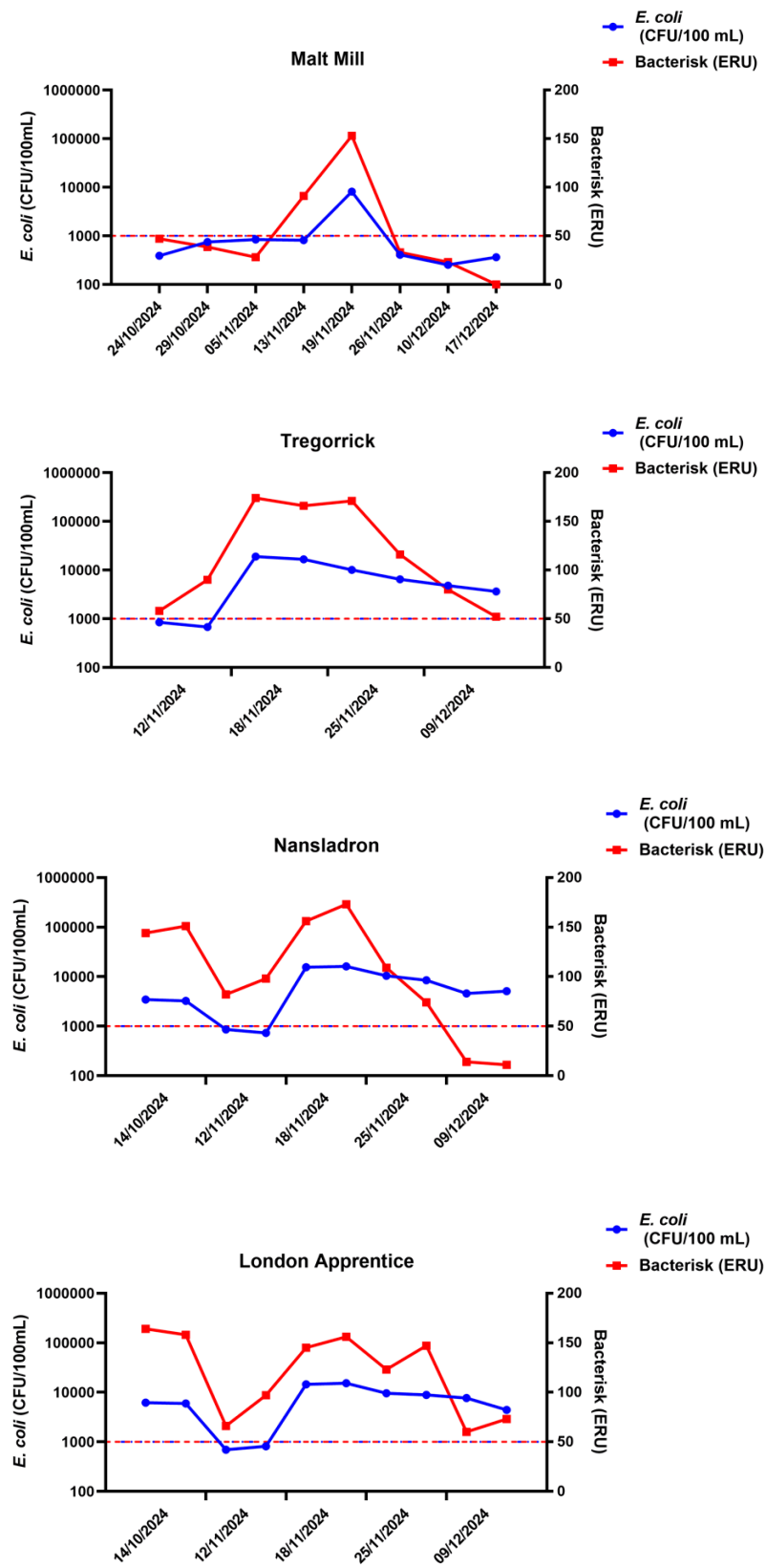


Figure 4. Water quality tracking using Bacterisk to measure ERU compared with *E. coli* along different rivers in SW England at different times.

Discussion

While FIB remain an essential tool for assessing faecal contamination, they are insufficient for evaluating the broader spectrum of waterborne health risks. A more holistic approach to monitoring and managing recreational water quality is needed to ensure the safety of all users and to address the emerging challenges posed by non-faecally transmitted pathogens. Also, the current ISO reference methods of enteric pathogenic bacteria detection are time-consuming, expensive, and often insensitive even in fresh faeces (Liu et al., 2014). Considering that many beach recreational and professional users may suffer from a certain degree of immunological compromise, it is crucial for beach managers to inform the public as thoroughly as possible of any risks associated with exposure to pathogens and opportunists at the beach (Stec et al. 2022). There is thus an urgent need for validated rapid methods to assess bacterial water quality that can be used *in situ* and cover a broader range of potential pathogens. This will enable proactive measures to be taken in the event of water contamination thereby protecting human health before use. Additionally, the recent water reuse policies regulated, for example, in the European Union by the recast of the Directive (EU) 2024/3019 of the European Parliament and of the Council of 27 November 2024 concerning urban wastewater treatment, aims at the protection of the environment from pathogens and is based on the same slow reference ISO methods. These methods show an incomplete picture and take up time, which sometimes does not exist, when facing an extreme weather event or an environmental disaster assessment (Halcomb et al. 2020).

The current study utilised a rapid bacterial assessment method (Bacterisk) and compared it with *E. coli* and enterococci culture to determine the bacterial water quality at different inland river and coastal sites in SW England. Bacterisk, which detects the endotoxin present in Gram-negative bacteria, has been used extensively and validated as a useful method to assess the bacterial contamination of coastal recreational waters (Sattar et al., 2022; Good et al., 2024). In the latter reference, we have shown that while Bacterisk assay results could be used to obtain risk groups that differentiate different levels of water quality, we have used it as a binary classification model to determine whether a water source is either polluted ('poor') or clean ('sufficient or better'), based on the regulatory levels of *E. coli* (EU bathing directive 2006) from thresholds of the ERU data from Bacterisk. Statistical analysis in the present study showed that the uncertainty of measurement for Bacterisk assay gave a 95% confidence in measurements above 25ERU and this level was set as the lower limit of detection for this method. From ROC analysis, compared with *E. coli* detection by membrane filtration, the assay provides a high sensitivity (92%) but a lower specificity (46%) – this is expected as the assay is not specific for *E. coli* but will also detect other Gram-negative bacteria including potential pathogens that are abundant in river water.

The present study has shown the applicability of this method for the analysis of recreational freshwaters as was shown previously for coastal waters (Good et al., 2024). In each case, the detection of endotoxin correlates well with conventional FIB *E. coli* and enterococci and can provide thresholds or cut-offs using the EU bathing water directive guidelines (EU, 2006). Bacterisk results, obtained in 15 minutes *in situ*, correlated well with conventional membrane filtration culture results. As Bacterisk is not restricted to detect only *E. coli* or enterococci, the present faecal indicator bacteria, it highlighted the presence of appreciable levels of other Gram-negative bacteria in river water. DNA sequence and qPCR analysis of the river water samples testing positive by Bacterisk endotoxin detection but low for *E. coli* by culture confirmed the presence of pathogenic bacteria including *Pseudomonas aeruginosa* and *Salmonella typhi* (Mena and Gerba, 2009). This shows the advantage of being able to detect bacteria other than the current FIB, as other pathogens might be present in water identified as 'good or sufficient' by current regulatory standards. Moreover, evidence is presented for the presence of antimicrobial resistance genes in these water samples, highlighting the need to be able to detect the presence of such bacteria.

In addition to the long time to results for bacterial culture, limitations of the faecal indicator paradigm have long been acknowledged (Field et al., 2007; Stewart et al., 2008). Researchers have identified many challenges and limitations to the effective use of both traditional and alternative faecal indicators to characterize risk, identify sources, and evaluate interventions (Stewart et al., 2013;

Fewtrell and Kay 2015). Arguably, one of the most significant limitations is the inconsistent relationships between FIB occurrence, enteric pathogens, and health risks (Fewtrell and Kay, 2015; Korajkic et al., 2018).

The FIB found to correlate with health risks vary widely by site (Griffith et al., 2016). Our data presented here (Figure 4) show that the levels of bacterial contamination vary greatly by site and by date. The co-occurrence of enteric pathogens and FIB in ambient waters is inconsistent at best (Korajkic et al., 2018; Wu et al., 2011) and commonly used FIB are known to persist and grow in the environment (Byappanahalli et al., 2003; Oh et al., 2012). Upon introduction to the environment, microbial contaminants are subject to highly variable dispersal and decay processes (Korajkic et al., 2019; Stewart et al., 2013). Pathogens aside, there is also propagation of resistance genes via recreational water; some of which in *E. coli*. (Farrell et al., 2023; Leonard et al., 2015; Singh et al., 2022; Han et al., 2022; Li et al., 2022) This study also highlighted the presence of AMR genes in the river samples and Farrell et al. (2023) link recreational water use to an increased carriage of antimicrobial-resistant organisms and Singh et al. (2022) describe how wastewater and natural water systems act as vectors for the spread of ARGs into recreational waters.

The need to differentiate faecal sources in recreational waters led to the emergence of microbial source tracking (MST) methods in the early 2000s, most notably the PCR-based assays that target the 16S rRNA gene in *Bacteroides* spp. (Bernhard et al., 2000; Dick et al., 2005; Schriewer et al., 2013). Some studies have found strong relationships between the MST markers and enterococci (Schriewer et al. 2010), while other studies have found either weak or no relationships (Flood et al., 2010; Santoro and Boehm, 2007), many of which are discussed in a review by Harwood et al. (Harwood et al., 2014). One main factor affecting the relationship between enterococci and the relative strength of different sources of fecal contamination is that enterococci can persist and grow in the environment, which can significantly influence their concentrations in recreational water (Byappanahalli et al., 2012). Enterococci have been shown to persist in fresh water sediments and marine sediments and in some cases, their relative concentrations in sediments are several orders of magnitude higher than that in the overlying water (Anderson et al., 2005; Ferguson et al., 2005).

Previous studies have shown that Bacterisk ERU data also correlates well with enterococci culture data (Good et al., 2024). Results presented here also confirm that Bacterisk ERU data correlate with enterococci data from fresh waters. While enterococci do not contain endotoxin, this relationship probably reflects the decay of *E. coli* and other Gram-negative bacteria providing endotoxin that is detected while enterococci persist. Thus, detection of endotoxin is a useful indirect proxy for the presence of enterococci.

There is clearly a need for more comprehensive water quality monitoring that extends beyond traditional FIB assessments. Techniques such as nanofluidic quantitative real-time PCR (qPCR) could allow simultaneous detection of multiple FIB, MST, and pathogen genes in under 4 hours (Shahraki et al., 2019). However, the expense, complexity and necessary expertise likely preclude the routine application of such methods for direct pathogen detection. Protecting public health in recreational waters remains an important goal.

While FIB remain an essential tool for assessing faecal contamination, they are insufficient for evaluating the broader spectrum of waterborne health risks. A more holistic approach to monitoring and managing recreational water quality is needed to ensure the safety of all users and to address the emerging challenges posed by non-fecally transmitted pathogens. Results from the current study confirm the applicability of endotoxin detection as a rapid method for the risk assessment of recreational water quality. It is not only rapid (15 minutes) but will alert to the presence of non-FIB pathogens vital to the protection of human health. We suggest that such a method could form a key component in the development of Quantitative microbial risk assessment (QMRA) to provide a comprehensive evaluation of recreational water quality.

Conclusion

Faecal contamination of water continues to be a major public health concern, with new challenges necessitating a renewed urgency in developing rapid and reliable methods to detect contamination and prevent human exposures. Non-faecally transmitted bacteria in recreational waters pose significant health risks to users, underscoring the limitations of relying solely on faecal indicator bacteria (FIB) to assess water quality. Outbreaks linked to these bacterial pathogens often expose gaps in standard monitoring practices, which can leave recreational users unaware of potential dangers. As a result, a water body considered safe based on FIB levels may still harbour significant health risks from environmental bacteria. This study highlights the need for more comprehensive water quality monitoring to extend beyond traditional FIB quantifications. Following the rapid screening and detection especially of 'non-FIB', pathogen-specific testing should be incorporated in monitoring programmes, particularly in high-risk recreational waters. Environmental surveillance is also crucial, with factors such as temperature, salinity, and nutrient levels monitored to predict conditions conducive to pathogen proliferation. This study has shown that bacterial endotoxin, measured in near real-time with the Bacterisk assay, is a reliable marker not only of faecal contamination but also of the presence of potential pathogens of non-faecal origin. The portability and ease of use of this assay allows its convenient use to provide data on water quality at different locations and at different times providing a comprehensive surveillance tool.

Supplementary Materials: The following supporting information can be downloaded at the website of this paper posted on Preprints.org.

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