



Article Studies on Simultaneous Enrichment and Detection of Escherichia coli O157:H7 during Sample Shipment

Chuyan Chen¹, Claudia P. Coronel-Aguilera¹, Bruce M. Applegate^{1,2,3,4,5,6,*}, Andrew G. Gehring⁷, Arun K. Bhunia^{1,3,4,5,6} and George C. Paoli^{7,*}

- ¹ Department of Food Science, Purdue University, West Lafayette, IN 47907, USA
- ² Department of Biological Sciences, Purdue University, West Lafayette, IN 47907, USA
- ³ Center for Food Safety Engineering, Purdue University, West Lafayette, IN 47907, USA
- ⁴ Purdue Institute of Inflammation, Immunology and Infectious Disease, Purdue University, West Lafayette, IN 47907, USA
- ⁵ Interdisciplinary Life Science Program (PULSe), Purdue University, West Lafayette, IN 47907, USA
- ⁶ Department of Comparative Pathobiology, Purdue University, West Lafayette, IN 47907, USA
- ⁷ Molecular Characterization of Foodborne Pathogens Research Unit, Eastern Regional Research Center,
- Agricultural Research Service, U.S. Department of Agriculture, Wyndmoor, PA 19038, USA
- * Correspondence: applegab@purdue.edu (B.M.A.); george.paoli@usda.gov (G.C.P.)

Abstract: The USDA-FSIS has zero tolerance for *E. coli* O157:H7 in raw ground beef. Currently, FSIS collects samples from beef processing facilities and ships them overnight to regional testing laboratories. Pathogen detection requires robust methods that employ an initial 15–24 h culture enrichment. This study assessed the potential of using the Φ V10*nluc* phage-based luminescence detection assay during enrichment while the sample is in transit. Parameters including phage concentrations, temperature, and media-to-sample ratios were evaluated. Results in liquid media showed that 1.73×10^3 pfu/mL of Φ V10*nluc* was able to detect 2 CFU in 10 h. The detection of *E. coli* O157:H7 was further evaluated in kinetic studies using ratios of 1:3, 1:2, and 1:1 ground beef in about 15 h at 37 °C. These results suggest that this approach is feasible, allowing the detection of a presumptive positive upon arrival of the sample to the testing lab. As the current cargo hold controlled temperature is required to be 15–25 °C, the need for elevated temperature should be easily addressed. If successful, this approach could be expanded to other pathogens and foods.

Keywords: *E. coli* O157:H7; Shiga toxin-producing *E. coli*; bacteriophage; culture enrichment; foodborne pathogen; pathogen detection

1. Introduction

Foodborne pathogens cause an estimated 9.4 million cases of illness annually in the United States. Of those, there are over 175,000 cases of illnesses caused by Shiga toxinproducing *Escherichia coli* (STEC) infections [1]. STEC commonly cause gastroenteritis and those that result in enterohaemorrhagic gastroenteritis are called (EHEC). The EHEC pathovar has a known pathology by attaching and effacing (A/E) lesions on intestinal epithelial cells, destroying microvilli and inhibiting actin function, and forming pedestals below the site of attachment. Infection by EHEC manifests into symptoms such as bloody diarrhea [2]. STEC infections are defined by the specific production of one or both Shiga toxins (Stx₁ or Stx₂), causing symptoms that range from mild to life-threatening, including stomach cramps, vomiting, and diarrhea [3]. *E. coli* O157:H7 infections are a main cause of hemolytic uremic syndrome, which is associated with severe health implications, including kidney failure, hemolytic anemia, and potential death [4]. The primary reservoir for STEC has been recognized to be cattle [5]. From 1982 to 2002, outbreaks found to be associated with the Shiga toxin producing *E. coli* O157:H7 were primarily foodborne, transmitted by



Citation: Chen, C.; Coronel-Aguilera, C.P.; Applegate, B.M.; Gehring, A.G.; Bhunia, A.K.; Paoli, G.C. Studies on Simultaneous Enrichment and Detection of *Escherichia coli* O157:H7 during Sample Shipment. *Foods* **2022**, *11*, 3653. https://doi.org/10.3390/ foods11223653

Academic Editor: Michel Federighi

Received: 30 September 2022 Accepted: 24 October 2022 Published: 15 November 2022

Publisher's Note: MDPI stays neutral with regard to jurisdictional claims in published maps and institutional affiliations.



Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). ground beef [6]. In addition, other sources of outbreaks have involved raw milk, vegetables, sandwiches, and water [7–10].

Since 1994, STEC O157:H7 has been under strict "zero-tolerance" enforcement in ground beef by the U.S. Department of Agriculture (USDA) Food Safety and Inspection Service (FSIS) [11]. This "zero-tolerance" policy means that any sample testing positive for STEC O157;H7 (even a single cell in a 325 g sample) is considered adulterated and not fit for consumption. In 2012, this policy expanded to include the six serotypes of non-O157 STEC, due to their increasing prevalence in causing foodborne illness in the U.S. [12]. With a low infectious dose of an estimated 10–100 cells, *E. coli* O157:H7 poses a considerable threat to public health [13,14]. Thus, it is critical that *E. coli* O157:H7 detection in food is carried out rapidly and accurately.

Traditional methods for detection of STEC O157:H7 and other foodborne pathogens involve culture media enrichment, isolation on solid selective/differential media, and biochemical identification of isolates, which can take several days to weeks to complete. Over the past few decades, numerous rapid methods have been developed that are typically immunologically (antibody)-based or nucleic acid-based (e.g., PCR) [15,16].

While these methods have grown very popular, government regulatory agencies, such as the USDA-FSIS, continue to employ microbiological methods to acquire the required bacterial isolate. For example, the FSIS method for detection of STEC O157:H7 and other regulated STEC in meat products involves an initial 15–24 h culture enrichment of a 325 g sample in 975 mL of modified tryptone soy broth (mTSB) [17,18], screening of the enrichment by PCR, separation of STEC by immunomagnetic separation, plating on modified Rainbow Agar, and testing isolated by O-antigen agglutination to acquire a presumptive positive isolate. Presumptive positive isolates are then grown on sheep blood agar, tested again for O antigen agglutination, identified using the Bruker MALDI Biotyper, and once again screened by PCR [17].

In October 1994, the USDA-FSIS commenced a verification sampling program for *E. coli* O157:H7 in order to verify process control, to stimulate industry testing, and to reduce the pathogen presence in raw ground beef. Over time, the testing program expanded to include other beef parts, such as trim and other components. In 2010, over 85% of samples analyzed were of raw ground beef [19], and in 2022, more than half of beef samples tested (62%) still consisted of ground beef [20]. Food samples for testing at FSIS labs are collected at the production facility by FSIS inspectors. The inspectors randomly and aseptically collect raw ground beef samples from the current day's production. The raw ground beef samples (325 g each) are placed into three Whirl-Pak bags, chilled, and sent to the testing laboratory. Assuming samples are shipped at 5 PM daily and received 8 AM the following morning, 15 h is the minimum time that a ground beef sample will spend in transit. The testing laboratory uses one or two samples for pathogen testing, and the third is held for additional analysis in the event of a positive sample [21].

We have previously described a bacteriophage with an integrated luminescence reporter for the detection of *E. coli* O157:H7 [22]. This reporter phage Φ V10*nluc* was constructed by cloning the genes for NanoLuc[®] luciferase [23] into the *E. coli* O157:H7-specific temperate phage Φ V10 [24,25]. NanoLuc[®] produces ATP-independent luminescence after reacting with furimazine substrate. While bacterial luciferase is 77 kDa, NanoLuc[®] luciferase is a small protein subunit which is only 19 kDa [26]. Its smaller size gives it flexibility in bioreporter construction. The light intensity produced by NanoLuc[®] has been found to be roughly 150-fold greater than either Firefly or *Renilla* luciferases [23]. The Φ V10*nluc* reporter is added during culture growth, and is capable of detecting as few as five cells of *E. coli* O157:H7 in 40 mL of ground meat slurry within 9 h [22].

This study assessed the potential of using the Φ V10*nluc* reporter system to detect *E. coli* O157:H7 in mTSB, leveraging the 15 or more hours that the sample spends in transit from the production facility to the FSIS lab. The potential to exploit shipment time for enrichment could allow presumptive positive samples to be rapidly identified soon after samples are received in the testing laboratories. In order to assess the potential of this

method, key factors including phage concentrations, temperature, and sample to media ratios were evaluated.

In addition, prior research indicated that $\Phi V10$ *nluc* phage preparations may be contaminated with NanoLuc[®] luciferase released from the cells during lytic propagation. This co-purification of phage and luciferase results in a significant background luminescence upon addition of the enzymatic substrate. In this study, a previously reported method for phage purification was modified and evaluated for its efficacy in removing the luciferase contaminant.

2. Materials and Methods

2.1. Media Preparation

All media components were purchased from Fisher Scientific (Waltham, MA, USA) unless otherwise noted. Lysogeny broth (LB) was prepared by adding 10 g of tryptone powder, 10 g of sodium chloride, and 5 g of yeast per liter of deionized water. All LB media were adjusted to a pH of 7.5. The LB plates and top agar were made by adding 17 g and 6 g, respectively, of agar to 1 L of the LB media. When used, kanamycin (IBI Scientific, Las Vegas, NV, USA) was added to the LB plates at a final concentration of 50 μ g/mL. Modified tryptone soya broth (mTSB) was made according to the USDA-FSIS Media and Reagents guideline (USDA-FSIS 2022).

2.2. E. coli O157:H7 Culture Preparation

The *E. coli* strain C7927 used in this study is a human STEC serotype O157:H7 isolate from an apple cider outbreak [27]. A single colony of *E. coli* O157:H7 C7927 from an LB agar plate was inoculated in 100 mL of LB liquid media and grown overnight at 37 °C with shaking (100 rpm). After overnight incubation, 100 μ L of serial dilutions (1:10) were spread onto LB agar plates to enumerate the number of colony-forming units (CFU)/mL of the overnight culture.

2.3. Phage Purification Procedure

In order to prepare the bacteriophage, 50 μ L of a Φ V10*nluc* lysogen of *E. coli* O157:H7 C7927 [22] was inoculated in one liter of LB containing kanamycin (50 μ g/mL) and incubated overnight on a shaker at 100 rpm at 37 °C. Following overnight incubation, 4 mL of chloroform (Fisher Scientific, Waltham, MA, USA) was added to the culture in order to permeabilize the cell membrane and to complete cell lysis. In addition, 48 g of sodium chloride was added and dissolved with constant stirring. The solution was then centrifuged at $15,000 \times g$ for 10 min in order to collect bacteria and cell debris into a pellet. The supernatant containing the phage was collected, and 80 g of polypropylene glycol (Fisher Scientific, Waltham, MA USA) per liter was dissolved into the solution using slow agitation. The solution was stored at 4 °C overnight to precipitate the phage. The phage solution was then centrifuged at $17,000 \times g$ for 15 min in order to collect the phage. The supernatant was carefully decanted, and the phage pellet was resuspended in phage buffer (50 mM Tris, 100 mM MgCl₂, pH 7.6) and vacuum filtered through a 0.45 μ m pore size membrane (Fisher Scientific, Waltham, MA, USA). Subsequently, the filtrate was passed through a second filter with pore size of 0.1 µm (Fisher Scientific, Waltham, MA, USA). The resulting solution was collected and stored in a sterile 50 mL Falcon screw cap tube at 4C.

When using the Φ V10*nluc* phage prepared as described above, phage infection assays yielded significant background light upon addition of the Nano-Glo[®] reagent (Promega Madison WI; see Section 2.4.). Furthermore, addition of the Nano-Glo[®] reagent directly to the Φ V10*nluc* phage preparations resulted in light emission, suggesting the presence of free luciferase protein in the phage preparations. Thus, a subsequent purification procedure was conducted to remove the free luciferase and reduce the background light emission from the Φ V10*nluc* phage preparations, as briefly described here. An Amicon[®] Ultra-15 (MilliporeSigma, Burlington, MA, USA) centrifugal filter device that contained a membrane with a 100,000 kDa cutoff was used to separate the luciferase from the Φ V10*nluc* phage. Twelve (12) mL of phage preparation was added to each centrifugal filter device, which was then placed in a fixed-angle rotor centrifuge and spun at $5000 \times g$ for 15 min per cycle, for a total of 7 cycles. Since the NanoLuc[®] (19 kDa) protein is smaller in size compared to the phage (>100 kDa), the phage is concentrated on the filter while the protein is washed out through the filter. After every cycle the filtrate was measured for NanoLuc[®] luminescent activity (described below) and the resuspended phage concentration was determined. The purified and concentrated Φ V10*nluc* phage was then resuspended in phage buffer for storage at 4 °C until further analysis.

2.4. ΦV10nluc Phage Titer Determination

Coomassie Brilliant Blue G-250 dye (Bio-Rad laboratories, CA USA) was added at 1% wt./vol to top agar in order to enhance the contrast of phage plaques, increasing their visibility and making them easier to count [22]. For plaque assays, 200 μ L of an overnight culture of *E. coli* O157:H7 C7927 and 100 μ L of serial dilutions of the Φ V10*nluc* phage solution were added to melted blue top agar, vortexed, and then poured onto LB plates. Plates were incubated in a 37 °C incubator for 18 h and plaques were enumerated the next day for phage titer determination. Plaque assays were conducted in triplicate.

2.4. Nano-Glo[®] Reagent Preparation

The Nano-Glo[®] reagent (Promega, Madison, WI, USA), which produces ATP-independent luminescence upon the oxidation of furimazine by the NanLuc luciferase, was used to measure presence of NanLuc luciferase according to the instructions from the manufacturer. An aliquot of 20 μ L of Nano-Glo[®] substrate was added to 1 mL of Nano-Glo[®] buffer. The resulting reagent was vortexed and either used immediately or stored at 4 °C for later use.

2.5. Effects of Phage Concentration on Time to Detection

Initial experiments were performed in LB to determine the relationship between phage concentrations (10^2-10^5 pfu/mL) and time to detection with *E. coli* O157:H7, ranging from approximately 2 to 2 × 10⁵ CFU per assay. Each assay consisted of a total of 40 mL.

Ten-fold dilutions of cells from 10^{S5} to 10^{-11} were prepared from an overnight culture for phage assays. One hundred microliters of each dilution were spread on LB plates in triplicate and incubated overnight at 37 °C to determine the initial number of CFU/mL in the overnight culture.

2.6. Characterization of the Growth of E. coli O157:H7 and Corresponding Φ V10nluc Lysogen

Growth of the wild-type *E. coli* O157:H7 C7927 and the C7927 Φ V10*nluc* lysogen was characterized by measuring the OD₆₀₀ of cultures grown in mTSB. A colony of C7927 Φ V10*nluc* lysogen was tested in 1 mL of LB and 10 µL Nano-Glo[®] reagent for luminescence prior to inoculation of an overnight growth. A 20µL aliquot of an overnight culture of wild-type and Φ V10*nluc* lysogen strains was inoculated in 100 mL mTSB flasks, then incubated at 37 °C with shaking at 100 rpm. Growth curves were performed in triplicate. Optical density measurements (OD₆₀₀) were taken using a BioPhotometer (Eppendorf North America, Enfield, CT, USA) at inoculation and every 25 min for 8 h.

2.7. E. coli O157:H7 Detection in Raw Ground Beef

In order to evaluate the ability of the purified Φ V10*nluc* phage to detect *E. coli* O157:H7 cells in raw ground beef, matrices assays were performed following standard and modified FSIS protocols. Approximately 1.73×10^3 pfu/mL final phage concentrations were used with ground beef in stomacher bags and Nalgene bottles, with a final volume of 1 L. This concentration was achieved by the addition of 100 µL of 1.73×10^7 pfu/mL phage per assay. All raw ground beef used in this study was purchased at a local grocery store and consisted of 83% lean meat, 17% fat. All ground beef was used immediately after purchase.

2.7.1. Detection of E. coli O157:H7 in Raw Ground Beef in Stomacher Bags

Samples of 325 g of ground beef were put in 975 mL of mTSB broth (1:3 sample to enrichment media) inside a sterile, plain, clear polypropylene stomacher bag without a filter mesh. *E. coli* O157:H7, ranging from approximately 3 to 3×10^4 CFU per bag, were inoculated into each ground beef slurry. Ground beef with mTSB broth and wild-type Φ V10 phage was used as the negative control. Each sample was hand-massaged for 30 s at the initial time of inoculation. Each assay was performed in triplicate. Luminescence measurements of 1 mL samples were taken after the addition of 10 µL of previously prepared Nano-Glo[®] reagent using a Sirius luminometer (Berthold Detection Systems, Bad Wildbad, Germany) once per hour.

2.7.2. Detection of E. coli O157:H7 in Raw Ground Beef in Nalgene Bottles

In order to determine the efficacy of the phage-based detection during shipping of ground beef samples, different ratios of beef sample to mTSB were evaluated in selected sample containers. I-chem Brand N311-1000 Nalgene 300 Series HDPE Wide Mouth Bottles (Thermo Fisher Waltham, MA, USA) were chosen due to size, low leak potential, and pressure requirements for shipping.

Initial experimentation included 325 g of raw ground beef added to 650 mL of mTSB for a 1:2 sample to media ratio. The same amount of raw ground beef was used with 325 mL of mTSB for a separate experiment at a 1:1 ratio. Each sample was inverted by hand for 30 s at the initial time of inoculation. All Nalgene bottles used were sterile. *E. coli* O157:H7 cell inocula were approximately 3 to 3×10^2 CFU per bottle in the 1:2 sample to media ratio and approximately 2 to 2×10^2 CFU per bottle in the 1:1 sample to media ratio.

3. Results and Discussion

3.1. Phage Purification

When the bacteria are infected by $\Phi V10nluc$, the cells may be lysed or become lysogens. When the lysogens are stressed by environmental factors, they may go into the lytic phase. They can spontaneously go lytic as well, or, during preparation, the addition of chloroform and sodium chloride can affect the lytic process. While intact and injured cells and cell debris are removed by centrifugation and filtration in the initial phage preparation, the phage as well as the NanoLuc[®] luciferase remain in solution, resulting in high levels of background light emission upon addition of the Nano-Glo® substrate. The presences of the NanoLuc[®] luciferase resulted in high background luminescence in *E. coli* O157:H7 detection assays. In an attempt to remove the NanoLuc[®] luciferase from the phage, three purified phage preparations were subjected to further purification using Amicon[®] Ultra-15 centrifugal filtration devices [28]. Removal of the NanoLuc[®] luciferase was determined by measuring the luminescence generated by the Nano-Glo® reagent in the filtrate after each centrifugation/filtration cycle. Reduced light emission in sequential rounds of filtration demonstrated the removal of the luciferase from the phage preparation (Table 1). The luciferase activity in the filtrate was reduced by approximately five orders of magnitude through the first six cycles of filtration, but plateaued to about two to four thousand RLUs after the seventh round of filtration (Table 1). The total background light emission in the phage retentate was also significantly reduced by roughly five orders of magnitude, from an average of 7.03×10^8 RLU/s to 9.33×10^3 RLU/s per 10 μ L sample assay (Table 2). The residual light generated after filtration appeared to indicate physical association of luciferase with the phage. Although this background light emission is not negligible, it is sufficiently low as to not interfere with E. coli O157:H7 detection assays and may serve as an indicator that the $\Phi V10nluc$ phage was indeed added to negative samples. It is worth noting that there was a consistent 10- to 100-fold reduction in phage concentration after the filtration steps. Plaque assays confirmed that a minimal amount of phage was found in the filtrate at the end of the purification procedure, suggesting that the phage was not being lost through the membrane.

Centrifugation Cycle	Replicates Filtrate Luminescence (RLU/s)		
	1	2	3
1	419,040,100	371,692,000	349,226,900
2	76,547,500	64,513,800	58,367,400
3	9,490,580	5,744,485	8,108,047
4	382,266	228,219	532,749
5	21,660	16,011	30,201
6	2954	3181	3045
7	3486	4083	2552

Table 1. Removal of NanoLuc[®] Luciferase from Φ V10*nluc* through seven rounds of centrifugal filtration. Luminescence (relative light units per second, RLU/s) was measured for 10 µL of Φ V10*nluc* filtrate after each consecutive centrifugation/filtration cycle for three separate phage preparations.

Table 2. Luminescence in Φ V10*nluc* phage preparations before and after purification by centrifugal filtration. Luminescence (RLU/s) was measured in 10 µL of Φ V10*nluc* phage preparation before and after purification using the Amicon[®] centrifuge filter to remove the NanoLuc[®] luciferase.

Phage Concentration (pfu/mL)	Luminescence Prior to Purification (RLU/s)	Luminescence After Purification (RLU/s)
7.20×10^{5} 1.62×10^{5} 1.62×10^{5}	$5.39 imes 10^{8}$ $7.08 imes 10^{8}$ $8.62 imes 10^{8}$	$6.08 imes 10^3 \ 8.08 imes 10^3 \ 1.38 imes 10^4$

3.2. E. coli O157:H7 Detection by Φ V10nluc Phage in LB

The ability of various concentrations of Φ V10*nluc* phage to detect *E. coli* O157:H7 cells was assessed in LB. These experiments were performed in order to test whether a higher concentration of $\Phi V10$ *nluc* phage would contribute to a shorter time to detection than previously shown by Zhang et al. [22]. $\Phi V10nluc$ phage concentrations ranging from 10^2 to 10^5 pfu/mL were assessed for time to detection of 0 to 10^5 *E. coli* O157:H7 CFU per assay. Results shown in Figure 1A indicate that 1.76×10^2 pfu/mL phage concentration was able to detect the presence of approximately 2 CFU in 40 mL of LB in the shortest amount of time—just over 6 h with no prior incubation. Additionally, the 10^3 pfu/mL phage concentration detected approximately 2 cells in under 10 h, while 10⁴ phage concentration was able to detect the same in roughly 8.5 h. Φ V10*nluc* phage concentration of 4.5×10^5 pfu/mL was able to detect approximately 200 cells in roughly 9 h (Figure 1C). The time to detection using both 4.33×10^3 pfu/mL and 4.33×10^4 pfu/mL phage concentrations decreased as cell numbers increased (Figure 1A–C). The phage concentration of 4.33×10^4 pfu/mL could detect approximately 2 cells in 8.5 h, 20 cells at 7 h, and 200 cells at 6 h. The phage of 4.33×10^3 pfu/mL detected approximately 2 cells in just under 10 h, while it was able to detect approximately 20 cells in 8.5 h, and approximately 200 cells in just under 7 h. Φ V10*nluc* of 1.73×10^3 pfu/mL concentration was used in subsequent experiments due to short time to detection and a low background luminescence. In addition, a higher phage concentration did not necessarily correlate with lower time to detection, exemplified by the inability of 10^5 pfu phage to detect either 2 or 20 cells within 12 h (Figure 1A,B).





3.3. Characterization of the Growth of E. coli O157:H7 and Φ V10nluc Lysogen

The growth of *E. coli* O157:H7 was compared with the Φ V10*nluc* lysogen growth in mTSB at 37 °C by measuring culture optical density. The wild-type *E. coli* O157:H7 culture has an approximate growth of 0.87 OD₆₀₀/h, while the Φ V10*nluc* lysogen has an estimated growth rate that is less than half of that of the wild-type, at 0.38 OD₆₀₀/h (Figure 2). After the initial lag of about 3.5 h, the optical density of *E. coli* O157:H7 cells doubled roughly every 50 min, while the Φ V10*nluc* lysogens showed a lag of about 6 h and an estimated doubling time of approximately 110 min. At room temperature (25 °C), the growth rate of the wild-type *E. coli* O157:H Φ V10*nluc* lysogen decreased to about 0.26 OD₆₀₀/h (doubling time of ~160 min) after a lag of approximately 20 h (Figure 3).



Figure 2. Growth of *E. coli* O157:H7 C7927 and C7927 Φ V10*nluc* lysogen at 37 °C.



Figure 3. Growth of *E. coli* O157:H7 C7927 ΦV10*nluc* lysogen at 37 °C and 25 °C.

3.4. E. coli O157:H7 Detection in Raw Ground Beef

The concentration of 1.73×10^3 pfu/mL of Φ V10*nluc* was used to evaluate time to detection of varying amounts of *E. coli* O157H7 C7927 in raw ground beef slurry. First, the FSIS protocol was followed using 325 g inoculated raw ground beef containing added phage in 975 mL of mTSB enrichment media within stomacher bags. Luminescence measurements were recorded hourly for 15 h and are shown in Figure 4. Increased light production correlated to increasing cell concentrations.



Figure 4. Time course of Luminescence detection of *E. coli* O157:H7 using the standard 1:3 ground beef to mTSB broth ratio in a stomacher bag enrichment format over time.

3.5. Growth of E. coli O157:H7 C7927

It is worth noting that the luminescence in the control samples not inoculated with *E. coli* O157:H7 gradually decreased over time. This gradual decrease was not observed in control experiments conducted in LB, suggesting this phenomenon may be due to proteases naturally present in the raw ground beef that may be degrading the NanoLuc[®] luciferase, leading to lower observed background luminescence. In addition, the source of variability in time to detection is most likely due to the increased enrichment volume (1300 mL) compared to the previous report by Zhang et al. [22], in which 40 mL ground beef slurries were used. For comparison, the cell concentration in the previously reported 40 mL assays [22] would be approximately equivalent to the 30 CFU inoculation shown in Figure 4.

3.6. Detection of E. coli O157:H7 in Raw Ground Beef in Nalgene Bottles

Due to the potential for leaks using an enrichment volume of 1300 mL, enrichment during shipment is not reasonably conducted in stomacher bags; therefore, the capability of the Φ V10*nluc* phage to detect *E. coli* O157:H7 in one-liter Nalgene bottles was investigated. One-liter bottles were chosen due to cost and shipment weight. Previous work by Cai and Cabezas ([29], unpublished) demonstrated positive results with ground meat samples which were simply sprayed with Φ V10*nluc*. This suggested that we could combine the use of the one-liter Nalgene bottles with lowered mTSB media volume (650 mL, 325 mL) with the ground meat sample (325 g), which would facilitate enrichment during transport (Figure 5).

Ground beef slurries (650 mL mTSB and 325 g ground beef) in Nalgene bottles containing approximately 0, 3, 30, and 300 CFU of *E. coli* O157:H7 were subjected to the phage-based luminescence assay. In all three experimental replicates, 300 CFU were detected at roughly 11 h (Figure 6). Results also showed that 30 CFU was detected in all three replicates in roughly 14 h. However, for three CFU, only the first replicate resulted in detection at about 15 h (Figure 6A). This fractional positive result at low inoculation levels is not surprising, given the likelihood that some samples may not have been inoculated with any cells.



Figure 5. Ground beef enrichment and phage enrichment during shipping. In the examples shown, 650 mL mTSB and 325 g ground beef were added to Nalgene bottles with 1.73×10^3 pfu/mL Φ V10*nluc* (final concentration) for simultaneous enrichment and detection.



Figure 6. Φ V10*nluc* infectivity in 650 mL mTSB Nalgene bottles. Replicates (A) 1, (B) 2, and (C) 3.

In this study, pressure was observed to have built up in the bottles during enrichment. This may be due to the shortage of dissolved and headspace oxygen initially available within the bottle, resulting in fermentative growth and gas production by *E. coli* O157:H7 or the beef microbiota. Though no leaks were observed in this study, this observation raises a potential safety concern related to sample integrity (i.e., bottle leakage) for enrichment during shipment.

An additional experiment was conducted using a 1:1 ratio of mTSB media to raw ground beef in the Nalgene bottles over 20 h in order to evaluate if a decrease in media would lead to higher available oxygen for the bacteria, potentially resulting in a shorter time to detection. Observations during this experiment included that there was still gas release upon opening the bottles after 8 h; however, there was less pressure and less gas released compared with the 2:1 ratio experiment. Results indicate that the Φ V10*nluc* phage detected approximately two cells in one out of three assays (Figure 7B). Figure 7 shows that a higher cell concentration correlated to higher measured luminescence over time. Similar to the 2:1 ratio Nalgene bottle experiment, 200 CFU was detected at roughly 11 h in all 3 replications while 20 CFU was detected at roughly 15 h. No luminescence above the control response was detected in any Nalgene bottle studies containing 0.2 to 0.3 CFU (Figures 6 and 7).



Figure 7. ΦV10*nluc* infectivity in 325 mL mTSB Nalgene bottles. Replicates (A) 1, (B) 2, and (C) 3.

4. Conclusions

Previous work has confirmed Φ V10*nluc* specificity to *E. coli* O157:H7 isolates [22]. This research showed that Φ V10*nluc* phage can detect roughly five *E. coli* O157:H7 cells in 40 mL raw ground beef slurry after approximately 9 h of enrichment/incubation. Such performance indicates the potential for exploiting this phage infection system as a detection platform coincident with ground beef regulatory test sample shipment. However, the

previous work conducted in 40 mL assays was not representative of current FSIS sampling methods. Therefore, this study evaluated the FSIS-recommended sample to media volume of a 325 g raw ground beef sample in 975 mL of mTSB enrichment media, and showed that Φ V10*nluc* was able to detect *E. coli* O157:H7, showing higher cell concentrations correlated with high luminescence in real-life application sample sizes. Within government regulatory protocols which are recommended to ensure food safety, the enrichment step is ubiquitous across all official and rapid methods used to detect foodborne pathogens. While culture enrichment can take up to 24–48 h after receipt of the sample at the regulatory lab, there is an opportunity to exploit this time during shipping for sample enrichment. Shipment of samples from meat production facilities to testing laboratories takes approximately 15-18 h. While it is known that FedEx Express Controlled Room Temperature holds samples at 15–25 °C during shipping, growth curves in this study show that this phage-based assay will not function optimally even at the upper limit of the cargo hold of 25 °C. Growth at 37 °C will be necessary to maximize the growth of STEC and, at least partially, to limit the growth of some of the foodborne microbiota. Competition studies are recommended to evaluate how background microbiota may affect the sensitivity of this detection method. In addition, the temperature recommendation of 37 °C for optimal performance should be addressed through further engineering studies. Potentially, a simple and inexpensive heat source, akin to a dry chemical or gel pack handwarmer or insulated boxes, may be assessed for maintaining the required temperature of samples during shipment.

Author Contributions: Conceptualization, B.M.A., A.G.G., and G.C.P.; methodology, B.M.A., C.C., and C.P.C.-A.; formal analysis, C.C.; investigation, C.C. and C.P.C.-A.; resources, B.M.A., A.G.G., and G.C.P.; data curation, B.M.A.; writing—original draft preparation, C.C.; writing—review and editing, B.M.A., A.K.B., A.G.G., and G.C.P.; supervision, B.M.A.; project administration, B.M.A.; funding acquisition, B.M.A. All authors have read and agreed to the published version of the manuscript.

Funding: This research was supported by the U.S. Department of Agriculture, Agricultural Research Service, Food Safety program in-house projects 8072-42000-093-00D (A.G.G.) and 8072-42000-094-00D (G.C.P.), and Agreement No. 59-8072-1-002 to the Purdue University Center for Food Safety Engineering (B.M.A.). This work was also supported by the USDA National Institute of Food and Agriculture, Hatch project number 1004830.

Data Availability Statement: The original contributions presented in the study are included herein.

Acknowledgments: Mention of trade names or commercial products in this article is solely for the purpose of providing specific information and does not imply recommendation or endorsement by the U.S.D.A. Opinions, findings conclusions, or recommendations expressed in this publication are those of the author(s) and do not necessarily reflect the view of the U.S.DA. U.S.D.A. is an equal opportunity provider and employer.

Conflicts of Interest: B.M.A. is the founder of Phicrobe an LLC located in West Lafayette Indiana, which has licensed technology related to using Φ V10 and its derivatives for the detection of *E. coli* O157:H7. All other authors declare no conflicts of interest.

References

- Scallan, E.; Hoekstra, R.M.; Angulo, F.J.; Tauxe, R.V.; Widdowson, M.; Roy, S.L.; Jones, J.L.; Griffin, P.M. Foodborne Illness Acquired in the United States—Major Pathogens. *Emerg. Infect. Dis.* 2011, *17*, 7–15. [CrossRef] [PubMed]
- Croxen, M.A.; Finlay, B.B. Molecular Mechanisms of *Escherichia coli* Pathogenicity. *Nat. Rev. Microbiol.* 2010, *8*, 26–38. [CrossRef] [PubMed]
- 3. Centers of Disease Control and Prevention. *E. coli* (*Escherichia coli*). U.S. Department of Health and Human Services. 2018. Available online: https://www.cdc.gov/ecoli/general/index.html (accessed on 15 September 2022).
- 4. Mele, C.; Remuzzi, G.; Noris, M. Hemolytic Uremic syndrome. Sem. Immunopathol. 2014, 36, 399–420. [CrossRef] [PubMed]
- EFSA. The European Union Summary Report on Trends and Sources of Zoonoses, Zoonotic Agents and Food-borne Outbreaks in 2012. EFSA J. 2014, 12, 3547. [CrossRef]
- Rangel, J.M.; Sparling, P.H.; Crowe, C.; Griffin, P.M.; Swerdlow, D.L. Epidemiology of *Escherichia coli* O157:H7 Outbreaks, United States, 1982–2002. *Emerg. Infect. Dis.* 2005, 11, 603–609. [CrossRef]
- 7. Karmali, M.A. Infection by Verocytotoxin-Producing Escherichia coli. Clin. Microbiol. Rev. 1989, 2, 15–38. [CrossRef] [PubMed]

- 8. Griffin, P.M.; Tauxe, R.V. The Epidemiology of Infections Caused by *Escherichia coli* O157:H7, Other Enterohemorrhagic *E. coli*, and the Associated Hemolytic Uremic Syndrome. *Epidemiol. Rev.* **1991**, *13*, 60–98. [CrossRef] [PubMed]
- 9. McGowan, K.L.; Wickersham, E.; Strockbine, N.A. Escherichia coli O157:H7 from water (Letter). Lancet 1989, 1, 967–968. [CrossRef]
- Swerdlow, D.L.; Woodruff, B.A.; Brady, R.C.; Griffin, P.M.; Tippen, S.; Donnell, H.D., Jr.; Geldreich, E.; Payne, B.J.; Neyer, A., Jr.; Wells, J.G.; et al. A Waterborne Outbreak in Missouri of *Escherichia coli* O157:H7 Associated with Bloody Diarrhea and Death. *Ann. Intern. Med.* 1992, 117, 812–819. [CrossRef] [PubMed]
- U.S. Department of Agriculture Food Safety and Inspection Service. Report on the Food Safety and Inspection Service's Microbiological and Residue Sampling Programs. 2011. Available online: https://www.fsis.usda.gov/wps/wcm/connect/0816 b926-c7ee-4c24-9222-34ac674ec047/FSIS_Sampling_Programs_Report.pdf?MOD=AJPERES (accessed on 15 September 2022).
- 12. Paoli, G.C.; Wijey, C.; Uhlich, G.A. Genetically Marked Strains of Shiga Toxin-Producing O157:H7 and Non-O157 *Escherichia coli*: Tools for Detection and Modeling. *J. Food Prot.* **2015**, *78*, 888–901. [CrossRef] [PubMed]
- Cloke, J.; Crowley, E.; Bird, P.; Bastin, B.; Flanngery, J.; Agin, J.; Goins, D.; Clark, D., Jr.; Radcliff, R.; Wickstrand, N.; et al. Validation of the Thermo Scientific SureTect *Escherichia coli* O157:H7 Real-Time PCR Assay for Raw Beef and Produce Matrixes. *J.* AOAC Int. 2015, 98, 1301–1314. [CrossRef] [PubMed]
- Feng, P.; Weagant, S.D.; Jinneman, K. Diarrheagenic *Escherichia coli*. Bacteriological Analytical Manual Chapter 4A. U.S. Food & Drug Administration. 2020. Available online: https://www.fda.gov/food/laboratory-methods-food/bam-chapter-4adiarrheagenic-escherichia-coli (accessed on 15 September 2022).
- Panwar, S.; Duggirala, K.S.; Yadav, P.; Debnath, N.; Yadav, A.K.; Kumar, A. Advanced Diagnostic Methods for Identification of Bacterial Foodborne Pathogens: Contemporary and Upcoming Challenges. *Crit. Rev. Biotechnol.* 2022. *advanced online publication*. [CrossRef] [PubMed]
- 16. Rohde, A.; Hammerl, J.A.; Boone, I.; Jansen, W.; Fohler, S.; Klein, G.; Diekmann, R.; Al Dahouk, S. Overview of Validated Alternative Methods for the Detection of Foodborne Bacterial Pathogens. *Trends Food Sci. Technol.* **2017**, *62*, 113–118. [CrossRef]
- U.S. Department of Agriculture-Food Safety and Inspection Service. MLG 5C.02. Detection, Isolation, and Identification of Top Seven Shiga Toxin-Producing *Escherichia coli* (STEC) from Meat Products and Carcass and Environmental Sponges. 2021. Available online: https://www.fsis.usda.gov/sites/default/files/media_file/2021-08/MLG-5C.02.pdf (accessed on 15 September 2022).
- U.S. Department of Agriculture-Food Safety and Inspection Service. MLG Appendix 1.10. Media and Reagents. 2022. Available online: https://www.fsis.usda.gov/sites/default/files/media_file/2022-03/MLG_Appendix_1.10.pdf (accessed on 15 September 2022).
- U.S. Department of Agriculture-Food Safety and Inspection Service. Use of FSIS Regulatory Verification Sampling to Generate Prevalence Estimates. DCC Prevalence Estimate Workgroup. 2012. Available online: https://www.fsis.usda.gov/sites/default/ files/media_file/2020-09/Prevalence_Estimates_Report.pdf (accessed on 15 September 2022).
- U.S. Department of Agriculture-Food Safety Inspection Service. Sampling Results for FSIS Regulated Products. 2022. Available online: http://www.fsis.usda.gov/sites/default/files/media_file/documents/Sampling_Project_Results_Data_20210701_202 20630.pdf (accessed on 15 September 2022).
- U.S. Department of Agriculture-Food Safety and Inspection Service. Raw Beef Product Sampling. Inspection Methods. 2021. Available online: https://www.fsis.usda.gov/sites/default/files/media_file/2021-11/18_IM_Raw-Beef-Product-Sampling-11 032021.pdf (accessed on 15 September 2022).
- Zhang, D.; Coronel-Aguilera, C.P.; Romero, P.L.; Perry, L.; Minocha, U.; Rosenfield, C.; Gehring, A.; Paoli, G.C.; Bhunia, A.K.; Applegate, B. The Use of a Novel NanoLuc-Based Reporter Phage for the Detection of *Escherichia coli* O157:H7. *Sci. Rep.* 2016, 6, 33235. [CrossRef] [PubMed]
- Hall, M.P.; Unch, J.; Binkowski, B.F.; Valley, M.P.; Butler, B.L.; Wood, M.G.; Otto, P.; Zimmerman, K.; Vidugiris, G.; Machleidt, T.; et al. Engineered Luciferease Reporter from a Deep Sea Shrimp Utilizing a Novel Imidazopyrazinone Substrate. ACS Chem. Biol. 2012, 7, 1848–1857. [CrossRef] [PubMed]
- Khakhria, R.; Duck, D.; Lior, H. Extended Phage-Typing Scheme for Escherichia coli O157:H7. Epidemiol. Infect. 1990, 105, 511–520. [CrossRef] [PubMed]
- Perry, L.L.; SanMiguel, P.; Minocha, U.; Terekhov, A.I.; Shroyer, M.L.; Farris, L.A.; Bright, N.; Reuhs, B.L.; Applegate, B.M. Sequence Analysis of *Escherichia coli* O157:H7 Bacteriophage ΦV10 and Identification of a Phage-encoded Immunity Protein that Modifies the O157 Antigen. *FEMS Microbiol. Let.* 2009, 292, 182–186. [CrossRef] [PubMed]
- Stewart, G.S.A.; Williams, P. Lux Genes and the Applications of Bacterial Bioluminescence. J. Gen. Microbiol. 1992, 138, 1289–1300. [CrossRef] [PubMed]
- 27. Zhao, T.; Doyle, M.; Besser, R. Fate of Enterohemorrhagic *Escherichia coli* O157:H7 in Apple Cider with and without Preservatives. *Appl. Environ. Microbiol.* **1993**, *59*, 2526–2530. [CrossRef] [PubMed]
- Bonilla, N.; Rojas, M.I.; Cruz, G.N.F.; Hung, S.; Rohwer, F.; Barr, J.J. Phage on Tap—A Quick and Efficient Protocol for the Preparation of Bacteriophage Laboratory Stocks. *PeerJ* 2016, *4*, e2261. [CrossRef] [PubMed]
- Cai, S.; Cabezas, E. Bacteriophage Infection in Ground Beef. In Undergraduate Research Symposium; Purdue University: West Lafayette, IN, USA, 2016.