Evaluation of the novel substrate RUG™ for the detection of *Escherichia coli* in water from temperate (Zurich, Switzerland) and tropical (Bushenyi, Uganda) field sites†

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Direct testing of water quality to promote drinking water safety contributes to the sustainable development goals, which call for universal access to safely-managed drinking water services by 2030. Enzyme-substrate tests offer a potentially simple and reliable approach for the detection and quantification of fecal indicator bacteria, including *Escherichia coli* (*E. coli*). The novel aquatest (AT) based on resorufin-β-D-glucuronide methyl ester (RUG™) (AT-RUG) is an enzyme-substrate test that overcomes several drawbacks of other established tests. In this study, AT-RUG was used to detect and quantify *E. coli* in water from temperate (Zurich, Switzerland) and tropical (Bushenyi, Uganda) regions. Quantitative results of AT-RUG were compared with IDEXX Colilert-18® (C-18), m-TEC and m-ColiBlue24®. In temperate waters, AT-RUG was found to be as sensitive as m-TEC (97.0%) and C-18 (98.5%) and showed strong agreement with the reference methods. The false-positive rate for *E. coli* detection in temperate waters using AT-RUG was 6%. AT-RUG performed well at incubation temperatures of 37 °C and 45 °C, but not at 24 °C. In tropical waters, AT-RUG sensitivity was 94.1% compared to m-ColiBlue24®. AT-RUG detected significantly more *E. coli* than m-ColiBlue24®, suggesting it is a more conservative estimate. At both field sites, AT-RUG was able to effectively indicate categorical concentrations of *E. coli* in water samples indicating the level of risks of fecal contamination of water supplies. This study indicates that AT-RUG is a reliable and accurate medium for the detection and quantification of *E. coli* in temperate and tropical waters.

1 Introduction

Over two billion people did not have access to safely-managed drinking water services in 2015.1 According to the Joint Monitoring Programme, safely-managed drinking water services are indicated in part by the absence of “fecal and priority contamination”. The recommended measure of fecal contamination by the World Health Organisation (WHO) is the presence of fecal indicator bacteria such as *Escherichia coli* (*E. coli*) in a 100 mL water sample tested at the point of delivery.2 When fecal indicator bacteria are present in a sample, the concentration is also suggested to be an indicator of health risks. For example, the WHO classifies risks as “safe or low”, “intermediate”, “high”, or “very high” for waters with <1, 1–10, 11–100, >100 fecal coliform count per 100 mL, respectively.3,4 There is a need for simple, reliable and inexpensive microbial tests capable of determining the levels of fecal contamination to reduce waterborne disease.5

Enzyme-substrate tests offer a potentially simple approach to detect and quantify *E. coli* in drinking water.6 Enzyme-substrate tests typically rely on the combination of an enzyme substrate with a medium that supports the selective recovery
and growth for Gram-negative organisms.\textsuperscript{7} Often the tests are based on the metabolism of 4-methylumbelliferyl-β-D-glucuronide (MUG) to generate fluorescence.\textsuperscript{8} The targeted enzyme is β-glucuronidase, which is produced by most strains of \textit{E. coli} but only by a few other bacteria.\textsuperscript{7,9}

One of the more common enzyme–substrate assays for the detection and quantification of \textit{E. coli} is the IDEXX Colilert with Quanti-Tray\textsuperscript{®}/2000 system (IDEXX, Ludwigsburg, Germany), which quantifies \textit{E. coli} using the most-probable-number (MPN) method based on metabolism of MUG.\textsuperscript{10} Tests for detection of \textit{E. coli} based on MUG are very sensitive\textsuperscript{7} but have a number of disadvantages.\textsuperscript{11} \textit{E. coli} detection is based on fluorescence under ultraviolet (UV). The UV lamp requires access to electricity or batteries and entails a extra cost and a physical burden for field testing. Additionally, non-target microorganisms such as \textit{Aeromonas} spp. and \textit{Pseudomonas fluorescens} interfere with the fluorescence signal and can lead to false positive results.\textsuperscript{12,13} To reduce false positives, antibiotics (\textit{i.e.} ceftulodin) are typically included in MUG-based media, but antibiotics lead to further complications for field-based methods (\textit{i.e.}, supply chains, limited shelf life).\textsuperscript{11} An alternative to fluorescent substrates is the chromogenic substrate resorufin-β-D-glucuronide methyl ester (RUG\textsuperscript{TM}). RUG\textsuperscript{TM} is the methyl ester analog to resorufin-β-D-glucuronide (REG). REG has previously been shown to detect \textit{E. coli} at very low concentrations\textsuperscript{14} using the same enzyme (β-glucuronidase enzyme) that is used for \textit{E. coli} detection with MUG. The β-D-glucuronidase enzyme cleaves REG producing the highly fluorescent and pink molecule resorufin,\textsuperscript{11} that is easily visually detected by eye without the need for a UV source.\textsuperscript{15} Although the free acid agent REG is expensive to produce, the methyl ester analog RUG\textsuperscript{TM} is not (250 times less expensive than REG), and therefore has potential applications in low cost water quality testing.\textsuperscript{15} Additionally, RUG\textsuperscript{TM} has been shown to work as an \textit{E. coli} indicator with efficiency comparable to REG. Nevertheless, one limitation of both RUG\textsuperscript{TM} and REG is the requirement of oxygen to prevent the bleaching of resorufin through reduction which reduces the intensity of the color change.\textsuperscript{11} Therefore RUG and REG are not compatible with commonly used testing vessels that do not allow oxygen diffusion, \textit{e.g.}, Quanti-Tray\textsuperscript{®}/2000 system.

The objective of this study was to evaluate the performance of a media containing RUG\textsuperscript{TM} (AT-RUG) for the detection and quantification of \textit{E. coli} in water samples. AT-RUG was evaluated for the detectability of \textit{E. coli} and other bacterial strains (\textit{i.e.}, \textit{Proteus mirabilis}, \textit{Enterococcus faecalis}) that are commonly found in natural or drinking waters and may lead to false positives. Quantification performance of AT-RUG was further evaluated through direct comparison with other commonly used \textit{E. coli} detection reference methods (IDEXX Colilert Quanti-Tray/2000\textsuperscript{®}, m-TEC) in a temperate (Zurich, Switzerland) and (m-ColiBlue24\textsuperscript{®}) tropical (Bushenyi, Uganda) field site. For comparison with other methods, a field-portable MPN protocol was developed to overcome the limitation of an oxygen requirement for RUG\textsuperscript{TM}.

2 Material and methods

2.1 Preparation of AT-RUG

\textit{E. coli} was detected using AT-RUG (Biosynth\textsuperscript{®} AG, Staad, Switzerland; Thermo Fisher Scientific, Waltham, MA, USA) at concentration of 1 g per 100 mL sample, based on the findings of Magro \textit{et al.} 2014\textsuperscript{11} that 30 μM REG yields a very clear signal for the detection of \textit{E. coli}. In brief, 1 g ± 0.05 AT-RUG was poured into Whirl-Pak\textsuperscript{®} bags (120 mL capacity, Nasco, Fort Atkinson, WI, USA) along with 100 mL pipetted water samples. The samples were hand mixed to dissolve AT-RUG.

2.2 Strain challenge

The growth of \textit{E. coli} and other bacterial strains commonly found in natural waters was assessed in AT-RUG as compared to Colilert-18\textsuperscript{®} reagent (C-18) (IDEXX Laboratories, Ludwigsburg, Germany), lysogeny broth (LB) (Merck, Darmstadt, Germany) and phosphate-buffered saline water (PBS)\textsuperscript{16} as control. The bacterial strains tested included \textit{E. coli} (ATCC 25922), \textit{Enterococcus faecalis} (ATCC 47044), \textit{Klebsiella pneumoniae} (B02624), \textit{Pseudomonas aeruginosa} (G445), \textit{Enterobacter cloacae} (R1288), \textit{Proteus mirabilis} (G464), \textit{Klebsiella aerogenes} (DSM 30053) and \textit{Aeromonas hydrophila} (DSM 30187). All bacterial strains were grown overnight (15–20 hours) in LB at 37 °C and then five times serially diluted 10-fold. A 20 μL inoculum was added to 180 μL growth medium (AT-RUG, C-18, LB, or PBS) within a 96-well plate (Greiner Bio-One, Kremsmunster, Austria) and incubated for 20 hours at 37 °C within a plate reader (Software Gen5 2.04, BioTek, Sursee, Switzerland). The 96-well plate was sealed with a membrane of polyurethane (Breathe-Easy\textsuperscript{®}, Merck, Darmstadt, Germany) during incubation. Growth curves were generated by measuring the optical density (OD) at wavelengths of 365, 450, 565 and 600 nm every 20 minutes. After incubation, wells were scored visually for β-glucuronidase activity as positive or negative. Positive wells showed pink color or blue fluorescence for AT-RUG or C-18 respectively. Each experiment included five biological replicates. The experiment was repeated twice (two technical replicates). The mean and standard errors for each OD reading were determined from the biological replicates. The OD limit of detection (LOD) was determined to be 0.1, corresponding to the lowest measured OD value of the blank (PBS).

2.3 Temperate site (Zurich, Switzerland)

2.3.1 Samples. Seventy samples of approximately 0.5 L were collected in autoclaved glass bottles from rivers, lakes and wetlands in and around Zurich, Switzerland in May and June 2018 (Table 1). Samples were stored at 2–8 °C for transport and processed within six hours at Eawag in Duebendorf. The pH of water samples was measured on-site using pH indicator strips (Merck, Darmstadt, Germany). Alongside collected samples, a negative (PBS) control and a positive
2.3.2 Quantification of E. coli using AT-RUG as compared to IDEXX Colilert Quanti-Tray®/2000® and membrane filtration with m-TEC. In Zurich, the performance of AT-RUG was compared to two established reference methods for the quantification of E. coli in water: C-18 with IDEXX Quanti-Tray®/2000® system and membrane filtration with m-TEC. Aliquots were taken from the same sample (0.5 L) and processed in parallel by all three methods.

E. coli quantification with AT-RUG was performed using an adaptation of the IDEXX Quanti-Tray® MPN method. After samples were mixed with AT-RUG, the sample was distributed into the IDEXX Quanti-Tray®/Legiolert. As mentioned, detection of E. coli with AT-RUG requires exposure to oxygen to maintain colorimetric detection. Unlike the traditional Quanti-Trays used with C-18, the Quanti-Tray®/Legiolert has a semi-permeable membrane that allows gas exchange – including oxygen – during incubation. After 18 to 20 hours incubation at 37 °C, each well of the tray was scored as negative or positive for the presence of E. coli. Positive wells were identified as those which turned from orange to pink. Negative and positive controls were used as references when counting positive wells. Based on the number of positive and negative wells, the MPN value was determined using the MPN table for Quanti-Trays®/Legiolert provided by IDEXX. The number of E. coli was reported as MPN per 100 mL with lower and upper 95% confidence limits. The lower and upper LOD for the Quanti-Tray®/Legiolert system is 1 MPN per 100 mL.

Membrane filtration with m-TEC ChromoSelect agar (Merck, Darmstadt, Germany) was also used to process samples (100 mL and 10 mL aliquots) according to standard methods using 0.45 μm pore size membrane S-PAK-filters (47 mm, Merck, Darmstadt, Germany). Incubation for m-TEC was 44.5 °C for 22 to 24 hours. E. coli were quantified and reported as colony forming units (CFU) per 100 mL for the membrane filtration method. The upper counting limit was 200 CFU per filter. Thus, with two sample volumes (10 mL and 100 mL), the lower and upper LODs were 1 CFU per 100 mL and 2000 CFU per 100 mL respectively.

2.3.3 Laboratory comparison of E. coli growth in media. The performance of Quanti-Tray®/Legiolert system with AT-RUG for the quantitative detection of E. coli was also compared to m-TEC and C-18 with Quanti-Tray®/2000® system in the laboratory using E. coli (ATCC 25922) reference strain. In brief, a two-fold dilution series in PBS of overnight E. coli culture was quantified simultaneously using AT-RUG and C-18 as compared to m-TEC. For comparison, the range of detection was set to that obtained by m-TEC, with a lower and upper LOD of 1 CFU per 100 mL and 200 CFU per 100 mL respectively. The lower limit of quantification (LLOQ) was calculated based on the residual standard deviation and the slope of the linear regression of C-18 with m-TEC. The residual standard error was 0.144 and the slope 1.13 leading to a LLOQ of 1.2 log10 MPN per 100 mL (15.8 MPN per 100 mL).

2.3.4 Presumptive E. coli isolation and species-level identification. Presumptive E. coli isolates exhibiting color change in the AT-RUG media were further tested to determine (diluted E. coli stock solution) control were processed by all methods on each sampling day.

### Table 1: Study sites and methods with number of samples (n) below and above the LOD for the correlation analysis

| Study sites | n       | Sample types (n) | pH       | Methods                                           | Lower LOD | Upper LOD | n < lower LOD | n > upper LOD |
|------------|---------|-----------------|----------|--------------------------------------------------|-----------|-----------|---------------|---------------|
| Zurich, Switzerland (temperate) | 70      | Rivers (30)     | 7.4–7.7  | AT-RUG with Quanti-Tray®/Legiolert (MPN per 100 mL) | 1         | 2272.6    | 59*           | 2             | 3             |
|            |         | Lakes (35)      |          | C-18 with Quanti-Tray®/2000® (MPN per 100 mL)     | 1         | 2419.6    |               | 2             | 13            |
|            |         | Wetlands (5)    |          | m-TEC with membrane filtration (CFU per 100 mL)   | 1         | 200       |               | 3             | 26            |
| Bushenyi, Uganda (tropical) | 85      | Springs (50)    | 5.0–6.8  | AT-RUG with deep 96-well plates (MPN per 100 mL)  | 1         | 430       | 85            | 18            | 5             |
|            |         | Swamps (6)      | 5.3      |                                                  |           |           |               |               |               |
|            |         | Lakes (15)      | 7.0      |                                                  |           |           |               |               |               |
|            |         | Wetlands (8)    | 5.3–5.5  |                                                  |           |           |               |               |               |
|            |         | Rivers (2)      | 6.9      |                                                  |           |           |               |               |               |
|            |         | Dams (4)        | 5.0–5.3  |                                                  |           |           |               |               |               |

* Only 59 of the 70 collected samples were analyzed because one sample was not readable and ten samples were above the upper LOD.

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specificity of AT-RUG, using a modification of the methods described in Julian et al. 2015. In brief, 1 mL sample was collected from up to two presumptive positive large wells from each Quanti-Tray®/Legiolert. The sample was streaked onto either MacConkey agar (Merck, Darmstadt, Germany), \( n = 52 \), or m-TEC ChromoSelect agar, \( n = 60 \). Samples were incubated for 18 to 24 hours at 37°C (MacConkey) or 45.5°C (m-TEC). After incubation, colony morphology types were recorded and between one and five morphology types from each agar plate were inoculated into 2 mL of AT-RUG in oxygen permeable culture tubes to confirm the isolate was the cause of the previously observed color change. The tubes were incubated at 37°C for 18 to 20 hours. Both MacConkey and m-TEC were used because isolation with MacConkey was insufficiently selective; bacterial isolates grown on MacConkey frequently failed to produce color change in subsequent incubation in AT-RUG. The \( E. coli \) isolates that exhibited pink color in the AT-RUG media again were streaked to MacConkey agar and incubated at 37°C for 18 to 24 hours. If more than one isolate from the same positive large well met this criterion, only one was considered. After incubation, the plates were observed for uniform colony morphology before proceeding.

One colony from each plate was then tested using an oxidase test followed by an API-20E biochemical test strips (bioMérieux Inc., Marcy l’Etoile, France) according to the manufacturer’s instructions. Control strains for the API-20E included \( E. coli \) (ATCC 25922), \( K. pneumoniae \) (B02624), \( E. cloacae \) (R1288) and \( P. mirabilis \) (G464).

2.3.5 Incubation temperature sensitivity of AT-RUG. Aliquots of 18 water samples were used in combination with the Quanti-Tray®/Legiolert with AT-RUG media to test the sensitivity of AT-RUG to different temperatures. Samples were incubated at either 24, 37 or 45.5°C for up to 42 hours. The MPN was scored after 18 to 20 hours, 24 and 42 hours. \( E. coli \) estimates at 24°C and 45°C were compared to \( E. coli \) estimates at 37°C for different incubation times.

2.4 Tropical site (Bushenyi, Uganda)

2.4.1 Sample collection. Eighty-five water samples were collected in autoclaved plastic bottles from lakes, springs, wetlands, rivers, dams and swamps in Bushenyi district, Uganda, in June and July 2018 (Table 1). Samples were stored at 2–8°C and processed within six hours at the laboratory of the National Water and Sewerage Corporation at the Nyaruzinga water treatment plant in Bushenyi district. The pH was measured at each sampling site using pH indicator strips (Merck, Darmstadt, Germany). A negative (distilled water) control and a positive (diluted chicken feces) control were processed by all methods alongside collected samples on each sampling day.

2.4.2 Quantification of \( E. coli \) using AT-RUG as compared to m-ColiBlue24® with membrane filtration. To quantify \( E. coli \) using AT-RUG in Bushenyi, Uganda, the MPN approach was adapted for use with deep 96-well plates (BRAND, Wertheim, Germany). Deep 96-well plates were used because neither Quanti-Tray®/Legiolert nor an appropriate tray sealer were available in Bushenyi at the time of the study. Each well within the 96-well plate holds a maximum volume of 1.2 mL. The 96-well plates also allow sufficient oxygen exchange (none of the wells turning colourless during incubation), which is a prerequisite for use of AT-RUG.

In brief, 100 mL water samples were transferred into the wells (approximately 1.05 mL per well) using a 10 mL serological pipette with 0.1 mL graduations (Greiner Bio-One, Kremsmunster, Austria). The plates were incubated at 37°C for 18 to 20 hours. After incubation, each well of the tray was scored as negative (orange color) or positive (pink color) for the presence of \( E. coli \), and the \( E. coli \) concentration was estimated using an MPN calculator in Microsoft Office Excel 2010, available from Jarvis et al. 2010.20 For reporting, \( E. coli \) concentrations were adjusted to MPN per 100 mL with lower and upper 95% confidence limits. The lower and upper LOD for deep 96-well plates is <1 MPN per 100 mL and 430 MPN per 100 mL respectively.

The performance of AT-RUG was compared to the reference medium m-ColiBlue24® (Hach, Duesseldorf, Germany) using membrane filtration according to standard methods.21 In brief, 100 mL water samples were filtered through 0.45 μm membrane S-PAK-filter (47 mm, Merck, Darmstadt, Germany). The m-ColiBlue24 Petri dishes were incubated at 37°C for 22 to 24 hours. \( E. coli \) colonies were quantified and reported as CFU per 100 mL for the membrane filtration method. The upper and lower LOD for the enumeration of \( E. coli \) by membrane filtration were 200 CFU per 100 mL and 1 CFU per 100 mL respectively. The countable range of agar plates (1–200 CFU) was expanded beyond the recommended range of 20–80 CFU18 to reduce the resources (both material and time) needed. The expanded countable range allows for a 10-fold dilution series as oppose to the recommended 5-fold dilution series, reducing resource use by 30–50%. Bacterial counts below 20 CFU may lead to increased variation in counts; bacterial counts above 80 CFU per plate may lead to underestimation as at higher counts bacterial colonies crowd.

2.5 Statistical analysis

Data for each water sample were recorded and entered into Microsoft Office Excel 2010 prior to analysis in the statistical analysis software R (version 3.4.1, R Foundation for Statistical Computing, Vienna, Austria). Scatter plots were generated using Microsoft Office Excel 2010. Censored data were set to half the lower LOD or to the upper LOD (Table 1) according to Cole et al. 2009.22 Correlation analysis between log-transformed microbial counts was assessed using linear regression or Spearman’s rank correlation. Paired sample \( t \)-tests were used to determine the statistical significance of the mean difference \( (\Delta d) \) in log-transformed \( E. coli \) concentrations between AT-RUG and the reference method.23 Statistical significance was defined at \( \alpha = 0.05 \).
Sensitivity and specificity metrics of the methods were calculated as defined in the United States Environmental Protection Agency (US EPA) protocol.24 False positives were defined here as positive for E. coli (≥1 E. coli count per 100 mL) when tested with the AT-RUG method, but negative for E. coli (<1 E. coli count per 100 mL) when tested with the reference method (either m-TEC for Zurich or m-ColiBlue24® for Busheyeni). False negatives are negative for E. coli (<1 E. coli count per 100 mL) when tested with the AT-RUG method, but positive for E. coli (≥1 E. coli count per 100 mL) when tested with the reference methods. For correlation analysis, samples were excluded when the results were not readable (n = 1), when results were not quantitative because the E. coli concentrations exceeded the upper LOD for a 100 mL sample, and when no dilution was performed (n = 10). For Zurich, 59 out of 70 samples were analyzed; in Busheyeni, 85 out of 85 samples were analyzed.

3 Results

3.1 Strain challenge

Out of the eight bacterial strains tested, only the E. coli strain tested was positive for AT-RUG (pink coloration) or C-18 (blue fluorescence). All other strains tested showed no detectable color or fluorescence change. E. coli reference strain growth had similar time-to-detection for AT-RUG, C-18 and LB broth (Fig. 1, with time-to-detection defined as when optical density exceeded the background OD of 0.1). Notably, all tested strains grew in AT-RUG; all strains except Enterococcus faecalis and Aeromonas hydrophila grew in C-18 (Fig. S1–S7†). The suppression of two strains in C-18 is attributable to the presence of an antibiotic as a growth inhibitor. Growth did not coincide with color (AT-RUG) or fluorescence (C-18) demonstrating specificity of the detection methods to the presence of the β-glucuronidase enzyme. Comparison of the growth curves for a specific strain generated by different wavelengths showed that 365 nm is the most suitable for all media. However, growth was observable for all wavelengths tested (450, 565, 600 nm).

3.2 Temperate site (Zurich, Switzerland)

AT-RUG performance was similar to both m-TEC and C-18 at the temperate site. E. coli concentrations measured using AT-RUG correlated linearly with both m-TEC ($r^2 = 0.84$, slope = 0.99, 95% confidence interval = [0.88, 1.10], Fig. 2a) and C-18 ($r^2 = 0.96$, slope = 0.99, 95% confidence interval = [0.94, 1.05], Fig. 2b). The Spearman’s rank correlation coefficient was 0.907 and 0.950 for AT-RUG with m-TEC and C-18, respectively. A correlation of $r^2 = 0.82$ (slope = 0.85, 95% confidence

![Fig. 1](image1.png)

Fig. 1. Optical density (365 nm) of E. coli grown over 20 hours in LB, AT-RUG, C-18, and PBS. The LOD for the OD was set to 0.1.

![Fig. 2](image2.png)

Fig. 2. Scatter plots of E. coli estimates using AT-RUG versus (a) m-TEC and (b) C-18. Error bars show the 95% confidence interval of the MPN assays for AT-RUG and C-18.
interval = [0.74, 0.96]) and a Spearman's rank coefficient of 0.915 was obtained for C-18 with m-TEC (Fig. S8†). The mean difference in the estimated E. coli concentrations (log_{10} MPN per 100 mL) between AT-RUG and m-TEC or C-18 was not statistically significant (paired sample t-test, two-tailed, \( \mu_d = 0.066, t = 1.013, p = 0.315 \) for m-TEC and \( \mu_d = -0.033, t = -1.122, p = 0.267 \) for C-18) (Tables S1 and S2†). On the basis of categorical E. coli concentrations, AT-RUG generally agreed with both m-TEC and C-18 (Tables 2, 3 and S3†). The agreement based on the categorical E. coli concentrations was 0%, 15.9%, 14.5% and 55.1% between AT-RUG and m-TEC and 1.5%, 14.5%, 17.4% and 55.1% between AT-RUG and C-18 for the concentration categories of <1, 1–10, 11–100 and >100 MPN or CFU per 100 mL respectively. The low agreement in the <1 MPN or CFU per 100 mL category was due to the low number of water samples collected with <1 MPN or CFU per 100 mL (no sample for m-TEC, the defined reference method).

Notably, of the 69 m-TEC agar plates processed, 20 (29%) had colony counts between 1–20 CFU and 27 (39%) between 80–200, whereas 1–20 and 80–200 represent colony counts outside of the recommended range of the U.S. EPA method 1603.18 For the 69 samples analyzed, sensitivity was high but specificity was low due to the low number of samples without E. coli. Specifically, sensitivity of AT-RUG was 97.0% (95% confidence interval = [89.5, 99.6]) for m-TEC and 98.5% (95% confidence interval = [92.0, 100]) for C-18. Specificity for AT-RUG was 0% (95% confidence interval = [0, 70.8]) for m-TEC since no samples were negative for both methods, and as 50.0% (95% confidence interval = [1.3, 98.7]) for C-18 since only one sample was negative for both AT-RUG and C-18 (Tables S4 and S5†). C-18 compared to m-TEC (Table S6†) gave same results regarding sample positivity for E. coli, sensitivity and specificity as AT-RUG compared to m-TEC.

3.2.1 Laboratory comparison of E. coli growth in media. Evaluation of AT-RUG in Quanti-Tray®/Legiolert using a dilution series of cultured E. coli demonstrated good correlation with m-TEC in the quantification (15.8–200 CFU per 100 mL) and detection (1–200 CFU per 100 mL) range with \( r^2 = 0.99 \) (slope = 0.97, 95% confidence interval = [0.86, 1.09]) and \( r^2 = 0.90 \) (slope = 0.78, 95% confidence interval = [0.48, 1.08]) respectively (Fig. S9†). The E. coli concentrations were significantly correlated between AT-RUG and m-TEC (Spearman’s rank correlation coefficient of 0.955 in the detection range). Similar correlation was obtained for AT-RUG with C-18 with \( r^2 = 1.00 \) and 0.81 (slope = 0.92 and 0.85, 95% confidence interval = [0.56, 1.65] and [0.38, 1.33]) for quantification and detection range respectively. The Spearman’s rank correlation coefficient was 0.929 in the detection range. C-18 correlated with m-TEC with \( r^2 = 1.00 \) and 0.97 (slope = 0.95 and 1.13, 95% confidence interval = [0.60, 1.66] and [0.91, 1.36]) for quantification and detection range respectively. The Spearman’s rank correlation coefficient was 0.991 in the detection range.

3.2.2 E. coli confirmation tests. From 60 samples, 112 positive AT-RUG Quanti-Tray®/Legiolert wells were plated on MacConkey (52) and m-TEC (60) agar. From these, up to five isolates were tested for subsequent fluorescence and 84 (75.0%) showed fluorescence. An isolate was identified using API-20E kit in 83 (74.1%) wells that caused fluorescence in AT-RUG. From 83 of the fluorescence-causing isolates, 73 (88.0%) of the isolates were identified as E. coli with \( \geq 80.0\% \) likelihood that the isolates were E. coli based on their biochemical profiles (corresponding to acceptable, good, very good or excellent confidence). Of the other ten isolates, five isolates (6.0%) were identified as E. coli with low confidence (\( \geq 66.0\% \)), two were identified as Enterobacter cloacae (97.7% and 99.3% confidence), one as Citrobacter freundii (99.3% confidence), one as Proteus mirabilis with very good identification to the genus (93.5% confidence), and one as Klebsiella pneumoniae with only 31.4% identification confidence. Of all the isolates tested, 94% (78 out of 83) were identified as E. coli with at least low confidence. This corresponds to a false-positive rate of 6%. Including only isolates with acceptable or higher confidence the false-positive rate was 12%.

Using m-TEC to isolate and confirm bacterial strains causing colorimetric changes in AT-RUG from the Quanti-Tray®/Legiolert wells performed better than MacConkey. From the 112 positive tested for subsequent fluorescence, the content of 52 and 60 wells were streaked on MacConkey and m-TEC agar respectively. Using m-TEC agar, 58 out of the 60 wells (96.7%) showed fluorescence. However, out of the 52 samples tested on MacConkey, only 26 (50.0%) showed fluorescence. From 26 of the fluorescent causing isolates on MacConkey, 24 (92.3%) of the isolates were identified as E. coli with at least low confidence. From 54 (93.1%) out of the 58 fluorescent causing isolates on m-TEC, 54 (93.1%) were identified as E. coli with at least low confidence.

3.2.3 Incubation temperature sensitivity of AT-RUG. AT-RUG incubation at 24 °C performed substantially worse than incubation at 37 °C. Paired samples t-test for the lower-tailed

| AT-RUG [MPN per 100 mL] | m-TEC (CFU per 100 mL) | 11–100 | >100 |
|--------------------------|------------------------|--------|------|
| <1                       | 0% (0)                 | 0% (0) | 0% (0) |
| 1–10                     | 2.90% (2)              | 15.34% (11) | 14.39% (10) |
| 11–100                   | 1.45% (1)              | 2.90% (2)   | 1.45% (1)   |
| >100                     | 0% (0)                 | 0% (0)     | 1.45% (1)  |
| Total                    | 4.35% (3)              | 21.74% (13) | 15.94% (11) |

Percentages refer to the total sampling number.
alternative hypothesis indicate that the \(E.\) \textit{coli} estimates are significantly lower for samples incubated at \(24\) °C compared to \(37\) °C regardless of incubation time (Table 4). When AT-RUG was incubated for 18 hours at \(24\) °C, \(E.\) \textit{coli} estimates were always (18 of 18 total) below the lower LOD (Fig. S10†). Longer incubation times (24 and 42 hours) improved the performance only slightly, as correlations with incubation at \(37\) °C for 24 hours were low (\(r^2 = 0.36\), slope = 1.42, 95% confidence interval = [0.42, 2.41] at 24 hours; \(r^2 = 0.23\), slope = 0.53, 95% confidence interval = [0.01, 1.04] at 42 hours). Spearman’s rank correlation coefficients were moderate (0.456 at 24 hours and 0.326 at 42 hours), but not statistically significant.

In contrast, results from AT-RUG incubation at \(45\) °C were comparable to those obtained at \(37\) °C. Paired samples \(t\)-test for the two-tailed alternative hypothesis demonstrated no significant (18 and 42 hours incubation) or only slightly significant (24 hours incubation) difference between AT-RUG \(E.\) \textit{coli} estimates for samples incubated at \(45\) °C and \(37\) °C (Table 3). Linear regression showed \(r^2 = 0.93, 0.94\) and 0.96 (slope =1.07, 0.99 and 1.02; 95% confidence interval = [0.92, 1.23], [0.86, 1.13], [0.91, 1.13]) for 18, 24 and 42 hours respectively (Fig. S11†). Spearman’s rank correlation coefficients were significant with 0.951, 0.936 and 0.920 respectively.

### 3.3 Tropical site (Bushenyi, Uganda)

AT-RUG compared to m-ColiBlue24® showed a high level of agreement (Fig. 3). The two methods demonstrated good correlation with \(r^2 = 0.82\) (slope = 0.83, 95% confidence interval = [0.75, 0.92]) and a Spearman’s rank correlation coefficient of 0.898. The \(p\)-value of the paired \(t\)-tests for the upper-tailed alternative hypothesis was less than the significance level, providing evidence that the mean difference between AT-RUG and m-ColiBlue24® \(E.\) \textit{coli} estimates were significantly greater than zero (Table S7†).

When comparing AT-RUG with m-ColiBlue24® based on sample positivity for \(E.\) \textit{coli}, four samples (22.2%) were false negative and three samples (4.5%) were false positive (Table S8†). AT-RUG produced a sensitivity of 94.1 % (95% confidence interval = [85.6, 98.4]) and a specificity of 82.4% (95% confidence interval = [56.6, 96.2]) when compared with m-ColiBlue24®. The agreement between the two methods based on the categorical \(E.\) \textit{coli} concentrations was 16.5%, 7.1%, 37.7% and 16.5% for the concentration categories of <1, 1–10, 11–100 and >100 per 100 mL respectively (Table 4).

Of the 85 m-ColiBlue24® agar plates processed, 40 (47%) had colony counts between 1–20 CFU and 18 (21%) between 80–200, where 1–20 and 80–200 represent colony counts outside of the recommended range of the U.S. EPA method 1603 specified for m-TEC agar.18

### 4 Discussion

AT-RUG appears to be a reliable and accurate medium for the detection and/or quantification of \(E.\) \textit{coli} in water. Quantification of \(E.\) \textit{coli} using AT-RUG aligned with standard \(E.\) \textit{coli} field detection methods (C-18 with Quanti-Trap®/2000, membrane filtration with m-TEC and m-ColiBlue24®) when AT-RUG was combined with MPN methods that allowed for gas exchange (Quanti-Trap®/Legiolert, deep 96-well plates). AT-RUG offers an inexpensive alternative to other colorimetric methods for \(E.\) \textit{coli} detection and quantification.

#### 4.1 Benefits and limitations of AT-RUG quantification testing

AT-RUG is field portable and provides a suitable low-cost growth medium for water-quality monitoring.11 To permit

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**Table 3** Comparison of categorical concentrations of \(E.\) \textit{coli} from AT-RUG using Quanti-Tray®/Legiolert system and C-18 using Quanti-Tray®/2000 system. Percentages refer to the total sampling number

| AT-RUG (MPN per 100 mL) | C-18 (MPN per 100 mL) |
|-------------------------|------------------------|
| <1                      | 1.45% (1)              |
| 1–10                    | 14.49% (10)            |
| 11–100                  | 17.39% (12)            |
| >100                    | 55.07% (38)            |
| Total                   | 2.90% (2)              |

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**Table 4** Results of paired samples \(t\)-test for the lower-tailed alternative hypothesis including \(t\)-value, \(p\)-value, 95% confidence interval (CI) and mean difference (\(\mu_d\)) of AT-RUG \(E.\) \textit{coli} concentrations (log\(_10\) MPN per 100 mL) at \(24\) °C as compared to \(37\) °C incubation

| 24 °C incubation as compared to 37 °C | 24 °C incubation as compared to 37 °C |
|--------------------------------------|--------------------------------------|
| \(t\)                                 | \(p\)-Value                          | \(\mu_d\) | 95% CI                               |
| 18 hours                             | -5.042                               | <0.001    | -1.730 [-Inf, -1.133]                |
| 24 hours                             | -5.174                               | <0.001    | -1.396 [-Inf, -0.926]                |
| 42 hours                             | -2.321                               | <0.001    | -0.754 [-Inf, -0.189]                |

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**Table 5** Paired samples \(t\)-test for the two-tailed alternative hypothesis including \(t\)-value, \(p\)-value, 95% confidence interval (CI) and mean difference (\(\mu_d\)) of AT-RUG \(E.\) \textit{coli} concentrations (log\(_10\) MPN per 100 mL) at \(45\) °C as compared to \(37\) °C incubation

| 45 °C incubation as compared to 37 °C | 45 °C incubation as compared to 37 °C |
|--------------------------------------|--------------------------------------|
| \(t\)                                 | \(p\)-Value                          | \(\mu_d\) | 95% CI                               |
| 18 hours                             | 0.223                                | 0.826    | 0.021 [-0.175, 0.216]                |
| 24 hours                             | 2.161                                | 0.045    | 0.178 [0.004, 0.352]                 |
| 42 hours                             | 1.010                                | 0.327    | 0.068 [-0.074, 0.209]                |
quantification, AT-RUG was combined with Quanti-Tray®/Legiolert systems and deep 96-well plates, resulting in comparatively precise estimates of the concentration. The distinction between negative (orange) and positive (pink) AT-RUG wells after incubation was clear and we consider inter-operator variability in positive/negative classification unlikely with the AT-RUG substrate. Occasionally, a color gradient of white-to-pink would be observed as a result of limited diffusion of oxygen which is required to prevent loss of color. Gentle shaking of the plates/trays to promote increased oxygen diffusion resolved the color gradient, which universally resulted in a clearly pink well. Positive and negative controls were used as a reference for this study. For field work, references for AT-RUG classification are likely not needed.

A drawback of using the Quanti-Tray®/Legiolert system is the requirement for large, expensive equipment (Quanti-Tray® sealer PLUS) and is therefore difficult to perform in field settings. The application of AT-RUG with deep 96-well plates seems to be a good alternative method for detection and quantification of E. coli in resource limited settings and overcomes the challenges related to the redox of the resorufin dye. At the time of the study, the costs per deep 96-well plate were around $4.70 (Merck, May 8, 2018), but it is likely possible to produce cheaper alternatives. Bain et al. 2012\textsuperscript{23} estimated costs for membrane filtration and Colilert MPN methods ranging from $0.50 to $6.00 per sample. However, the use of AT-RUG in combination with deep 96-well plates requires no expensive additional equipment for processing. Furthermore, the deep 96-well plates can be autoclaved and therefore reused. The use of AT-RUG in combination with the MPN method allows implementation with minimal training and more widespread water quality monitoring in resource limited settings.

4.2 E. coli confirmation tests

False positive results of the test method can lead to unnecessary corrective actions and wasting of resources.\textsuperscript{26} For AT-RUG, confirmation testing produced a false-positive rate of 6% during laboratory testing in Switzerland. This is in the range of reported false-positive rates for C-18 between 1 and 8%\textsuperscript{27–29} but lower than the reported rate for C-18 of 36.4% in tropical waters.\textsuperscript{30} In contrast to C-18, AT-RUG does not require the addition of an antibiotic inhibitor of natural fluorescence and thus potentially allows growth of non-target organisms. However, the detected AT-RUG false positives Enterobacter cloacae, Citrobacter freundii and Klebsiella pneumoniae corresponded to the detected C-18 false positives.\textsuperscript{10} More research is needed regarding whether the AT-RUG false positives are common false positives for glucuronide-based detection methods or specific for AT-RUG.

MacConkey agar resulted in isolation and growth of a large proportion of non-target bacteria, which made it difficult to find the microorganisms responsible for detection. Therefore, E. coli colonies might not have grown or selected on the MacConkey agar leading to the high false-positive rate. Grabow et al. 1979\textsuperscript{30} evaluated different media for membrane filtration counting of total coliform bacteria and reported that usually the lowest average coliform counts were recorded on the MacConkey agar compared to reference methods. Therefore, a more specific agar like m-TEC is needed for confirmation testing.

4.3 Comparison of AT-RUG and m-ColiBlue24®

The paired sample t-test showed statistically significant differences for E. coli counts obtained based on AT-RUG with
dures, such as tightly controlled incubation temperatures.\textsuperscript{33} This article is licensed under a Creative Commons Attribution 3.0 Unported Licence.

4.4 Incubation temperature sensitivity of AT-RUG

\textit{E. coli} counts were significantly lower when AT-RUG was incubated at 24 °C compared to incubation at 37 °C or 45 °C, even when the incubation period was extended from 24 to 42 hours. AT-RUG did not work at lower temperatures. However, AT-RUG performed well at a high incubation temperature of 45 °C. Since in low-resource settings testing can be restricted by the requirement for laboratory facilities and testing procedures, it can be difficult to conduct tests in a manner that complies with regulations and standard operating procedures, such as tightly controlled incubation temperatures.\textsuperscript{33} Ambient-temperature incubation\textsuperscript{24} or field incubators, such as those developed by Schertenleib \textit{et al.} 2019 offer a potential low-cost solution.\textsuperscript{35} More research is needed regarding the possibility of ambient-temperature incubation with temperatures higher than 24 °C.

### Conflicts of interest

Thermo Fisher Scientific (Waltham, MA, USA) provided the pilot batch of AT-RUG used in this study. RB is named as a coinventor on International Patent Application number PCT/GB2010/050728 entitled “Apparatus for testing the quality of a fluid sample” and International Patent Application number PCT/GB2012/050452 of the same name.

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