

University of Nevada, Reno

**Bacteriophage solutions as processing aids in beef production: An essential hurdle to  
reduce *E. coli* O157:H7 and the “Big Six” serotypes.**

A thesis submitted in partial fulfillment of the requirements for the  
degree of Master of Science in Animal and Rangeland Science

by

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THE GRADUATE SCHOOL

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## Abstract

The research presented herein demonstrates the applications of bacteriophage solutions against pathogenic *E. coli* on fresh beef. The first chapter demonstrates the effects of a novel seven-bacteriophage (Mello-Shebs 1 phages) cocktail application against the seven USDA adulterant Shiga Toxin-producing *E. coli* (STEC) O157:H7 and the “Big Six” in vitro and on beef trim before grinding. The in vitro killing efficiency of each phage against the strain used for isolation ranged from 96.2% to 99.9%. On beef, inoculated and treated samples were held for 30 min or 6 h and at 7°C or 25°C before grinding to simulate refrigeration and ambient air temperatures. Both phage application ( $P \leq 0.0001$ ) and temperature ( $P = 0.0451$ ) had significant effects on STEC reduction. After 30 minutes of lysing time, phage application led to a significant 0.76 Log at 7°C and 0.81 Log reduction at 25°C. This study shows that the application of a seven-bacteriophage cocktail that target STEC O157:H7 and ‘Big Six’ serotypes are effective in reducing STEC on ground beef.

The research presented in chapter two demonstrates the effects of a commercially available bacteriophage cocktail compared to lactic acid and peroxyacetic acid in reducing *E. coli* O157:H7 on fresh beef surface when kept under vacuum or aerobic conditions simulating a High Event Period, a period of high *E. coli* rates in beef trim. During which, beef subprimals are typically removed from vacuum and treated with antimicrobials, then re-packaged and later tested to verify the absence of STEC. Bacterial loads were evaluated 30 min and 6 h after antimicrobial applications. Under

aerobic conditions for 30 min and 6 h, phage reduced STEC in beef by approximately 1.4 Log CFU/cm<sup>2</sup> whereas organic acids resulted in approximately 0.5 Log CFU/cm<sup>2</sup> reduction. Under vacuum conditions for 30 min, bacteriophage significantly reduced STEC by 1 Log CFU/cm<sup>2</sup>, while no significant effects were observed from organic acid treatment. Under vacuum conditions for 6 h, bacteriophage reduced STEC loads by 1.4 Log CFU/cm<sup>2</sup>, lactic acid reduced by 0.6 Log CFU/cm<sup>2</sup>, and no significant effects were observed with peroxyacetic acid application.

The research presented in chapter three demonstrates the effects of multiple antimicrobial treatments against STEC O157:H7 and the “Big Six” on fresh beef surface: a novel seven-bacteriophage cocktail (P), peroxyacetic acid (PAA), ultraviolet light (UV), and acidified chlorite (ASC) each alone and in combinations of two under simulated High Event Period scenarios. All treatment combinations that included phage significantly decreased STEC populations ( $P < 0.0001$ ). Under vacuum conditions, Phage + PAA resulted in the lowest STEC load (1.49 Log CFU/cm<sup>2</sup> reduction), followed by Phage, Phage + UV, Phage + ASC, Phage + PAA and UV + ASC, respectively. Under aerobic conditions, P + UV provided the greatest reduction (1.46 Log CFU/cm<sup>2</sup>), followed by Phage + PAA, P, Phage + ASC, UV+ASC, and PAA, respectively. When analyzing contrasts, treatments with phage significantly decreased STEC loads when compared to other treatments ( $P < 0.0001$ ) and control ( $P < 0.0001$ ). Overall, bacteriophage application has been demonstrated to have a high efficacy in reducing STEC loads on beef alone or with other antimicrobial treatments during a simulated industry application.

Overall, bacteriophage applications targeting STEC provided an efficient control against all adulterant serotypes. It is clear that organic acids commonly used during meat processing are not efficient in controlling STEC. Bacteriophages emerged as an effective antimicrobial that can be applied in different processing steps and during different production scenarios. Currently, according to the United States Department of Agriculture (USDA), bacteriophage solutions allowed to be used in meat processing have Generally Recognized as Safe (GRAS) status only against O157:H7. This research showed that phages can also lower the “Big Six” serotypes. Therefore, when producing phage cocktails that successfully target the “Big Six”, biotechnology companies must request to the Food and Drug Administration (FDA) GRAS status for solutions not only targeting STEC O157:H7, but also the O145, O121, O111, O103, O45, and O26 serotypes.

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Table of Contents	
<b>Abstract</b> .....	i
<b>Acknowledgements</b> .....	iv
Table of Contents .....	v
<b>List of Tables</b> .....	vii
<b>List of Figures</b> .....	viii
<b>List of Appendices</b> .....	ix
<b>Introduction</b> .....	1
<b>Review of Literature</b> .....	3
<b>Foodborne diseases</b> .....	3
<i>Escherichia coli</i> .....	4
<b>Routes of contamination</b> .....	7
<b>Food Safety interventions</b> .....	9
<b>Organic Acids</b> .....	11
<b>Acid Resistance</b> .....	12
<b>Ultraviolet Light</b> .....	14
<b>Bacteriophage (phage)</b> .....	14
<b>Mechanism of Infection</b> .....	16
<b>Phage Therapy as Biocontrol in Food</b> .....	18
<b>Conclusion</b> .....	19
<b>References</b> .....	21
<b>CHAPTER ONE</b> .....	36
<b>Abstract</b> .....	37
<b>Introduction</b> .....	38
<b>Materials and methods</b> .....	40
<b>Results</b> .....	43
<b>Discussion</b> .....	44
<b>Conclusion</b> .....	49
<b>References</b> .....	50
<b>CHAPTER TWO</b> .....	63
<b>Abstract</b> .....	64
<b>Introduction</b> .....	65

<b>Materials and Methods</b> .....	67
<b>Results</b> .....	72
<b>Discussion</b> .....	73
<b>Conclusion</b> .....	79
<b>References</b> .....	80
<b>CHAPTER THREE</b> .....	88
<b>Abstract</b> .....	89
<b>Introduction</b> .....	90
<b>Materials and methods</b> .....	92
<b>Results</b> .....	97
<b>Discussion</b> .....	99
<b>Conclusion</b> .....	110
<b>References</b> .....	112

## List of Tables

### Chapter 1

Table 1. Lysing efficiency<sup>1</sup> of individual bacteriophages MS1-157:H7, MS1-145, MS1-121, MS1-111, MS1-103, MS1-45, and MS1-26 against *E. coli* O157:H7 (ATCC® 35150™), O26 (ATCC® BAA-2196™), O45 (ATCC® BAA-2193™), O103 (ATCC® BAA-2215™), O111 (ATCC® BAA-2440™), O121 (ATCC® BAA-2219™), and O145 (ATCC® BAA-2192™) in vitro.

Table 2. Least square means of Shiga Toxin *E. coli* contamination level (Log<sub>10</sub> CFU/g) strains in ground beef following application of control (BPW) or bacteriophage cocktail preparation (MS1-157:H7, MS1-145, MS1-121, MS1-111, MS1-103, MS1-45, and MS1-26) at two holding times 30 minutes and 6 hours at 7° C or 25° C on beef trim prior to grinding.

### Chapter 2

Table 1. Killing efficacy of PhageGuard E™ (Bacteriophages EP75 and EP335) for four *E. coli* O157:H7 strains.

Table 2. Effects of the application of organic acids and bacteriophage solutions under vacuum (V) and aerobic conditions (non-vacuum, NV) on beef contaminated with 4 *E. coli* strains (ATCC® 35,150™, ATCC® 43,895™, ATCC® 43,894™, and NCTC 13128). Log CFU/cm<sup>2</sup>.

### Chapter 3

Table 1. Significance of contrasts analysis for treatments applied under vacuum and aerobic conditions.

Table 2. Significance of contrasts analysis for treatments applied under vacuum and aerobic conditions.

## List of Figures

### Chapter 1

Figure 1. Least Square Means of Shiga Toxin *E. coli* populations (Log CFU/g) in ground beef following control and phage applications on beef trim with fixed effects of time (30 min, 6 h) and temperature (7° C, 25° C). Std error = 0.2477,  $P < 0.0001$ . Contrasts: Phage vs Control,  $P < 0.0001$ ; 7°C vs 25°C,  $P = 0.0497$ ; and 30 min vs 6 h,  $P = 0.2797$ .

### Chapter 3

Figure 1. Least square means of Shiga Toxin *E. coli* populations in beef samples (*m. Cutaneous trunci*) kept under vacuum and treated with BPW (C), Peroxyacetic acid (PAA, 400 ppm), Acidified Sodium Chlorite (ASC, 1200 ppm), Ultraviolet light - C (UV, 254 nm, 23°C for 30 s), MS1 Bacteriophages (P, 108 PFU/ml), PAA+ASC, PAA+UV, ASC+UV, P+PAA, P+ASC, and P+UV. Standard error of the mean = 0.2108.

Figure 2. Least square means of Shiga Toxin *E. coli* populations in beef samples (*m. Cutaneous trunci*) kept under aerobic conditions and treated with BPW (C), Peroxyacetic acid (PAA, 400 ppm), Acidified Sodium Chlorite (ASC, 1200 ppm), Ultraviolet light - C (UV, 254 nm, 23°C for 30 s), MS1 Bacteriophages (P, 108 PFU/ml), PAA+ASC, PAA+UV, ASC+UV, P+PAA, P+ASC, and P+UV. Standard error of the mean = 0.1919.

**List of Appendices**

Appendix I. Safe Minimum Internal Temperature Chart (USDA,2015)

Appendix II. Getting Started with an ATCC Bacterial Strain (ATCC, 2015)

Appendix III. Bacteriophage isolation from sewage

Appendix IV. Double-layer Agar Assay. Adams (1959), modified by Yeh et al. (2017)

Appendix V. Bacteriophage Amplification

Appendix VI. Measuring Optical Density ( $OD_{600}$ ) and diluting inoculum

## Introduction

Foodborne illness will affect 1 in 6 people in the United States every year, despite the numerous antimicrobial interventions used to reduce contamination of food products, particularly those of animal origin. In the United States, ground beef is the most widely consumed beef product at an estimated 42% of all beef cuts (Davis & Lin, 2005). Ground beef is also known by its associations with foodborne diseases caused by Shiga Toxin-producing *E. coli* (STEC) and *Salmonella* (Scallan et al., 2011). These pathogens are found within the intestines of animal hosts as well as within their environment. There is a high inherent risk of cross-contamination from animals to their raw food products during processing, requiring the need for food producers to have robust and effective food safety systems to ensure they produce safe food for the public.

*Escherichia coli* is known to be present on cattle hides before the slaughter process, unfortunately hide removal can create opportunity for cross-contamination from hide to carcass (Arthur et al., 2007). This is likely a main mechanism of transfer of enteric organisms, such as *E. coli*, from the gastrointestinal tract to fresh beef products. The contamination of carcasses can produce contaminated whole cuts (primals, subprimals, steaks) and contaminated trim. Trim from hundreds of carcasses can make up one lot of trim, with many lots combined for ground beef. The USDA does mandate sampling of Shiga Toxin-producing *E. coli* on beef carcasses and in beef trim, but

even the best sampling plans do not ensure identification or complete absence of pathogens.

In order to maintain the safety of our food there needs to be continuous efforts in improving antimicrobial food safety interventions. The inseparability of microorganisms from food animals and the environment will always pose a risk for contaminating final food products. From upstream preventative measures to direct sanitizing applications, antimicrobial interventions are extensively used in order to combat the presence of foodborne bacteria, viruses, and parasites that cause food poisoning (Marriott, Schilling, & Gravani, 2018). The recent acceptance of bacterial viruses (bacteriophages) as a natural method for food biocontrol has continued to show promise for their efficacy and versatility and could be the next advancement for food safety systems.

The continuous advancement of food safety interventions is essential to decrease and eliminate foodborne pathogens and to mitigate the burden of foodborne illnesses. The objectives within the presented research are to 1) Evaluate the efficacy of reducing all seven STEC serotypes in ground beef with a cocktail of seven environmentally isolated bacteriophages, 2) Evaluate the efficacy of a commercial bacteriophage preparation on beef surface under a High Event Period scenario compared to other commonly used antimicrobial solutions, and 3) Determine the optimal treatment of STEC during a simulated High Event Period between organic acids PAA and ASC, UV light, and a 7-bacteriophage cocktail alone and in combinations.

## Review of Literature

### Foodborne diseases

Foodborne illness affects 1 in 6 people in the United States every year and remains a major cause of hospitalization and death worldwide despite many advances in food sanitation techniques and pathogen surveillance (Moye, Woolston, & Sulakvelidze, 2018). The Centers for Disease Control and Prevention (CDC) estimates that every year 48 million people become ill, 128,000 are hospitalized, and 3,000 die from foodborne diseases in the United States (CDC, 2020a). There are 31 major foodborne pathogens that are acquired in the United States, the top five illness-causing microorganisms are norovirus (58%), followed by nontyphoidal *Salmonella* spp. (11%), *Clostridium perfringens* (10%), *Campylobacter* spp. (9%), and *Staphylococcus aureus* (3%). The five leading causes of hospitalization were nontyphoidal *Salmonella* spp. (35%), norovirus (26%), *Campylobacter* spp. (15%), *Toxoplasma gondii* (8%), and *E. coli* O157:H7 (4%). The five leading causes of death were nontyphoidal *Salmonella* spp. (28%), *T. gondii* (24%), *Listeria monocytogenes* (19%), norovirus (11%), and *Campylobacter* spp. (6%) (Scallan et al., 2011). In 2013, foodborne illnesses imposed over \$15.5 billion of economic burden onto the public, with 90% of the burden caused by only five pathogens. *Salmonella* nontyphoidal spp. poses the greatest economic burden at an estimated \$3.66 Billion, roughly 25% of the total burden by foodborne illnesses (Hoffmann, Macculloch, & Batz, 2015).

### ***Escherichia coli***

*Escherichia coli* are motile, gram negative, rod shaped bacteria belonging to the *Enterobacteriaceae* family. They are facultative anaerobes that make up a part of our gut microbiota as commensal bacterium living symbiotically within mammals such as humans, cattle, and deer. Most strains of *E. coli* are harmless; however, many strains have acquired DNA that code for virulence factors such as enterotoxins, invasins, or antibiotic resistance genes that allow them to cause severe gastrointestinal, urinary, or meningeal infections (Nataro & Kaper, 1998). Acquisition of new DNA is by horizontal gene transfer from extracellular DNA (transformation), other bacteria (conjugation), or bacteriophages (transduction). There are six well-known *E. coli* pathotypes that have acquired virulence factors and cause their own diseases: enteropathogenic *E. coli* (EPEC), enterotoxigenic *E. coli* (ETEC), enteroinvasive *E. coli* (EIEC), enterohaemorrhagic *E. coli* (EHEC), enteroaggregative *E. coli* (EAEC) and diffusely adherent *E. coli* (DAEC) (Nataro & Kaper, 1998).

The EHEC pathotype belong to the same group as verotoxigenic *E. coli* (VTEC) also known as Shiga toxin-producing *E. coli* (STEC), which are characterized by adhering to the intestinal walls of the colon and distal ileum (Nataro & Kaper, 1998; Phillips et al., 2000). Virulence factors found in this pathotype include the locus of enterocyte effacement, shiga toxins, and the pO157 plasmid, among others (Nataro & Kaper, 1998). The locus of enterocyte effacement (LEE) pathogenicity island in the chromosomal DNA (cDNA) codes for a type III secretion system (T3SS), secretory effector proteins, intimin, and Translocated intimin receptor (Tir) that are partially responsible for the attaching-

and-effacing (A/E) histopathology (Jarvis & Kaper, 1996; Mcdaniel, Jarvis, Donnenberg, & Kaper, 1995; Perna et al., 1998). Intimin is an adhesin encoded by the *eae* gene whose product is expressed on the bacterial cell surface. Intimin binds strongly to Tir (Translocated intimin receptor) which is inserted into the host enterocyte membrane by the bacterial T3SS, which also translocates many other bacterial proteins into the cytoplasm (Stevens, Galyov, & Stevens, 2006). The Shiga toxin produced by STEC is an AB<sub>5</sub> toxin coded by either *stx1* or *stx2* genes in the cDNA. The *stx1* is identical to the *Shigella dysenteriae* I gene (Nakao & Takeda, 2000; O'Brien et al., 1992) and *stx1* and *stx2* have 55-58% sequence identity (Jackson, Neill, O'Brien, Holmes, & Newland, 1987), while *stx2* has been shown to be 400 times more toxic (Siegler et al., 2003; Tesh et al., 1993). The secreted toxin is endocytosed into the host cell where it is able to cleave the 60S ribosomal subunit, inhibiting protein synthesis. All clinical isolates of *E. coli* O157:H7 possess a pO157 plasmid while most non-O157 EHEC isolates possess a similar plasmid (Hales, Hart, Batt, & Saunders, 1992; Lim, Yoon, & Hovde, 2010). The pO157 plasmid is a highly conserved F-like plasmid with many insertion sequences indicating that the genetic fragments are likely from many different origins (Lim et al., 2010). Some virulence factors contained within pO157 include hemolysins, type II secretion system, adhesins, and many functional enzymes (Brunner, Schmidt, & Karch, 1996, 1997; Lathem et al., 2002; Schmidt, Henkel, & Karch, 1997; Schmidt, Karch, & Beutin, 1994; Tatsuno et al., 2001). Such genes

could influence pathogenesis of the bacterial hosts by increasing abilities of adherence, colonization, and acid resistance (Lim et al., 2010).

Strains of *E. coli* within a pathotype are differentiated by serotyping, which evaluates lipopolysaccharides (LPS), a major component of cell wall structures. Serotyping identifies the unique combination of LPS structure, position, stereochemistry, and linkage of this highly variable region of the cell membrane (Liu et al., 2010; Stenutz, Weintraub, & Widmalm, 2006). The three components used for serotyping are the LPS O-antigen, flagellar H-antigen, and the capsule K-antigen. There are 183 identified O groups and 53 H groups (Joensen, Tetzschner, Iguchi, Aarestrup, & Scheutz, 2015) with O157:H7 being the most notorious. Other serotypes within the EHEC pathotype include O20, O25, O26, O44, O91, O111, O114, O126, O127, O128, O142, and O158 (Nataro & Kaper, 1998). Other notorious serotypes are those known to cause foodborne illness include STEC O145, O121, O111, O104:H4, O103, O45, and O26 whose infections can range from very mild to life-threatening (CDC, 2018). In 2014, there were 4,437 culture confirmed STEC infections (43.2% due to O157, 56.5% due to non-O157) and over \$298 million in medical care costs in the US (CDC, 2017; USDA-ERS, 2014). Common symptoms of STEC infections include diarrhea, most often bloody, abdominal cramps, and vomiting with the possible development of Hemolytic Uremic Syndrome (HUS), a type of kidney failure that develops in about 5-10% of STEC O157 infections that can be life-long or fatal (CDC, 2016). The infectious dose (ID) of STEC is very low with estimates of fewer than 700 cells to 10-100 cells ( $ID_{50} = < 100$  cells) that

can cause illness (Armstrong, Hollingsworth, & Morris, 1996; FDA, 2012; Griffin & Tauxe, 1991; Tuttle et al., 1999).

In 1994, the USDA deemed *E. coli* O157:H7 an adulterant in ground beef after a multistate outbreak from undercooked hamburger patties served at a national fast food chain that resulted in over 500 illnesses and four deaths (CDC, 1993). Years later in 2012, six more serotypes referred to as the “Bix Six” were declared adulterants by the USDA after causing 64% of the STEC foodborne illnesses (USDA-FSIS, 2012b). From 2010 to 2018, there were 8,802 STEC infections caused by O157 (48.6%), O26 (17.8%), O103 (15.2%), O111 (10.5%), O121 (3.9%), O145 (2.5%), and O45 (1.5%) (CDC, 2020b). The implementation of the USDA’s “Zero Tolerance” policy caused many food safety interventions to be established to eliminate the presence of foodborne *E. coli* in beef products before being sold to consumers (USDA-FSIS, 2012a). The USDA regulates sampling frequencies for the seven adulterant STEC serogroups and mandates that no STEC positive products are sold into commerce but are instead able to be sold to a cooker for further processing at a reduced profit.

### **Routes of contamination**

It is known that *E. coli* can colonize the colorectal junction of cattle where it can be excreted in manure potentially contaminating the environment, their hides, and the hides of other cattle (Burnens et al., 1995; Callaway et al., 2008; Clarke, 1994; Griffin & Tauxe, 1991; Hancock et al., 1994; Wells et al., 1991). The

presence of STEC on cattle hides has been shown to be a major source of bacterial contamination of beef carcasses due to direct or cross contamination during harvesting and processing, particularly at dehiding and evisceration steps (Arthur et al., 2007, 2008, 2004, 2010; Bacon et al., 2000; Bell, 1997; Brichta-Harhay et al., 2008; Byrne, Bolton, Sheridan, McDowell, & Blair, 2000; McEvoy et al., 2000; Nou et al., 2003; Rhoades, Duffy, & Koutsoumanis, 2009). Evaluating the prevalence of *E. coli* within a beef harvesting facility, Barkocy-Gallagher et al. (2003) recovered *E. coli* O157:H7 from 5.9% of fecal samples, 60.6% of hide samples, and 26.7% of carcasses sampled before the pre-evisceration wash and from 1.2% of post-intervention carcasses at chilling at low levels of <math><3.0</math> cells per 100 cm<sup>2</sup>. Slaughter facilities must implement extra preventative measures and increase sampling during warmer months (spring to mid-autumn) which have shown to have a higher incidence of *E. coli* (Barkocy-Gallagher et al., 2001; USDA-FSIS, 2014, 2017). Barkocy-Gallagher et al. (2003) also observed that non-O157 STEC *stx*-positive isolates were detected by PCR in fecal (34.3%), hides (92.0%), pre-evisceration carcasses (96.6%), and post-evisceration carcasses (16.2%) with about 4% of post-intervention carcasses having  $\geq 3.0$  cells per 100 cm<sup>2</sup> of non-O157 STEC and *stx*-harboring cells. Bacon et al. (2000) also demonstrated that the cattle hides can have approximately 5-7 Log<sub>10</sub> CFU/100 cm<sup>2</sup> of *E. coli* prior to the slaughter process, with hide removal resulting in 3-5 Log<sub>10</sub> CFU/100 cm<sup>2</sup> on the carcass, and 0.9 Log<sub>10</sub> CFU/100 cm<sup>2</sup> on the post-chilling carcass. Arthur et al. (2004) observed a positive relationship between contamination level of hides and pre-evisceration carcass. Not only are individual carcasses at risk for contamination, but contamination can also be

disseminated throughout fabrication lines, food contact surfaces, equipment, and other beef products by cross contamination during poor harvesting and dressing procedures (Yang, He, Badoni, Tran, & Wang, 2017). As a result, it was found that the most prevalent serogroups in beef trimmings at a midwestern plant were: O121(17.9%), O45 (8.6%), and O111 (8.6%), while beef trimmings from another plant were positive for O103 (31%), O121 (24.6%), and O45 (21.8%) (Kanankege et al., 2017). The Zero Tolerance policy mandates that no trace of the adulterant serogroups should be detected in fresh beef, however, a realistic and feasible sampling plan cannot ensure the complete absence of a specific organism from all products (Huffman, 2006; ICMSF, 2002).

### **Food Safety interventions**

The implementation of food safety regulations and interventions by industry and regulatory agencies has shown to reduce the overall number of retail food associated illnesses (Pollari et al., 2017). All antimicrobial interventions applied onto meat and poultry products are reviewed and approved by the Food Safety Inspection Service (FSIS) and listed in FSIS Directive 7120.1 (USDA-FSIS, 2019). There are many interventions that are effective in reducing microorganisms that can be implemented pre-harvest, during lairage, during harvest, and post-harvest (Wheeler, Kalchayanand, & Bosilevac, 2014). The best adopted pre-harvest interventions are probiotic feed supplementation and application of phages onto cattle hides, with vaccines against *E. coli* O157

also being well researched (Callaway, 2010; Dean-Nystrom, Gansheroff, Mills, Moon, & O'Brien, 2002; Fingermann et al., 2018; Judge, Mason, & O'Brien, 2004; Rabinovitz et al., 2012; Smith, 2014; Snedeker, Campbell, & Sargeant, 2012; Wheeler et al., 2014; Wolf et al., 2012). In lairage, phage products for hide sprays applied prior to entering the slaughter facility have become commercially available (Brashears & Chaves, 2017). During harvest, a multi-hurdle approach combining washing and sanitizing solutions has been effective in reducing bacterial populations on carcasses as long as the bacterial load does not exceed the capacity of the interventions (Koochmaraie et al., 2007). The most common physical interventions during harvest include steam-vacuuming (hocks and initial hide cuts), knife trimming (visible contamination), and water rinsing (Dorsa, Cutter, & Siragusa, 1996, 1997). Chemical interventions used during and after harvest mainly consist of organic acids with acetic, citric, and lactic acids being some of the most common agents (Belk, 2001). Other antimicrobial compounds used during or after harvest include oxidizing agents such as peroxyacetic acid (PAA), acidified sodium chlorite (ASC), ozone, or hypobromous acid (Wheeler et al., 2014).

Thermal decontamination methods are commonly used and have shown to be very effective (Olson & Nottingham, 1980). Steam vacuuming, hot, pressurized water sprays, and steam pasteurization are very common thermal interventions with nearly all large U.S. beef processing plants have automated hot water wash cabinets for pre-evisceration and final carcass washes (Wheeler et al., 2014). Spraying hot water (85 °C) for 15 s at 15 psi has shown to reduce all seven STEC serotypes between 3.2 and 4.2 Log CFU/cm<sup>2</sup> on inoculated beef surfaces (Kalchayanand et al., 2012). Even though thermal

treatments can cause the meat to have a cooked or whitened appearance, but the discoloration lessens after chilling (Castillo, Hardin, Acuff, & Dickson, 2002). Non-thermal interventions do not affect quality attributes such as color, making them a good alternative to thermal interventions and chemical interventions. The beef industry uses electron beams (e-beams), cold atmospheric plasma, UV light, UV and ozone combination, and high-pressure processing (Wheeler et al., 2014). Electron beams utilize streams of electrons to damage microbial DNA, inhibiting replication (Silindir & Özer, 2009). UV light, like e beams, damages microbial DNA causing replication inhibition. Cold atmospheric plasma is a weak ionized gas that generates reactive molecules like atoms, ions, and radicals that damage the cell wall (Laroussi, 2005; Stoffels, Sakiyama, & Graves, 2008). Lastly, high pressure processing is able to decontaminate food throughout instead of only surface decontamination while in packaging, however, it does cause discoloration (Gross & Jaenicke, 1994; Wheeler et al., 2014). No single intervention results in complete sterilization, making a multi-hurdle approach necessary to produce safe food and to maximize reduction of foodborne microorganisms (Wheeler et al., 2014).

### **Organic Acids**

Weak acidic solutions such as lactic acid or acetic acid diffuse across the bacterial cell wall and dissociate in the cytoplasm, disrupting the chemiosmotic functions of the cell membrane, and the transmembrane proton gradient

causing the cytoplasmic pH to decrease (Baldry & Fraser, 1988; Leaper, 1984; Liberti & Notarnicola, 1999). The efficacy of these chemical interventions has been demonstrated with high reductions (Harris, Miller, Loneragan, & Brashears, 2006; Laury et al., 2009; Zhao, Zhao, & Doyle, 2009), but lower or inconsistent efficacy has also been observed (Bosilevac, Nou, Barkocy-Gallagher, Arthur, & Koohmaraie, 2006; King et al., 2005; Yeh, de Moura, Van Den Broek, & de Mello, 2018). These interventions are advantageous due to their cost, availability, and simple application (Mataragas, Skandamis, & Drosinos, 2008). However, their potential decrease in efficacy has been demonstrated with the development of acid resistant bacterial pathogens when exposed to sub-lethal applications (Bearson, Bearson, & Foster, 1997; Brown, Ross, McMeekin, & Nichols, 1997; Capita, Riesco-Peláez, Alonso-Hernando, & Alonso-Calleja, 2014; Cheng, Yu, & Chou, 2003; Foster & Hall, 1990; Goodson & Rowbury, 1989; Leyer, Wang, & Johnson, 1995; Wesche, Gurtler, Marks, & Ryser, 2009).

### **Acid Resistance**

*E. coli* O157:H7 outbreaks associated with acidic foods such as apple cider and fermented sausage have brought attention to the ability of *E. coli* to survive in foods previously believed to be preserved by low pH (Leyer et al., 1995; Tsai & Ingham, 1997). Furthermore, acid resistance has also been described as having a cross-protection effect against heat, salt, and irradiation preservation of foods (Buchanan & Edelson, 1999; Cheng & Kaspar, 1998; Garren, Harrison, & Russel, 1998) and that acid tolerance of *E. coli* O157:H7 is sustained or even enhanced within refrigerated storage (Cheng &

Kaspar, 1998; Clavero & Beuchat, 1996; Lin et al., 1996; Miller & Kaspar, 1994).

There are two main acid tolerance mechanisms that EHEC are known to have: a pH-dependent system (acid-induced) involving arginine decarboxylase and/or glutamate decarboxylase (Hersh, Farooq, Barstad, Blankenhorn, & Slonczewski, 1996; Lin et al., 1996) and a pH-independent system (general stress protection system) regulated by  $\sigma^{38}$  encoded by *rpoS* (RNA polymerase, sigma) (Cheville, Arnold, Buchrieser, Cheng, & Kaspar, 1996; Kolter, 1993; Matin, Auger, Blum, & Schultz, 1989). *rpoS* regulates the transcription of many proteins, some of which have protective functions that could increase bacterial survival (Loewen & Triggs, 1984; Sak, Eisenstark, & Touati, 1989). Interruption of *rpoS* significantly reduces the acid tolerance of *E. coli* O157:H7, *Salmonella typhimurium*, and *Shigella flexneri* (Cheville et al., 1996; Lee, Slonczewski, & Foster, 1994; Small, Blankenhorn, Welty, Zinser, & Slonczewski, 1994). Gorden & Small (1993) found that 80% of tested *E. coli* isolates were considered acid resistant and that the acid resistance mechanism can be triggered at pH 5.0, which allows for up to nearly 1,000-fold enhanced survival of *E. coli* O157:H7 at a second exposure (Brudzinski & Harrison, 1998). Some non-O157:H7 serotypes (O26, O45, O104) have been shown to be more acid resistant than O157 (Kim, Breidt, Fratamico, & Oh, 2015). Acid resistance may enhance virulence of pathogenic strains (Arnold & Kaspar, 1995; Gorden & Small, 1993; O'Driscoll, Gahan, & Hill, 1996) which increases the need for effective, multi-hurdle decontamination efforts for animal products.

## **Ultraviolet Light**

Ultraviolet (UV) light is a common method for disinfection used in many different fields. Ultraviolet light is approved for use by the U.S. Food and Drug Administration (FDA) to decontaminate food, food contact surfaces, juices, and water because of its non-thermal and environmentally friendly nature (Bintsis, Litopoulou-Tzanetaki, & Robinson, 2000; FDA, 2000; Sommers, Sites, & Musgrove, 2010). UV light applied at 250-270 nm is considered germicidal UV-C with 254 nm the most common wavelength used for disinfection, compared to UV-A (315-400 nm) and UV-B (280-315 nm) that have limited microbial properties (Bintsis et al., 2000; Guerrero-Beltrán & Barbosa-Cánovas, 2004). The antimicrobial ability of UV-C light is based on the ability for the light to be absorbed by the DNA of microorganisms to cause pyrimidine dimerization through a thiamine and cytosine cross-linkage which cannot be replicated, leading to cell death (Escalona, Aguayo, Martínez-Hernández, & Artés, 2010; Guerrero-Beltrán & Barbosa-Cánovas, 2004; Hijnen, Beerendonk, & Medema, 2006; Sastry, Datta, & Worobo, 2000; Snowball & Hornsey, 1988). The amount of DNA damage depends on the UV absorption characteristics, the target micro-organism's resistance to UV exposure, and the applied UV dose (Ngadi, Smith, & Cayouette, 2003).

## **Bacteriophage (phage)**

Bacteriophages are bacterial viruses, discovered independently by Fredrick Twort in 1915 and Felix D' Herelle in 1917, but it was D'Herelle that coined the term "bacteriophage", from Greek *phagein* meaning "to devour" (Kutter, Kuhl, Alavidze, & Bs,

2013). Phages are nature's most abundant microorganism with over  $10^{30}$  on Earth, outnumbering bacteria 10 to 1 (Chibani-Chennoufi, Bruttin, Dillmann, & Brüssow, 2004; Wommack & Colwell, 2000). They are obligate intracellular parasites, which are organisms that require a host cell to replicate. Phages were researched in the early 20th century for their antibacterial properties to be used for bacterial infections, however, trouble creating phage therapies and the introduction of Penicillin in 1928 led to the near disappearance of phage therapy in Western countries (Kortright, Chan, Koff, & Turner, 2019). In recent time, the development of antibiotic resistant bacteria and 'superbugs' along with the decline in research and development of new antibiotics has raised a global alarm for new and alternative methods to combat life threatening resistant infections that kill over 35,000 people each year in the United States alone (CDC, 2019).

Tailed phages belong to the order *Caudovirales*, that are then divided into three families based on the tail morphology: *Myoviridae* (long contractile tail), *Siphoviridae* (long non contractile tail), and *Podoviridae* (short non contractile tail) (Ackermann, 2003). *E. coli* infecting bacteriophages, known as T4 phages belonging to the *Myoviridae* family, are relatively large viruses with complex viral structures at about 90 nm wide and 200 nm long with around 168 kbp size double stranded DNA genome stored inside their icosahedral capsid (Fuller, Raymer, Kottadiel, Rao, & Smith, 2007; Miller et al., 2003; Petrov et al., 2006). Connected to the capsid is the collar region that has short whiskers called whisker antigen control, followed by the long contractile tail consisting of a

hollow tail tube covered by a contracting tail sheath. The tail is connected to the complex baseplate at the end that has a set of six short and six long tail fibers (Cerritelli, Wall, Simon, Conway, & Steven, 1996; Coombs & Eiserling, 1977).

### **Mechanism of Infection**

There are two mechanisms of infection used by phages: the lytic cycle and the lysogenic cycle. The lysogenic cycle is typically seen in temperate phages which can integrate their DNA into the host genome and can remain there for multiple replication cycles as a prophage, but can also undergo the lytic cycle (Echols, 1972). *Escherichia coli* infecting T4 phages are strictly lytic, replicating only by the lytic cycle. Briefly, the lytic begins by 1) attachment of the long and short tail fibers to the surface of a bacterial cell via receptor specificity 2) tail sheath contraction and tail tube penetration occur in a syringe-like motion that injects phage DNA into host cytoplasm 3) replication and synthesis of phage genome and proteins using host mechanisms 4) newly synthesized phage proteins and DNA assemble into new virions 5) virion release via host cell lysis (Fineran, Petty, & Salmond, 2009).

Phage attachment is initiated by the reversible binding of at least three long tail fibers to cell surface structures via receptor specificity determined by protein gp37, a 14 amino acid residue region on the tail fibers (Heller, 1992; Montag, Hashemolhosseini, & Henning, 1990). Tail fibers bind to bacterial structures including proteins, lipopolysaccharides (LPS), teichoic acids, pili, and flagella, with the most important being OmpC porins and the O-chain on LPS (Heller, 1992; Rakhuba, Kolomiets, Szwajcer-Dey, &

Novik, 2010; Yu & Mizushima, 1982). *E. coli* and *Salmonella* have been shown to share 98% of OmpC amino acid sequence identity and immunogenic structure (Singh, Singh, Williams, Jones, & Abdullah, 1995; Singh, Upshaw, Abdullah, Singh, & Klebba, 1992). Bacteriophages are able to recognize similar core structures of LPS of related bacteria since they are relatively conserved across gram-negative species allowing for a potentially broad host range (Rakhuba et al., 2010). Long tail fiber binding causes a conformational change in the baseplate from a hexagonal dome-shaped structure to a six-point star that allows the short tail fibers to unfold and bind irreversibly to the host cell (Le et al., 2013; Yap & Rossmann, 2014). Short tail fiber binding causes the tail sheath to undergo a conformational change causing it to contract to about one-third of its original length while extending the hollow tail tube underneath, in a syringe-like motion, through the outer membrane (Kostyuchenko et al., 2005; Yap & Rossmann, 2014). To assist with tail tube penetration and cell wall lysis, the lysozyme gp5 protein on the spike-form structure at the tip of the tail dissolves the outer membrane before dissociating into the periplasmic space (Hu, Margolin, Molineux, & Liu, 2015; Kanamaru, Ishiwata, Suzuki, Rossmann, & Arisaka, 2005; Kanamaru et al., 2002). After gp5 dissociates, gp27 is able to contact the inner membrane to form a transmembrane channel, allowing the phage genome to be adsorbed directly into the host cytoplasm (Hu et al., 2015; Kanamaru et al., 2002). Adsorption of phage DNA into the host cell has been shown to be faster at higher bacterial growth rates which is positively correlated to cell size and

directly proportional to total cell surface area (Hadas, Einav, Fishov, & Zaritsky, 1997; Rabinovitch, Fishov, Hadas, Einav, & Zaritsky, 2002). In conditions where there is a high number of phages per bacterial cell (>100 phages per cell), a phenomenon called 'lysis from without' occurs where the cumulative effects of gp5 on the cell wall cause sufficient damage and quick lysis without the need for phage replication (Abedon, 2011). Upon DNA adsorption into the host cytoplasm, replication and translation begin using hijacked cytosolic machinery such as ribosomes until virion proteins are assembled and the host cell wall lyses via lysosomal activity, releasing new phages that are able to begin the cycle on to new and susceptible bacteria (Kao & McClain, 1980). Host lysis can release 10 to 400 new phages depending on host factors such as growth rate, rates of synthesis and assembly of phage components, and by lysis time which seem to depend on the content of the protein-synthesizing system and cellular dimensions (Hadas et al., 1997).

### **Phage Therapy as Biocontrol in Food**

Phages have been studied for their antibacterial ability in the medical field, and recently, have been approved for a clinical trial investigating their use as an intravenous therapy against multi-drug resistant bacteria (LaFee & Buschman, 2019). Phages have been well studied in other environments and have been used in fields such as aquaculture (Nakai & Park, 2002; Oliveira, Castilho, Cunha, & Pereira, 2012), dairy (Brüssow, 2001), and agriculture (Svircev, Roach, & Castle, 2018) due to their ability to lyse only targeted bacteria without disrupting other non-targeted microbiota (Carlton,

1999; Meaden & Koskella, 2013; Sulakvelidze, Alavidze, & Morris, 2001). Since the approval of List-Shield™ by the FDA in 2006 to target *L. monocytogenes* on ready-to-eat (RTE) foods, there have been many commercial phage products including Ecoshield™ and Listshield™ by Intralyix, Inc that target *E. coli* and *L. monocytogenes* respectively and Listex™ and PhageGuard S™ are products of Microcos Food Safety, Inc that target *Listeria* and *Salmonella* for direct food application (Bren, 2007; Kazi & Annapure, 2016). Phage use in food safety has been well investigated for reducing foodborne pathogens including *E. coli*, *Salmonella*, and *Listeria*. Phages are viewed as a natural alternative to traditional antimicrobial approaches to food safety and preservation, and have many advantages as well as notable disadvantages (Goodridge & Abedon, 2003; Greer, 2005). Research is still needed to thoroughly understand the mechanisms of phage resistance that can be acquired by the hosts and the development of methods to overcome phage resistance. Although the results of the studies appear to be encouraging, they should be interpreted with caution (Kazi & Annapure, 2016; Mahony, McAuliffe, Ross, & van Sinderen, 2011). Using bacteriophages for specific pathogen biocontrol offers an additional hurdle for robust food safety management systems (Endersen et al., 2014).

## **Conclusion**

There are many food safety interventions that can be used to improve the safety of our food supply. With the continued evolution of pathogens, our

antimicrobial interventions also need to evolve and adapt to address the changing demands. New and alternative interventions must be introduced in order to maintain an effective food safety system. Good hygienic slaughtering practices reduce fecal contamination on carcasses but does not guarantee the absence of STEC from products. The only effective method of eliminating STEC from foods is to introduce antimicrobial treatments in a multi-hurdle system (Wheeler et al., 2014). New interventions such as bacteriophages have been studied for their potential as antibacterial agents showing promise to combat acid resistant and virulent pathogens such as Shiga Toxin-producing *E. coli*.

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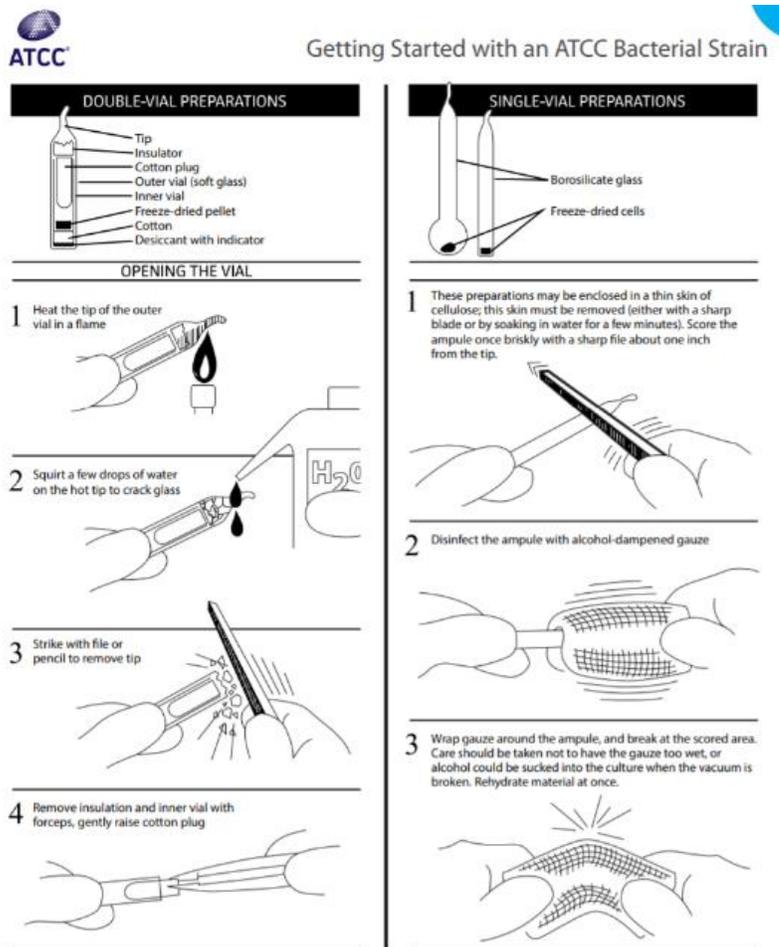
Appendix I. Safe Minimum Cooking Temperatures Charts (adapted from FoodSafety.gov)

Follow the guidelines below for minimum cooking temperatures and rest time for meat, poultry, seafood, and other cooked foods. Be sure to use a food thermometer to check whether meat has reached a safe internal temperature that is hot enough to kill harmful germs that cause food poisoning.

<b>Food</b>	<b>Type</b>	<b>Internal Temperature (°F)</b>
Ground meat and meat mixtures	Beef, pork, veal, lamb	160
	Turkey, chicken	165
Fresh beef, veal, lamb	Steaks, roasts, chops Rest time: 3 minutes	145
Poultry	All Poultry (breasts, whole bird, legs, thighs, wings, ground poultry, giblets, and stuffing)	165
Pork and ham	Fresh pork, including fresh ham Rest time: 3 minutes	145
	Precooked ham (to reheat) Note: Reheat cooked hams packaged in USDA-inspected plants to 140°F	165
Eggs and egg dishes	Eggs	Cook until yolk and white are firm
	Egg dishes (such as frittata, quiche)	160
Leftovers and casseroles	Leftovers and casseroles	165
Seafood	Fish with fins	145 or cook until flesh is opaque and separates easily with a fork
	Shrimp, lobster, crab, and scallops	Cook until flesh is pearly or white, and opaque
	Clams, oysters, mussels	Cook until shells open during cooking

Date Last Reviewed April 12, 2019

## Appendix II. Getting Started with an ATCC Bacterial Strain (ATCC, 2015)



**CHAPTER ONE**

**Application of MS bacteriophages on trimmings reduce O157:H7 and the “Big Six”  
STEC in ground beef**

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**Abstract**

According to the United States Food and Drug Administration (FDA) agency, bacteriophage solutions targeting only the serotype O157:H7 are Generally Recognized as Safe (GRAS) to control STEC during beef processing. However, outbreaks involving the “Big Six” STEC increased the industry concern about those serotypes. In this study, we evaluated the efficacy of Mello-Shebs (MS) bacteriophages targeting the “Big Six” O26, O45, O103, O111, O121, and O145 STEC. The in vitro killing efficiency of phages isolated for each specific serotype varied from 96.2% to 99.9%. When applied to contaminated trim, a 90% reduction of all STEC was observed in ground beef. Therefore, bacteriophage applications may provide an additional barrier against the “Big Six” STEC in robust food safety systems. Results of this research provide support documentation to the FDA to extend GRAS status for bacteriophages as processing aids against all adulterant STEC.

## Introduction

Shiga-toxin *E. coli* (STEC) continues to be a major public health concern due to the severity of its foodborne infections that range from very mild to life-threatening (CDC, 2018). In 2014, there were 4,437 culture-confirmed STEC infections (43.2% due to O157, 56.5% due to non-O157) and caused more than \$298 million in medical care costs in the United States (CDC, 2017; USDA-ERS, 2014). Animal products, especially ground beef, have demonstrated to be a common source of contamination of STEC (CDC, 2016). Previous research showed that STEC can be transferred to beef carcasses from hides during the harvesting process and persist on fresh beef after fabrication (Ferens & Hovde, 2011; Arthur et al., 2010; Bell, 1997; Byrne, Bolton, Sheridan, McDowell, & Blair, 2000; McEvoy et al., 2000).

The United States Department of Agriculture (USDA) deemed STEC O157:H7 adulterant in ground beef in 1994 after a multistate outbreak from undercooked hamburger from a fast food chain. In 2012, six more serotypes including STEC O145, O121, O111, O103, O45, and O26 (the “Big Six”) were also deemed adulterant due to their association with foodborne outbreaks (USDA-FSIS, 2012). According to the USDA-Food Safety Inspection Service (FSIS), beef trim tested positive for STEC still can be used for human consumption, but it must be further processed into heat-treated products to eliminate bacteria (USDA-FSIS, 2014). Positive-tested trim is sold at discounted price, leading to economic losses for the industry. Therefore, due to human health and economic impacts, regulatory agencies developed processing policies whereas private

companies implemented food safety interventions to decrease the incidence of STEC in beef products. In the United States, large beef processing facilities commonly use hot water carcass washing during harvest and applications of acidic solutions during harvest and fabrication as antimicrobial interventions (Wheeler, Kalchayanand, & Bosilevac, 2014). Although the effectiveness of acid solutions as a food safety intervention against *E. coli* is supported by some research (Harris et al., 2006; Laury et al., 2009; Schmidt et al., 2014; Wolf et al., 2012; Zhao, Zhao, & Doyle, 2009), several studies have shown that *E. coli* may become resistant to low pH after exposure to acid stress (Anderson & Marshall, 1989; Cheng et al., 2003; Kim et al., 2015; Leyer et al., 1995; Samelis, Sofos, Kendall, & Smith, 2005; Smulders & Greer, 1998). Therefore, there is an immediate need for developing more efficient aids to decontaminate beef.

Bacteriophage (phage) solutions are known as bacteria-specific interventions that can be applied on food products (Atterbury, Connerton, Dodd, Rees, & Connerton, 2003; Endersen et al., 2014; Mahony et al., 2011) Recently, phage applications were reported to significantly decrease *Salmonella* contamination in ground meats (Grant, Parveen, Schwarz, Hashem, & Vimini, 2017; Jorquera et al., 2015; Sharma, Dhakal, & Nannapaneni, 2015; Yeh et al., 2017). When in contact with a bacteria cell, phages attach to specific cell structures to introduce its genome into the bacteria. The lysis of the host is achieved by phage replication and release (Lindberg, 1973; Montag et al., 1990; Rakhuba et al., 2010; Yu & Mizushima, 1982) In the US, the Food Safety and

Inspection Service (FSIS) Directive 7120.1 lists bacteriophage solutions as a processing aid to control only one STEC serotype (O157:H7) (USDA-FSIS, 2019). The objectives of this study were: i) to demonstrate that phages can decrease “Big Six” contamination in beef; and ii) provide technical evidence to the FDA to extend GRAS status for bacteriophages as processing aids against all adulterant STEC.

## **Materials and methods**

### ***E. coli* strains and inoculum preparation**

Strains used in this study included *E. coli* O26 (ATCC® BAA-2196™), O45 (ATCC® BAA-2193™), O103 (ATCC® BAA-2215™), O111 (ATCC® BAA-2440™), O121 (ATCC® BAA-2219™), O145 (ATCC® BAA-2192™), and O157:H7 (ATCC® 35150™). Frozen stocks from each strain were briefly thawed and a loopfull of culture was aseptically streaked onto 1.6% agar LB agar plates in duplicate and incubated for 24 h at 37°C to ensure cultures were live and viable. For each strain, a single isolated colony was transferred into independent sterile tubes containing 10 mL sterile LB broth and incubated at 37°C overnight with shaking (200 rpm). To prepare the inoculum, 15 mL of LB broth was inoculated with 750 µL of an individual culture and incubated with shaking at 37°C until culture absorbance was measured to be between 0.5 and 0.6 OD<sub>600</sub>. Once each strain achieved this absorbance, equal amounts of the cultures were combined while on ice to avoid overgrowth. This *E. coli* cocktail was then diluted with sterile 0.1% Buffered Peptone Water (BPW) to create a 25% culture cocktail solution for inoculation (approximately  $1.1 \times 10^8$  CFU/mL).

### **Bacteriophage preparation and stock titer**

Solutions of bacteriophages Mello-Shebs 1 (MS1) 026, MS1 045, MS1 103, MS1 111, MS1 121, MS1 145, and MS1 157 were obtained directly from the University of Nevada, Reno - Food Safety Microbiology Laboratory Library. The concentration of phage solutions was determined by the double-layer agar methodology described by Adams (1959) and modified by Yeh et al. (2017). In short, 100  $\mu$ L of 10-fold diluted phage and 100  $\mu$ L of Log-phase *E. coli* culture was added into four mL of molten LB agar (0.6% agar) tempered at 45°C. Solution was mixed by inverting twice then evenly distributed onto hard LB agar (1.6%) plates until solidified. Duplicated plates were incubated at room temperature overnight for plaque enumeration. Phage concentration of individual solutions and the cocktail were determined to be  $10^8$  PFU/mL.

### **In vitro bacteriophage killing efficiency**

Killing efficiency of the seven bacteriophages was determined by plating each phage against each of the seven STEC serotypes. Briefly, 100  $\mu$ L of diluted overnight cultures were plated in quadruplicate onto 1.6% LB agar followed by 100  $\mu$ L of each phage solution. Plates were allowed to fully dry before being inverted and incubated overnight at 37°C. Resulting colonies were counted and compared to quadruplicate controls plated without phage.

### **Sample preparation and treatment design**

A 15 kg batch of beef trim (80% lean) was sourced from a USDA-inspected beef processing facility, screened for background *E. coli* and frozen until all analyses could be done. Upon thawing, 100 g aliquots of beef trim were randomly assigned to 8 treatments: Control 7°C 30 min, Control 7°C 6 h, Control 25°C 30 min, Control 25°C 6 h, Phage 7°C 30 min, Phage 7°C 6 h, Phage 25°C 30 min, and Phage 25°C 6 h. Control inoculated samples (Control) were treated with BPW, and Phage samples were treated with the phage cocktail. The experiment was designed as a Completely Randomized Design and the following model was used:  $Y_{ij} = \mu + \alpha_i + \varepsilon_{ij}$ , where  $Y_{ij}$  was STEC count,  $\mu$  was the overall mean across the treatments, and  $\alpha_i$  was the effect of treatment from the grand mean specific to the 8 treatment levels. The model errors ( $\varepsilon_{ij}$ ) were assumed to be random and independent of each other.

### **Beef trim inoculation, bacteriophage application, bacterial enumeration**

Trim samples were inoculated with a cocktail comprising of all STEC strains (O26, O45, O103, O111, O121, O145, and O157:H7) to yield in a contamination level of about 5 Log CFU/g in the ground product. A volume of 1 mL of the inoculum was applied by dripping to each side of trim samples, for a total of 2 mL for each 100 g sample. After 30 minutes at 7°C, samples were treated with 2 mL of bacteriophage cocktail solution ( $10^8$  PFU/mL of MS1 026, MS1 045, MS1 103, MS1 111, MS1 121, MS1 145, and MS1 157 phages) or with 2 mL of BPW. One (1) mL was distributed on each side of the trim. Subsequently, samples were covered and held for either 30 min or 6 h while at 25°C or

7°C. After holding, samples were aseptically ground using a tabletop electric grinder (model 33-0201-w, Weston China). A 25g aliquot of each sample was collected in a sterile sampling bag and masticated with 225ml of BPW for two minutes at 230 rpm. For *E. coli* enumeration, 10 mL of the masticated homogenate was centrifuged at 10,000 x *g* for 6 minutes and the supernatant discarded to avoid plating phages. The pellet was resuspended with 10 mL of BPW, vortexed, serial diluted in BPW, and 100 µL was plated onto LB agar in duplicate. The plates were inverted and incubated at 37°C for 24 h and the resulting colonies counted (CFU/g).

### **Statistical analysis**

The experiment was replicated 5 times with 2 repetitions per treatment (n=80 total e.u., n=10 e.u. per treatment). Data was analyzed using the GLIMMIX procedure of SAS (® 9.3 package, SAS Institute, Inc., USA). Contrast analyses were also performed to evaluate single effects of bacteriophage application, and lysing temperature and time. When significance ( $P \leq 0.05$ ) was detected, means separations and significance for main treatments were performed by using the LSMEANS and DIFF function statements.

### **Results**

The results of killing efficiency of each bacteriophage against all STEC serotypes is demonstrated in Table 1. When looking to effects of serotype-

specific phage against its respective STEC (gray highlighted diagonal in the table), phages demonstrated at least 96.2% of killing efficiency (MS1 103 against STEC O103). The highest killing efficiency (99.9%) was observed for MS1 026 and MS1 157. Overall, when tested against their non-specific STEC serotypes, killing efficiency of MS1 phages varied from 34.1% (MS1 103 against STEC O111) to 99.8% (MS1 157 against STEC O26).

Effects of the phage cocktail application on contaminated beef trim and resulting STEC populations after grinding are presented in Figure 1. Overall, bacteriophage application on trim significantly decreased STEC populations in ground beef in all temperature and lysing time scenarios. Reductions from 4.68<sup>AB</sup> to 3.87<sup>DE</sup>, 5.12<sup>A</sup> to 3.82<sup>DE</sup>, 4.74<sup>AB</sup> to 3.98<sup>DC</sup>, and 4.39<sup>BC</sup> to 3.40<sup>E</sup> Log CFU/g, were observed for 25°C 30 min, 25°C 6 h, 7°C 30 min, and 7°C 6 h, respectively. Reductions in Log are numerically presented in Table 2. When analyzing contrasts, there was no effect of lysing time ( $P = 0.2797$ ), but a significant effect of lysing temperature was observed ( $P = 0.0497$ ).

## Discussion

Bacteriophages MS1 exerted high killing efficiency in vitro against their respective serotypes (above 1 Log). All MS1 bacteriophages were isolated from sewage as described by Mirzaei, Eriksson, Kasuga, Haggård-Ljungquist, & Nilsson (2014), Hudson et al. (2013), Carey-Smith, Billington, Cornelius, Hudson, & Heinemann (2006), Hooton, Atterbury, & Connerton (2011), and Pereira et al. (2016) by using the same STEC serotypes applied on beef trim in this study. The ability for each phage to lyse other non-specific STEC serotypes is possible associated to its genetic similarity. Although complete characterization of MS1 phages is still currently in progress, genomic data

deposited at GenBank® showed that MS1 phages have similar nucleotide sequences when compared to other *E. coli* phages. The MS1 phages were isolated using the same strain (*E. coli*) but different serotypes (O26, O45, O103, O111, O121, O145, and O157:H7), which explains the similarity and possible differences in their genome. Bacteriophages targeting the same strain usually show similar nucleotide sequences, however, genomic differences that make them more efficient against different serotypes is possibly associated to the ability of phages to move between serotypes in the environment and acquire different host genomic material, which potentially broadens their host range (Bergman, Fineran, Petty, & Salmond, 2019).

To our knowledge this is the only study that evaluates the efficacy of a 7-bacteriophage cocktail to reduce STEC in ground beef. The use of a phage cocktail prevents the emergence and phage resistance mutants as the different phages are able to bind to different host cell surface receptors adding to the versatility and overall effectiveness of a bacteriophage solution (Sulakvelidze & Barrow, 2005). The ability for each phage to lyse every STEC strain that it was not isolated against, increases the efficacy of a cocktail that targets all 7 adulterant serotypes that may be present on beef products, rather than a single serotype. There are many reported *E. coli* targeting bacteriophage cocktails, but none contain more than six individual phages (Carter et al., 2012; Coffey et al., 2011; Greer, 2005; Magnone, Marek, Sulakvelidze, & Senecal, 2013; O'Flynn et al., 2004; Tomat et al., 2018; Viazis, Akhtar, Feirtag, & Diez-Gonzalez, 2011). A commercially available bacteriophage preparation with 3 phages was evaluated by

(Carter et al., 2012) against *E. coli* O157:H7 and demonstrated over 94% and 87% reduction on beef and lettuce, respectively. O'Flynn et al. (2004) showed that a three-phage cocktail targeting STEC O157:H7 completely eliminated bacteria cells on beef surface after 1 h at 37°C.

Regarding lysing temperature, Tomat et al. (2018) evaluated a six-phage cocktail against four strains of *E. coli*, an enteropathogenic *E. coli* (EPEC) O157, DH5 $\alpha$ , and two STEC (O157:H7 and O18) and demonstrated that a phage cocktail can maintain low *E. coli* contamination levels on meat surface at 4°C, but not at 24°C or 37°C when regrowth was observed. Previously, Leverentz et al. (2001) demonstrated that phage efficacy was independent of temperature (5°C, 10°C, 20°C) against *Salmonella* on fresh fruit. Conversely, the majority of research reports, including this study, showed that bacteriophage applications have shown to have higher efficiency at higher temperatures, particularly at 24°C or at optimal bacteria growth range (37°C) (Bigwood, Hudson, Billington, Carey-Smith, & Heinemann, 2008; Hudson et al., 2013; Hudson, Billington, Wilson, & On, 2015; Liu et al., 2015; Sharma, Dhakal, & Nannapaneni, 2015; Spricigo et al., 2013). Optimal phage activity at higher temperatures is possibly related to the metabolism of the host and its effect on the lytic cycle. Hadas et al. (1997) described how phage adsorption depends on the growth rate of host *E. coli* cells which seems to be dependent on nutrient availability, the rate of the protein synthesizing system, and cellular dimensions. Additionally, Sillankorva et al. (2004) suggested that the optimal growth temperature for *E. coli* ranges between 37-42°C, whereas while no growth is usually observed at refrigeration temperatures, 5-8°C (Buchanan & Klawitter,

1992; Buchanan, Stahl, & Whiting, 1989; Gonthier, Guérin-Faubleé, Tilly, & Delignette-Muller, 2001; Nauta & Dufrenne, 1999; Rabinovitch et al., 2002; Sillankorva et al., 2004). However, studies performed by our team (Yeh et al., 2017; Yeh et al., 2018) are in agreement with previous research that reported significant bacteria reduction when phages were applied at low temperatures (Fiorentin, Júnior, Suínos, & Author, 2005; Guenther, Herzig, Fieseler, Klumpp, & Loessner, 2012). This was possible because bacteria thrive in cold temperatures by adopting a cold tolerance basal metabolism, allowing the cell to remain alive (Sengupta & Chattopadhyay, 2013). Authors then hypothesized that even in conditions where bacteria adopt this type of metabolism, phages are still able to perform their lytic cycle. In this study, the contrast analysis revealed significant differences ( $P = 0.0497$ ) between treatments at 7 and 25°C, suggesting that temperature speeds up the lyse of the host. A similar trend was suggested by Liu et al. (2015), who evaluated 3 temperatures including 4°C, 22°C and 37°C and demonstrated that phage applied at higher temperatures took less time to achieve the same level of *E. coli* O157 reduction compared to reduction rates observed under refrigeration temperatures (3.2 Log CFU/cm<sup>2</sup> reduction within 3 h at 37°C vs 144 h at 4°C).

Similar to temperature, previous studies have also demonstrated the effects of time during phage lysing activity. Carter et al. (2012) tested different lysing times for a phage cocktail against *E. coli* O157:H7 on beef from 5 min to 24 h and revealed that there was no significant bacteria reduction with longer

phage contact time. Sharma, Dhakal, & Nannapaneni (2015) also showed no effect of time in reducing *Salmonella* counts on raw turkey with phage application after 2, 12, or 24 h. Similarly, our team previously tested bacteriophage applications against *Salmonella* holding red meat products at 4°C for 6 and 18 h, and no effect of time was observed (Yeh et al., 2017). In this study, after we applied the phage cocktail on contaminated trim, samples were held for 30 min or 6 h prior to grinding. The contrast analysis showed no effects of lysing time on ground beef either held for 30 min or 6 h ( $P = 0.2797$ ). However, previous research showed that longer lysing times may lead to higher bacteria reduction. Seo et al., 2016 showed a gradual decrease in *E. coli* O157:H7 populations overtime with the application of bacteriophage, being reduced from 5.09 Log CFU/cm<sup>2</sup> to 1.37 Log CFU/cm<sup>2</sup> at 4 h and being completely inhibited after 8 h. In similar fashion, Spricigo, Bardina, Cortés, and Llagostera et al. (2013) tested phage applications on chicken breast contaminated with *Salmonella* at 4°C and observed a 1.6 Log CFU/g reduction after one day and 2.2 after 7 days. Conversely, Yang, Sadekuzzaman, and Ha (2017), demonstrated an initial 0.8 Log CFU/g reduction of *Listeria* when applying phages on contaminated chicken breast. However, when lysing time was extended to 72 h, *Listeria* gradually regrew achieving similar values reported for contaminated samples at 0 h.

Phage DNA adsorption occurs faster at higher bacterial growth rates, and lysing time is limited by the rate of protein synthesis that is dependent on the host protein systems (Hadas et al., 1997). Previous research showed that host inactivation begins within the first few hours after phage application, even though phages are able to

complete the first lytic cycle ending in the release of new phages in 20-40 minutes depending on the MOI (Atterbury et al., 2003; Campbell, 2003; Guosheng et al., 2003; Leverentz et al., 2001; Rabinovitch et al., 2002). Overall although phages are efficient in decreasing bacteria populations in refrigerated temperatures, it seems that in higher temperatures, higher reductions may be achieved.

## **Conclusion**

According to the FSIS Directive 7120.1, only phage solutions targeting *E. coli* O157:H7 are allowed to be used as a processing aid during meat production. In this study, we demonstrated that bacteriophages are also effective against the “Big Six” serotypes. This study provides evidence to the FDA and to biotechnology companies that bacteriophages targeting all adulterant STEC, not only O157:H7. Therefore, future FDA requests to obtain GRAS status for phages targeting the “Big Six” must be considered. In addition, results of this research demonstrated that bacteriophage applications reduce STEC loads when applied in warm and cold temperatures, suggesting that phage solutions may be successfully used in different processing points from harvest and fabrication.

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Table 1. Lysing efficiency<sup>1</sup> of individual bacteriophages MS1-O157:H7, MS1-O145, MS1-O121, MS1-O111, MS1-O103, MS1-O45, and MS1-O26 against STEC O157:H7 (ATCC<sup>®</sup> 35150<sup>™</sup>), O26 (ATCC<sup>®</sup> BAA-2196<sup>™</sup>), O45 (ATCC<sup>®</sup> BAA-2193<sup>™</sup>), O103 (ATCC<sup>®</sup> BAA-2215<sup>™</sup>), O111 (ATCC<sup>®</sup> BAA-2440<sup>™</sup>), O121 (ATCC<sup>®</sup> BAA-2219<sup>™</sup>), and O145 (ATCC<sup>®</sup> BAA-2192<sup>™</sup>) in vitro.

STEC serotype	Bacteriophage						
	MS1-O157:H7	MS1-O145	MS1-O121	MS1-O111	MS1-O103	MS1-O45	MS1-O26
<b>O157:H7 (ATCC 35150)</b>	99.9%	96.5%	99.1%	50.9%	45.8%	54.9%	77.2%
<b>O145 (ATCC BAA-2192)</b>	98.8%	99.6%	97.1%	89.0%	84.0%	62.8%	96.7%
<b>O121 (ATCC BAA-2219)</b>	87.0%	88.0%	96.6%	85.6%	79.1%	73.4%	58.5%
<b>O111 (ATCC BAA-2440)</b>	98.9%	99.9%	99.3%	99.4%	34.1%	82.2%	67.3%
<b>O103 (ATCC BAA-2215)</b>	99.7%	97.4%	79.7%	97.0%	96.2%	55.6%	93.7%
<b>O45 (ATCC BAA-2193)</b>	99.4%	96.2%	88.5%	93.3%	95.8%	98.6%	68.4%
<b>O26 (ATCC BAA-2196)</b>	99.8%	96.1%	95.6%	97.6%	80.7%	58.7%	99.9%

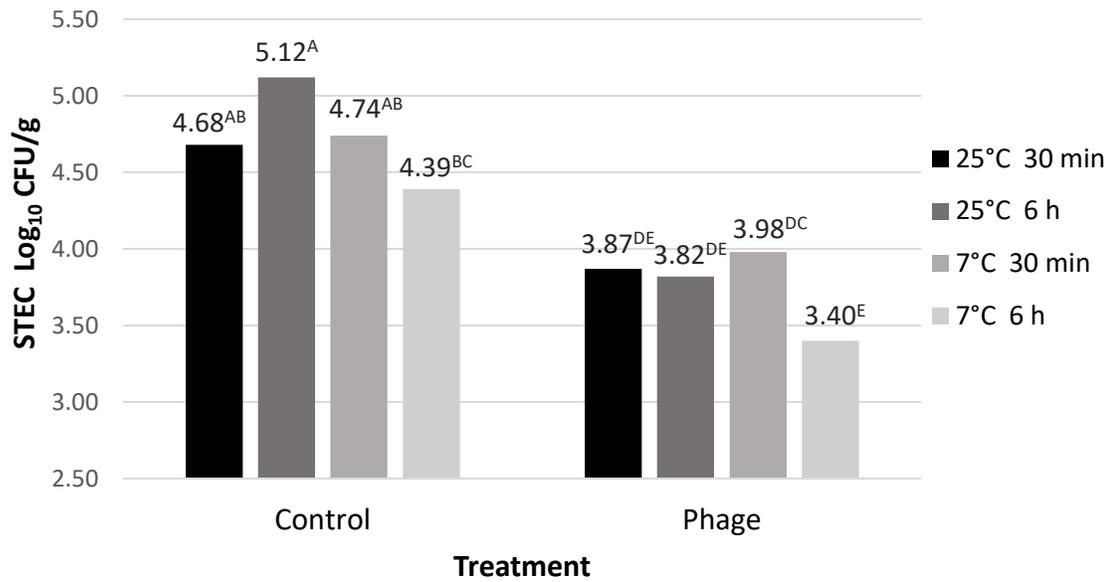
<sup>1</sup> In vitro lysing efficiency of individual phages based on the average CFU of 4 LB agar plates treated with phage versus inoculated.

Table 2. Means of Shiga Toxin *E. coli* contamination level (Log CFU/g) strains in ground beef following application of control or bacteriophage cocktail preparation MS1-O26, MS1O-45, MS1-O103, MS1-O111, MS1-O121, MS1-O145, and MS1-O157) at two lysing times 30 min and 6 h at 7°C or 25°C on beef trim prior to grinding.

<b>Holding Time</b>	<b>Temperature (°C)</b>	<b>Control*</b>	<b>Bacteriophage application</b>	<b>Reduction (Log<sub>10</sub> CFU/g)</b>
30 min	7	5.09	4.34	0.76
	25	5.03	4.22	0.81
6 h	7	4.74	3.76	0.98
	25	5.48	4.18	1.30

\* No bacteriophage applied.

Figure 1. Least Square Means of Shiga Toxin *E. coli* populations (Log CFU/g) in ground beef following control and phage applications on beef trim with fixed effects of time (30 min, 6 h) and temperature (7° C, 25° C). Std error = 0.2477,  $P < 0.0001$ . Contrasts: Phage vs Control,  $P < 0.0001$ ; 7°C vs 25°C,  $P = 0.0497$ ; and 30 min vs 6 h,  $P = 0.2797$ .



### Appendix III. Phage sewage isolation and purification

Raw sewage was acquired from Reno Municipal sewer wastewater treatment facility from both north and south sources. Aseptically, 40 mL of sewage (equal parts of both sources) were combined in sterile Erlenmeyer flasks and inoculated with 5 mL of overnight pure culture of a single strain of Shiga Toxin-producing *E. coli* (O157:H7, O145, O121, O111, O103, O45, O26). Flasks were covered and incubated with shaking (230 rpm) overnight at 37°C.

Subsequently, 10 mL of solution was centrifuged at 5,000 x g for 20 minutes and the supernatant was filtered with a 0.45 µM syringe filter twice. Soft agar (appendix IV) was performed on the resulting lysate. Plaques were evaluated visually, and a single, well-isolated, wide, and clear plaque was extracted from each serotype agar plate and placed in 1 mL of sterile Buffered Peptone Water (BPW) with one drop of chloroform before vortexing.

These isolated phages were amplified by combining 10 mL of sterile Luria Broth (LB) with 1 mL of overnight pure culture and 1 mL of the isolated phage extract and incubating overnight with shaking at 37°C. Subsequently, soft agar (appendix IV) assay was performed and a single, well-isolated, wide, and clear plaque was extracted from each serotype agar plate and placed into 1 mL of BPW and vortexed. This phage solution was amplified into an isolated working phage solution.

Appendix IV. Double-layer Agar Assay Adams (1959), modified by Yeh et al. (2017)

1. Prepare 0.6% soft agar (1 g Bacteriological agar and 4 g Luria Broth, Miller per 200 mL deionized water)
2. Sterilize the mixture in autoclave at 121 °C for at least 15 minutes.
3. Allow the solution to cool in a water bath tempered at 45°C.
4. Equilibrate 50 mL conical tubes in the water bath prior to aseptically adding 4 mL of the soft agar into each tube.
5. Add 100 µL of Log phase or overnight bacteria into the tube followed by 100 µL of serially diluted phage to achieve ~100 plaques per plate
6. Invert twice and pour slowly onto a pre-made 1.6% LB agar plate and spread soft agar across surface of plate and allow to solidify
7. Incubate overnight (not inverted) at 37°C for plaque enumeration

## **CHAPTER TWO**

### **Efficiency of bacteriophage and organic acids in decreasing STEC O157:H7 populations in beef kept under vacuum and aerobic conditions: A simulated High Event Period scenario**

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**Abstract**

During High Event Periods, beef subprimals are usually removed from vacuum and treated with antimicrobials. After re-packaging, subprimals are tested to verify the absence of STEC. In this study, bacteriophage and organic acids were applied on fresh beef surface contaminated with STEC O157:H7 to evaluate the efficiency of industry practices. Beef samples inoculated with STEC were treated with bacteriophage, lactic acid, and peroxyacetic acid and kept under vacuum or aerobic conditions. STEC loads were evaluated 30 min and 6 h after antimicrobial applications. Under aerobic conditions for 30 min and 6 h, phage reduced STEC in beef by approximately 1.4 Log CFU/cm<sup>2</sup> whereas organic acids led to approximately 0.5 Log CFU/cm<sup>2</sup> reduction. Under vacuum conditions for 30 min, bacteriophage significantly reduced STEC by 1 Log CFU/cm<sup>2</sup>, while no significant effects were observed in samples treated with organic acids. Under vacuum conditions for 6 h, bacteriophage reduced STEC loads by 1.4 Log CFU/cm<sup>2</sup>, lactic acid reduced by 0.6 Log CFU/cm<sup>2</sup>, and no significant effects were observed with peroxyacetic acid application.

**Keywords:** Bacteriophage, PAA, lactic acid, high event period, STEC, E. coli O157:H7, beef

## Introduction

In the United States, the production interval when beef processing establishments experience an elevated rate of STEC contamination in trimmings is referred to as a “High Event Period” (HEP). In large operations, contamination of trimmings is assessed by sampling combos using the N60 or N60 plus methods (USDA-FSIS, 2015). More recently, novel methodologies using a manual sampling device (MSD) and a continuous sampling device (CSD) have been also been implemented to sample combos (Wheeler & Arthur, 2018). The United States Department of Agriculture (USDA) – Food Safety Inspection Service (FSIS) considers an establishment’s process to be within a HEP if the percent of positive samples within a specified production interval is 5% or greater (USDA-FSIS, 2014). A HEP may indicate a localized out-of-control event or a systemic breakdown of slaughter dressing operations that led to contamination of beef carcass components including primals, subprimals, and trimmings (USDA-FSIS, 2014). Previous studies have identified that hides are the major source of contamination of carcasses during processing (Arthur et al., 2008; Bell, 1997; Byrne, Bolton, Sheridan, McDowell, & Blair, 2000; McEvoy et al., 2000). During hide removal, bacteria on hides can be transferred onto the carcass (Barkocy-Gallager et al., 2003; Nou et al., 2003) whereas contamination of trim and primals depend on incoming bacterial loads and the ability of establishments to prevent cross contamination between hides and carcasses during dressing

procedures (Brichta-Harhay et al., 2008). Although a decrease of STEC loads is achieved by implementing good dressing procedures and antimicrobial applications during slaughter (Brashers and Chavez, 2017), Yang, He, Badoni, Tran, and Wang (2017) showed that 62.2% of the *E. coli* found in meat are shared during processing and fabrication by mesh gloves, conveyor belts, and cutting tables. Therefore, if not eliminated or reduced by interventions during slaughter, higher STEC loads may be disseminated during fabrication to contaminate trimmings and primals, and lead to higher rates of positive STEC trim samples.

According to the USDA-FSIS, while STEC positive combos must be diverted to further processing to be heat treated, beef primals and subprimals produced from the same carcasses that generated positive trimmings must be tested before entering commerce (USDA-FSIS, 2014). After experiencing a HEP, establishments rework subprimals associated with contaminated trimmings by (1) removing them from vacuum sealed bags, (2) treating with antimicrobials, and (3) repackaging. Subsequently, in order to ensure that subprimals are not contaminated, the pieces are (4) removed from vacuum sealed bags again, and (5) the meat surface is swabbed to test for STEC contamination. Industry personnel estimate that the time spent from steps 1 to 4 may vary from 30 min to 6 h depending on processing speed and production volume.

In the United States, application of organic acids to decontaminate meat surfaces is the most common intervention used to decrease bacteria loads and eliminate STEC (Wheeler, Kalchayanand, & Bosilevac, 2014). Advantages of using those acids include availability, cost, and simplicity of usage (Mataragas, Skandamis, &

Drosinos, 2008). However, previous studies demonstrated that *E. coli* strains may develop resistance to commonly used organic acids including lactic acid (Smulders & Greer, 1998; Dickson & Kunduru, 1995; Anderson & Marshall, 1989). In this study, we evaluated the killing efficiency of organic acids including peroxyacetic and lactic acids and a bacteriophage solution on fresh beef contaminated with STEC, under vacuum and aerobic conditions, by simulating steps and time intervals where primals and subprimals are reworked and retreated after a HEP.

## **Materials and Methods**

### ***E. coli* strains and inoculum preparation**

Four strains of *E. coli* O157:H7 were used in this study to ensure that interventions were tested against different genotypes. Strains included ATCC® 35150™, ATCC® 43895™, ATCC® 43894™, and NCTC 13128. All ATCC strains were *stx1* and *stx2* positives whereas the NCTC strain was negative for both genes. ATCC strains were recovered by following the ATCC bacterial culture guidelines (ATCC, 2015) whereas the strain NCTC 13128 was directly recovered from a glycerol stock by streaking the content of the micro tube onto LB agar plates. Briefly, strains were thawed and a loop full of each stock was aseptically streaked on to 1.6% LB agar plates in duplicate. After incubating overnight at 37°C, one isolated colony was aseptically added to 10 mL of sterile LB broth and incubated overnight at 37°C with shaking (200 rpm). Subsequently, 1 mL was then transferred into a tube containing LB broth and cultured overnight at

37°C. After approximately 16 h, final absorbance was verified at 600 nm ( $OD_{600}$ ) and CFU concentration per mL was estimated by using the Agilent *E. coli* cell culture biocalculator (AGILENT, 2019). Cultures were individually diluted in 0.1% sterile buffered peptone water (BPW) to achieve  $1 \times 10^7$  CFU/ml. The final inoculum was prepared by combining equal amounts of all four individual *E. coli* strains.

### **Bacteriophage solution titer and killing efficiency**

The commercial preparation PhageGuard E<sup>TM</sup> containing two bacteriophages (EP75 and EP335) was provided by MICREOS Food Safety, Inc. (Wageningen, The Netherlands). Bacteriophage EP75 is a Vi1virus, that belongs to the *Caudovirale* order, *Ackermannviridae* family, and subfamily *Cvivirusinae* whereas the bacteriophage EP 335 is a Phi32virus that belongs to *Caudovirale* order and *Podoviridae* family (Van Mierlo et al., 2019). Phage stock concentrations was determined by following the double-layer agar method with some modifications (Adams, 1959). Briefly, the PhageGuard E<sup>TM</sup> stock solution was serially diluted in sterile potable water and added into Luria Broth (LB) soft agar (0.6% agar) tempered at 45°C after inoculation with individual fresh-log phase *E. coli* cultures. Tubes were vortexed twice and distributed onto the lawn of hard LB hard agar plates (1.6%) in duplicate. The soft agar was allowed to solidify at room temperature and subsequently, plates were inverted and incubated overnight at 37 °C. Concentration of the PhageGuard E<sup>TM</sup> was determined to be  $2 \times 10^{11}$  PFU/ml.

Killing efficiency of PhageGuard E<sup>TM</sup> was determined by testing the stock solution against individual *E. coli* strains. A volume of 100 µl of diluted overnight *E. coli* cultures

were spread onto LB agar plates. After the plates dried, the same volume of stock bacteriophage solution was applied onto the lawn of the plates. Plates were allowed to dry and were inverted and incubated overnight at 37 °C. Killing efficiency was determined in quadruplicate. Bacteriophage was not applied onto control plates.

### **Sample preparation and experimental design**

Beef Rose Meat (*m. cutaneous trunci*, IMPS 194) (USDA-AMS, 2014) was procured from a USDA-inspected facility and transported under refrigeration (4 °C) to the University of Nevada, Reno's Meat Quality Laboratory. Upon arrival, aliquots from individual coolers (n = 4, approximately 18 kg each) were screened for generic *E. coli*. A total of 160 samples measuring 10 x 10 cm<sup>2</sup> were subsampled, fabricated from rose meat pieces, and assigned to a completely randomized design (CRD). Fixed effects included antimicrobial treatment (CI = Control Inoculated, PGE = Bacteriophage-treated, PAA = peroxyacetic acid-treated, and LA= lactic acid-treated), packaging atmosphere (V=under vacuum and NV=wrapped with O<sub>2</sub> permeable film), and lysing time (30 min and 6 h).

### **Sample inoculation, antimicrobial applications, and bacterial enumeration**

Beef samples were inoculated with a cocktail containing all four *E. coli* strains at 1x10<sup>7</sup> CFU/ml. A volume of 500 µl of the inoculum was uniformly pipetted on meat and distributed throughout the whole surface area of the sample using a sterile plastic spreader. In order to simulate contamination of

beef trimmings and beef cuts in a HEP scenario, samples were kept under exposure of O<sub>2</sub> (NV) or in vacuum conditions (V). All samples were kept under refrigeration prior inoculation and during the trial (5±2°C). Samples wrapped with O<sub>2</sub> permeable film (NV) were: (1) inoculated, (2) wrapped with film and stored for 17-24 h, (3) removed from package and treated with antimicrobials (PGE, PAA, or LA), (4) wrapped with film and stored for 30 min or 6h, and (5) removed from package and swabbed for bacteria enumeration. Vacuum packaged samples were processed in similar fashion. Samples were (1) inoculated, (2) vacuum packaged (model VMC100P, VacPak-It, Lititz, PA, United States) and stored for 17-24 h, (3) removed from package and treated with antimicrobials (PGE, PAA, or LA), (4) vacuum packaged and stored for 30 min or 6h, and (5) removed from package and swabbed for bacteria enumeration. All antimicrobials were applied at a volume of 500 µl by pipetting and distributing the solution throughout the sample surface. PhageGuard E™ was diluted in potable water as recommended by the manufacturer and applied at a concentration of 1x10<sup>8</sup> PFU/ml. Lactic acid solution (2-hydroxypropionic acid, 87.5–88.5%, w/w) was applied at 5% at 50°C, whereas PAA (Ethaneperoxoic acid, stabilized, < 43%) was applied at 400 ppm at room temperature. Control inoculated samples were treated with BPW. For bacteria enumeration, all surface area was swabbed (QS1000, Hygiena, Camarillo, CA, United States) and the swab homogenate was vortexed, serially diluted, and 100 µL was spread-plated onto LB agar plates in duplicate. Plates were inverted and incubated overnight at 37°C, and *E. coli* colonies were enumerated (CFU/cm<sup>2</sup>). Differently than our previous experiments (Yeh et al., 2017; Yeh, de Moura, Van Den Broek & de Mello, 2018), the homogenate obtained

by swabbing the beef samples was not treated to eliminate possible residual phages or organic acids before plating. Our goal was to mimic real industry scenarios where phage separation and organic acid neutralization are not practiced by laboratories after receiving samples from processing facilities.

### Statistical analysis

The experiment was conducted as a CRD and was replicated 5 times whereas 1 replication had 1 repetition, 1 replication had 3 repetitions, and 3 replications had 2 repetitions (n=160 total e.u., n=10 per fixed effects combination). The following model was used:  $Y_{ij} = \mu + \alpha_i + \beta_j + \gamma_k + (\alpha\beta)_{ij} + (\alpha\gamma)_{ik} + (\beta\gamma)_{jk} + (\alpha\beta\gamma)_{ijk} + \varepsilon_{ijkl}$ , where  $Y_{ij}$  was *E. coli* count,  $\mu$  was the overall mean across the treatments included in the experiment,  $\alpha_i$  was the effect of antimicrobial from the grand mean specific to the  $i$  levels (CI, PGE, PAA, and LA),  $\beta_j$  was the effect of lysing time from the overall mean specific to  $k$  levels (30 min and 6 h), and  $\gamma_k$  was the effect of packaging atmosphere from the overall mean specific to  $j$  levels (V and NV). All interactions and respective errors were also added in the model. In addition, we identified two nested models of interest within packaging atmosphere ( $\gamma$ ). Therefore, data was also analyzed separately as a CRD for V and NV and the following model was used for both:  $Y_{ij} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \varepsilon_{ijl}$ . Data were analyzed using the GLIMMIX procedure of SAS® 9.3 package (SAS Institute, Inc., USA) as a 4 x 2 x 2 factorial for the full model and 4 x 2 for each nested

model. When significance ( $P \leq 0.05$ ) was indicated by ANOVA, means separations were performed by using the LSMEANS and DIFF function statements.

## Results

Killing efficiency of PhageGuard E™ is shown in table 1. Bacteriophages EP75 and EP335 reduced the growth of the strains ATCC® 35150™, ATCC® 43895™, ATCC® 43894™, and NCTC 13128 by approximately 98.2%, 97.2%, 96.7%, and 98.3%, respectively. A three-way interaction ( $P=0.035$ ) was observed when evaluating the full model including the three fixed effects (antimicrobial treatment, packaging atmosphere, and lysing time). However, when looking at individual effects, antimicrobial treatment and packaging atmosphere were highly significant ( $P<0.0001$ ) whereas lysing time did not individually drive the three-way interaction ( $P=0.226$ ) (table 2).

Results of the nested model evaluating samples kept under vacuum are shown in figure 1. Control inoculated samples (CI) treated with BPW yielded 2.07 and 2.37 Log CFU/cm<sup>2</sup> after 30 min and 6 h, respectively. The ANOVA showed an interaction between antimicrobial treatment and lysing time ( $P=0.006$ ). When samples were treated with PGE, 1.07 and 1.41 Log reductions were observed for lysing time of 30 min and 6h, respectively. Within lysing time of 30 min, no statistical difference was observed when comparing CI, LA, and PAA, whereas PGE applications significantly reduced bacterial load. Overall, samples treated with PGE yielded the lowest bacterial load when compared to CI and organic acids treatments.

Results for samples kept under aerobic conditions are shown in table 3. For this model, no interaction between antimicrobial treatment and lysing time was observed ( $P=0.986$ ). Individual effects of lysing time were also not significant ( $P=0.432$ ). However, the effects of antimicrobial treatment were statistically significant at  $P<0.0001$ . Under aerobic conditions, samples treated with organic acids showed lower *E. coli* counts when compared to CI samples. Lactic acid applications reduced bacteria loads by 0.47 Log CFU/cm<sup>2</sup> whereas PAA reduced 0.53 Log CFU/cm<sup>2</sup> when compared to CI samples. Bacteriophage applications led to the optimal reduction, lowering bacteria counts by 1.41 Log CFU/cm<sup>2</sup> when compared to CI samples. Therefore, in both packaging atmosphere conditions (V and NV), bacteriophage applications showed improved control of *E. coli*, when compared to LA and PAA.

## Discussion

The killing efficiency of the bacteriophage solution against the 4 strains was at least 96.7%. Both *caudovirales* phages were previously described by Mierlo et al. (2019) as STEC-specific phages with genome size of >50 nucleotides. The EP 75 shares its nucleotide identity of 96% with the *Escherichia* phage Phaxl, 92%, is homologous to *Escherichia* phage ECML-4, and have at least 91% nucleotide identity with other *Salmonella* phages. The bacteriophage EP 335 has nucleotide identity of 96% with the *Escherichia* phage KBNP1711 and 85.47% with *Escherichia* phage NJO1. The EP 335 is also homologous to two more *Escherichia* phages and to the vB\_EcoP\_SU10, a

bacteriophage that can infect a wide range of *E. coli* strains (Mirzaei, Erikson, Kasuga, Haggård- Ljungquist, and Nilsson, 2014). Bacteriophages move between different host species and acquire different genetic material, which possibly, broadens their host range.

To our knowledge, this study is the first to compare the effects of bacteriophage versus organic acid solutions on the reduction of *E. coli* contamination in vacuum and aerobically packaged beef. After a HEP, beef processors usually segregate subprimals corresponding to STEC positive trim combos to be treated with antimicrobials. These subprimals are only allowed to enter the commerce after tests are negative for STEC, otherwise they must be diverted to further processing as positive trimmings. In the United States, LA and PAA are two of the most common organic acids used to decontaminate beef carcasses and subprimals. Peroxyacetic acid is a combination of hydrogen peroxide and acetic acid that when diluted in water is capable to disrupt chemiostic functions of the membrane and inhibit bacterial biochemical metabolism (Baldry & Fraser, 1988; Leaper, 1984; Liberti & Notarnicola, 1999). The mode of action of LA is based on the disruption of cell regulation, which requires the bacterium to use additional amount of energy to maintain its internal optimal pH in order to avoid acid stress damage on cellular mechanisms (Desriac et al., 2013; Foster, 1995).

The efficacy of both organic acids has been reported by several authors, whose results showed inconsistent reductions varying from no reduction to 3.5 Log. Regarding PAA, Penney et al. (2007) evaluated spray applications at 180 ppm on beef and veal carcass surfaces and observed a 3.5 Log reduction of nonpathogenic *E. coli* O157:H7.

Ellebracht et al. (2005) detected no more than 1.2 Log reduction when dipping beef trimmings in PAA solutions at 200, 500, and 1000 ppm. Both studies used only one *E. coli* O157:H7 strain (Penney et al. used strain NZRCC 3614 whereas Ellebracht et al. used a pathogenic rifampicin-resistant strain) and tested the efficiency of PAA using different application methods. Based on results presented in both studies, it was expected to observe optimal reduction when samples were submerged in PAA solutions due to a better distribution of the antimicrobial on trimmings. Conversely, Penney et al. observed lower counts of *E. coli* counts when compared to results presented by Ellebracht et al. In addition, King et al. (2005), using the same strain as used by Ellebracht et al., did not observe a significant reduction of *E. coli* when PAA was applied on contaminated carcass surfaces.

For years, LA has been widely used in the United States to decontaminate beef carcasses and subprimals (Fouladkhah, Geonaras, Yang & Sofos, 2013). Bosilevac, Nou, Barkocy-Gallagher, Arthur and Koohmaraie (2006) showed a 1 Log reduction of *Enterobacteriaceae* counts when applying a solution of 2% LA at 42°C on carcasses. Castillo et al. (2001) tested the efficiency of LA on the same *E. coli* strain used by Ellebracht et al. and showed that a solution of 4% led to a 2-log reduction when a post chill spray was applied onto outside rounds. In addition, Pittman et al. (2012) tested the effectiveness of LA on five strains of STEC O157:H7 and achieved a 1.6 Log reduction of bacterial loads when applying LA at 5% under vacuum conditions with lysing time of 24 h. and in the same study, an additional 1.2 Log reduction was observed after a second application of LA after beef samples were removed from vacuum. However, Uytendaele

et al. (2001) showed no reduction of two (n=2) different *E. coli* strains (an acid resistant - II/45/4, and an acid sensitive - IX/8/16) when LA at 2% was applied onto fresh beef surface.

In our study, when simulating a rework procedure of subprimals after a HEP (treating with antimicrobials and re-vacuum packaging the meat), PAA did not significantly reduce bacteria loads (for 30 min, CI=2.07<sup>A</sup> vs PAA=1.80<sup>A</sup>; and for 6 h, CI=2.37<sup>A</sup> vs PAA=2.30<sup>A</sup> Log CFU/cm<sup>2</sup>). No reduction was also observed when samples were treated with LA for 30 min under vacuum conditions (CI=2.07<sup>A</sup> vs LA=2.12<sup>A</sup> Log CFU/cm<sup>2</sup>). However, when lysing time was extended to 6 h, LA led to a significant 0.65 Log CFU/cm<sup>2</sup> reduction in *E. coli* counts. Overall, *E. coli* O157:H7 strains have been reported to be able to self-adapt in acid conditions, including two strains tested in this research (Tsai & Ingham, 1997). Although Pittman et al. (2012) showed a large and significant reduction (1.6 Log) of one strain (ATCC 43895) combined with other strains in a cocktail, our results showed a small reduction when compared to bacteriophage applications. When the bacteriophage solution was applied, optimal *E. coli* reductions were observed when compared to both LA and PAA (for 30 min, CI=2.07<sup>A</sup> vs BA=1.00<sup>B</sup>; and for 6 h, CI=2.37<sup>A</sup> vs BA=0.96<sup>C</sup>).

Under aerobic conditions, lysing time did not affect *E. coli* counts. Although both organic acids significantly lowered contamination, the lowest counts were observed when beef was treated with PGE (CI=2.70<sup>A</sup>, LA=2.23<sup>B</sup>, PAA=2.17<sup>B</sup>, and BA=1.29<sup>C</sup>). Better activity of organic acids under aerobic conditions when compared to vacuum may be associated to different bacterial metabolisms. In vacuum conditions, *E. coli* metabolism

possibly shifts from aerobic to anaerobic due to low availability or absence of O<sub>2</sub> (Uden, Becker, Bongaerts, Schirawski & Six, 1994). Although vacuum packaging does not remove all O<sub>2</sub> from the package (Kelly, Cruz-Romero, Kerry & Papkovsky, 2018), *E. coli* strains possibly adopt a less effective metabolism alternating from aerobic to anaerobic when growing in low O<sub>2</sub> conditions as indicative of being a facultative anaerobe. It is known that anaerobic metabolism and fermentation is less efficient than aerobic metabolism (Wulffen, RecogNice-Team, Sawodny & Feuer, 2016), therefore, under aerobic conditions (NV), bacteria express a higher metabolic rate. As previously mentioned, the effects of both organic acids used in this research are directly dependent on bacterial metabolism, which in optimal aerobic conditions, could lead to a better bacterial load reduction when compared to the absence of antimicrobial effects observed in vacuum-packaged samples.

Optimal PGE effects on *E. coli* loads both under vacuum and in aerobic conditions were possibly achieved due to its mode of action. Lytic *caudovirales* bacteriophages attach and inject viral DNA into bacteria, hijacking the host replication and translation mechanisms to produce phage parts, which assembled into capsids with genetic material and released from the infected cell through the activity of endolysins, peptidoglycan hydrolases the cell wall at the end of the cycle (Salmond & Fineran, 2015). Although bacteriophage applications against STEC in vacuum conditions has yet to be discussed in scientific reports, results of this research suggest that bacteriophage activity seems to be more effective than organic acids, even when *E. coli* adopts a less efficient metabolism when low O<sub>2</sub> concentrations are available. When compared to

organic acids, bacteriophage also depends on bacterial metabolism to perform its lytic cycle. The bacteriophage's better efficiency in lysing host cells is possibly attributed to independent molecular mechanisms that initiate viral DNA replication. Weigel and Seitz (2005) suggest that *E. coli* bacteriophages are able to regenerate closed ends of bacterial DNA to generate the phage genome and consequently produce phage structures. Previous research has suggested that *E. coli* strains are still able to replicate during cold storage under vacuum atmosphere (Dykes, Moorhead & Roberts 2001). Therefore, even with low metabolic rates, the active and independent bacteriophage mode of action is still able to induce lysis of the host, differently than organic acids, which possibly depend on higher bacterial metabolic rates to induce bacteria cell death.

The efficacy of bacteriophage applications in lowering *Enterobacteriaceae* in meats has been demonstrated by several authors in different conditions (Duc, Son, Honjoh & Miyamoto, 2018; Hudson et al., 2013; Hungaro, Mendonça, Gouvêa, Vanetti & Pinto, 2013; Liu et al., 2015; Stratakos & Grant, 2018; Sukumaran, Nannapanenni, Kiess & Sharma, 2015; Thung et al., 2017; Tomat, Casabonne, Aquili, Balagué & Quiberoni, 2018; Yeh et al., 2017; Yeh, de Moura, Van Den Broek & de Mello, 2018; Zinno, Devirgillis, Ercolini, Ongeng & Mauriello 2014), but to our knowledge, no studies evaluating the effects of bacteriophage activity on STEC under vacuum conditions have been published. Results of this research suggest that bacteriophage applications targeting STEC contamination in beef is more efficient either under aerobic or when meat is vacuum packaged compared to commonly used organic acids.

## Conclusion

Bacteriophage applications lead to an optimal reduction of *E. coli* O157:H7 (including *stx 1* and *stx2* gene-positives) in contaminated beef when compared to organic acid applications. Lactic and peroxyacetic acids did not reduce *E. coli* counts under vacuum conditions and minimally decreased contamination under aerobic atmosphere. Bacteriophage applications are more effective than organic acids when decontaminating beef under aerobic conditions as it is performed during subprimal and trimming fabrication for example. When reworking subprimals associated to STEC-positive trimmings after a High Event Period, beef packing plants must consider using bacteriophage solutions to eliminate possible *E. coli* contamination in vacuum-packaged beef.

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Table 1. Killing efficacy of PhageGuard E™ (Bacteriophages EP75 and EP335) for four *E. coli* O157:H7 strains.

Strain	Bacteriophage application	Average CFU (4 plates)	Killing efficiency (%)
ATCC® 35150™	<sup>a</sup> Control	83.5	98.2
	2 × 10 <sup>11</sup> PFU/ml	1.5	
ATCC® 43895™	Control	181	97.2
	2 × 10 <sup>11</sup> PFU/ml	5	
ATCC® 43894™	Control	407	96.7
	2 × 10 <sup>11</sup> PFU/ml	13.5	
NCTC 13128	Control	103.5	98.3
	2 × 10 <sup>11</sup> PFU/ml	1.75	

<sup>a</sup> No bacteriophage applied

Table 2. Effects of the application of organic acids and bacteriophage solutions under vacuum (V) and aerobic conditions (non-vacuum, NV) on beef contaminated with 4 *E. coli* strains (ATCC® 35,150™, ATCC® 43,895™, ATCC® 43,894™, and NCTC 13128). Log CFU/cm<sup>2</sup>.

Nested model	Treatment time	Antimicrobial treatment*				Average***
		CI	LA	PAA	PGE	
Vacuum (V)	30 min	2.07A	2.12Aa	1.80Ab	1.00B	1.75
	6 hr	2.37A	1.72Bb	2.30Aa	0.96C	1.84
	Average***	2.23A	1.92AB	2.06B	0.68C	SEM = 0.13
Non-vacuum (NV)	30 min	2.68	2.17	2.14	1.26	2.07
	6 hr	2.71	2.29	2.21	1.31	2.13
	Average***	2.70A	2.23B	2.17B	1.29C	SEM = 0.11

A,B,C Means in the same row having different superscripts are significant at  $P = .0061$  for V and  $P < .0001$  for NV. a,b Means in the same column within V having different superscripts are significant at  $P < .0061$ . SEM = Standard error of the mean.

\* CI = Control Inoculated, LA = Lactic acid at 5%, PAA = Peroxyacetic acid at 400 ppm, and PGE = PhageGuard E™ at  $1 \times 10^8$  PFU/ml.

\*\* Average of means within treatment time ( $P = .432$ ).

\*\*\* Average of means within antimicrobial treatment ( $P < .0001$ ).

#### Appendix V. Measuring Optical Density (OD<sub>600</sub>) and diluting inoculum

1. Turn on the spectrophotometer and set the wavelength to measure 600 nm (OD<sub>600</sub>).
2. Place a cuvette with LB broth in the placeholder, close the lid, and zero out the absorbance. This will be the baseline for the inoculum.
3. Insert a cuvette containing the overnight inoculum, shut the lid, and record the absorbance.
4. Convert the OD<sub>600</sub> absorbance into CFU/mL (OD of 1.0 =  $8 \times 10^8$  CFU/mL).
  - a. Chem.agilent *E. coli* cell calculator is available online
5. Calculate the dilution from the measured overnight culture to the desired concentration (CFU/mL) using  $C_1V_1=C_2V_2$ .
6. Using sterile BPW, dilute the bacterial inoculum.

## CHAPTER THREE

### **Effects of MS Bacteriophages, Ultraviolet light, and Organic Acid Applications on Beef Surface Contaminated with STEC O157:H7 and the 'Big Six' Serotypes after a Simulated High Event Period Scenario**

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**Abstract**

A high event period (HEP) occurs when beef processing facilities experience an elevated rate of Shiga Toxin-producing *E. coli* (STEC) positive trim samples. In order to avoid trim-associated vacuum-packaged primals and subprimals entering into commerce, beef that may be contaminated are treated with antimicrobials, re-packaged, and re-tested for STEC to ensure product wholesomeness. In this study, we evaluated the efficacy of Bacteriophages (P), Peroxiacetic Acid (PAA), Acidified Sodium Chlorite (ASC), and Ultraviolet light (UV) to reduce STEC O157:H7 and the Big Six populations in beef, after a simulated HEP scenario. In vacuum conditions, phage applications led to the greatest STEC reduction ( $P < 0.0001$ ). Under aerobic conditions, a combination of P + UV led to the lowest STEC loads ( $P < 0.0001$ ). Overall, treatments including bacteriophage provided best reductions when compared to non-phage treatments ( $P < 0.0001$  and  $P < 0.0001$ , under vacuum and aerobic conditions, respectively). Bacteriophage solutions provided improved control of STEC O157:H7 and the Big Six serotypes when reworking product after a simulated HEP scenario.

**Keywords:** Acidified sodium chlorite, Bacteriophage, *E. coli* O157:H7, high event period, PAA ,STEC

## Introduction

In the United States, the Food Safety and Inspection Service (FSIS) declared *E. coli* serotype O157:H7 adulterant in raw non-intact beef products in 1994. In 2012, six more serotypes (O26, O45, O103, O111, O121, O145) known as the “Big Six” were also added in the adulterant list (USDA-FSIS, 2012b). Regulatory guidelines mandate sampling of beef trim products and that no STEC positive raw beef products can be sold into commerce (USDA-FSIS, 2012a, 2014). The implementation of in-plant applications has demonstrated to be efficacious in decontaminating carcasses and reducing the presence of foodborne pathogens (Carlson et al., 2008; Castillo, Lucia, Goodson, Savell, & Acuff, 1998, 1999; Castillo, Lucia, Kemp, & Acuff, 1999; Castillo et al., 2001; Harris, Brashears, Garmyn, Brooks, & Miller, 2012; Harris, Miller, Loneragan, & Brashears, 2006; Marriott, Schilling, & Gravani, 2018a; Wheeler, Kalchayanand, & Bosilevac, 2014). However, studies have demonstrated that commonly used antimicrobial interventions are have decreased efficacy due to the development of acid resistant bacteria (Berry & Cutter, 2000; Dickson & Kunduru, 1995; Garren et al., 1998; Lin et al., 1996; Stopforth, Skandamis, Geornaras, & Sofos, 2007; Stopforth et al., 2004; Youssef, Yang, Badoni, & Gill, 2013). When a beef processing facility experiences a greater than 5% rate of STEC positive beef trim samples in a single time period, it is known as a high event period (HEP) (USDA-FSIS, 2014). The occurrence of a HEP indicates either a localized, out-of-control event or a systemic breakdown of preventative slaughter procedures that has led to contamination of beef products including primals, subprimals, and trimmings. The FSIS recommends to rework of

primals (Chuck, Brisket, Shank, Rib, Short Plate, Loin, Flank, Round) and subprimals (cuts derived from primals) by applying additional antimicrobial treatments to ensure no adulterated product is released into commerce (USDA-FSIS, 2014). Reworking procedures of subprimals occurs by first removing them from vacuum packaging and treating them with antimicrobials before repackaging. Subsequently, an aliquot of all repackaged products is tested for STEC, relying on antimicrobial effects under vacuum conditions. This process has been estimated to take approximately 30 minutes to 6 hours to complete depending on processing speed and production volume (Shebs, Lukov, Giotto, Torres, & de Mello, 2020).

In the United States, organic acid solutions and hot water are the most commonly used antimicrobial interventions in harvesting and processing plants. Acidic solutions such as Peroxyacetic acid disrupt chemiosmotic functions and interfere with the host cell membrane thereby affecting cell biochemical regulation and energy metabolism (Baldry & Fraser, 1988; Leaper, 1984; Liberti & Notarnicola, 1999). The efficacy of these chemical interventions has been demonstrated with high reductions (Harris et al., 2006; Laury et al., 2009; Zhao, Zhao, & Doyle, 2009), but also has been under scrutiny with decreased efficacy against foodborne pathogens (Bosilevac, Nou, Barkocy-Gallagher, Arthur, & Koohmaraie, 2006; King et al., 2005; Yeh, de Moura, Van Den Broek, & de Mello, 2018). These interventions are advantageous due to their cost, availability, and simple application (Mataragas et al., 2008). However, their potential decrease in

efficacy has been demonstrated with the development of acid resistant bacteria due to its exposure to sub-lethal applications (Bearson, Bearson, & Foster, 1997; Brown, Ross, McMeekin, & Nichols, 1997; Capita, Riesco-Peláez, Alonso-Hernando, & Alonso-Calleja, 2014; Cheng, Yu, & Chou, 2003; Foster & Hall, 1990; Goodson & Rowbury, 1989; Leyer, Wang, & Johnson, 1995; Wesche, Gurtler, Marks, & Ryser, 2009). Although organic acids are commonly applied during beef during processing, there are still occurrences of STEC outbreaks demonstrating a need for improved interventions. In this study, we evaluated individual and combined effects of two commonly used organic acids (Peroxyacetic acid and Acidified Sodium Chlorite), UV light, and a 7-phage novel bacteriophage cocktail on beef contaminated with STEC O157:H7 and the “Big Six” in both aerobic and vacuum conditions, simulating rework after a HEP.

## **Materials and methods**

### ***E. coli* Strains and inoculum preparation**

The seven pure STEC strains that were used in this study included the *E. coli* O157:H7 (ATCC® 35150™), O26 (ATCC® BAA-2196™), O45 (ATCC® BAA-2193™), O103 (ATCC® BAA-2215™), O111 (ATCC® BAA-2440™), O121 (ATCC® BAA-2219™), and O145 (ATCC® BAA-2192™). The frozen glycerol stock of each strain was streaked onto 1.6% agar LB agar plates in duplicate and incubated for 24 h at 37°C to ensure cultures are live and viable. A single isolated colony from each strain was transferred into independent sterile tubes containing 10 mL sterile LB broth and incubated at 37°C overnight with shaking (200 rpm). To prepare the inoculum, 1 mL of overnight pure

culture was transferred into 20 mL of LB broth and cultured overnight at 37°C with shaking. Subsequently, absorbance was measured at OD<sub>600</sub> to calculate concentration (CFU/ml) by using the Agilent *E. coli* cell culture biocalculator (Agilent, 2019). Each culture was diluted to 1x10<sup>8</sup> CFU/ml with sterile 0.1% Buffered Peptone Water (BPW) and then combined in equal amounts to create the final inoculum.

### **Phage preparation and stock titer**

Stocks of bacteriophages Mello-Shebs 1 (MS1) 026, MS1 045, MS1 103, MS1 111, MS1 121, MS1 145, and MS1 157 were obtained directly from the University of Nevada, Reno - Food Safety Microbiology Laboratory Library.

Phage stock concentration was determined by the double-layer agar methodology described by Adams (1959) and modified by Yeh et al. (2017). In short, 100 µL of 10-fold diluted phage and 100 µL of Log-phase *E. coli* culture was added into 4 mL of molten LB agar (0.6% agar) tempered at 45°C. Solution was mixed by inverting twice then evenly distributed onto hard LB agar (1.6%) plates until solidified. Duplicated plates were incubated at room temperature overnight for plaque enumeration. Concentration of phage solutions were determined to be 10<sup>8</sup> PFU/mL.

### **Sample preparation and treatment design**

Fresh beef rose meat (*m. Cutaneous trunci*, IMPS 194) (USDA-AMS, 2014) was obtained from a USDA inspected facility and transported under refrigeration (4°C) to the University of Nevada, Reno's Meat Quality Laboratory. The whole batch was screened for generic *E. coli* using 3M™ Petrifilm™ *E. coli*/Coliform count plates. Meat samples were fabricated into 10x10 cm<sup>2</sup> (100 cm<sup>2</sup>) pieces (n=154) and randomly assigned to a Completely Randomized Design using the following model:  $Y_{ij} = \mu + \alpha_i + \epsilon_{ij}$ .  $Y_{ij}$  was STEC count,  $\mu$  was the grand mean across the treatments, and  $\alpha_i$  was the effect of treatment from the grand mean specific to the 11 treatment levels. Model errors ( $\epsilon_{ij}$ ) were assumed to be random and independent. Treatments included: Control treated with BPW (C), Peroxyacetic acid (PAA, 400 ppm), Acidified Sodium Chlorite (ASC, 1200 ppm), Ultraviolet light - C (UV, 254 nm, 23°C for 30 s), a cocktail of all 7 MS1 Bacteriophages (P, 10<sup>8</sup> PFU/ml), PAA+ASC, PAA+UV, ASC+UV, P+PAA, P+ASC, and P+UV. The experiment was conducted separately treating samples under vacuum (n=77) and aerobic conditions (n=77).

### **Sample inoculation, antimicrobial applications, and bacterial enumeration**

Beef samples were inoculated with a cocktail containing all seven STEC strains. The concentration of the inoculum was  $1 \times 10^8$  CFU/ml, to yield approximately 3 Log CFU/cm<sup>2</sup>. All samples were kept under refrigeration prior to inoculation and during the trial ( $5 \pm 2^\circ\text{C}$ ). A volume of 500  $\mu\text{L}$  of the inoculum was uniformly pipetted on meat and distributed throughout the whole surface area of the sample using a sterile plastic rod, samples were then wrapped with oxygen permeable film (aerobic conditions) or vacuum

packaged (model VMC100P, VacPak-It, Lititz, PA, United States) and stored at 5°C for 30 min for bacterial attachment.

Subsequently, samples were removed from packages and treated with 500 µL of the antimicrobial solutions. Ultraviolet light (UV) was applied 2.5 cm above the sample surface by using a wand containing a single emitter. For combined treatments, UV was always applied first than PAA, ASC, and P. For treatments containing P, PAA and ASC were applied first. After treatments were applied, samples were re-wrapped or re-vacuumed. Re-vacuumed samples were used to simulate rework of contaminated subprimals after a HEP. Samples were then kept under refrigeration (5°C). After one hour, samples were removed from their package and swabbed (QS1000, Hygiena, Camarillo, CA, United States) for bacteria enumeration. Swabs were homogenized and the homogenate was vortexed, serially diluted, and spread-plated onto LB agar plates in duplicate. Plates were incubated inverted overnight at 37 °C, and *E. coli* colonies were enumerated (CFU/cm<sup>2</sup>). In order to simulate an industry scenario, sample homogenates were not treated to neutralize possible residual effects of organic acids or phages. Antimicrobial neutralization procedures are not practiced by laboratories that provide service to meat processing facilities.

### **Color and lipid stability after UV light exposure**

In order to evaluate the effects of UV treatment on color and lipid stability of beef samples, a separate trial using a different set of samples was

performed. A total of 10 samples were randomly assigned to a Control (n=5) and to the same UV treatment applied on contaminated samples (n=5). Instrumental color was recorded for L\* (lightness; black = 0, white = 100), a\* (redness; red = positive values; green = negative values), b\* (yellowness; yellow = positive values; blue = negative values), hue angle ( $\text{hue} = \arctan(b^*/a^*)$ ), and chroma values [ $\text{chroma} = (a^{*2} + b^{*2})^{1/2}$ ] using a Minolta chromameter (CM-700D; Minolta Inc., Osaka, Japan) with an 8 mm-diameter after 0, 1, 12, and 24 h after UV light exposure with sample measurements averaged from three locations on each sample. Lipid stability was measured 24 h after UV light exposure by performing the Thiobarbituric Acid assay described by Buege & Aust (1978) modified by Ahn et al. (1998). Briefly, 5 g of pulverized sample and 14 mL of deionized water and 1 mL of 10% Butylated hydroxyanisole (BHA, 10g of BHA diluted in 90 mL of Alcohol 90% and 5 mL of distilled water) were mixed, homogenized and centrifuged at 200 rpm for 5 min. Subsequently, 1 mL of the homogenate was mixed with 2-Thiobarbituric Acid/Trichloroacetic Acid (TBA/TCA) and incubated at 70 °C for 30 min, prior to centrifugation at 2000 rpm for 15 min. Two-hundred (200)  $\mu\text{L}$  aliquots used to measure absorbance at 540 nm (Synergy HT, Biotek, USA). Lipid oxidation (TBA values) were expressed as malonaldehyde concentration (mg/Kg) and quantification was determined by comparing samples with solutions of known concentrations of TEP (1,1,3,3-tetraethoxypropane) in distilled water ranging from 0 to 50 nmol/mL.

### **Statistical analysis**

The experiment was replicated 7 times with 1 repetition per treatment (n=154 total e.u., 77 e.u. for aerobic and 77 e.u. for vacuum; n=7 e.u. per fixed effect). Data obtained to evaluate the effects of UV application on color was analyzed as a repeated measures. The smallest Akaike and Bayesian information criteria indicated that Compound Symmetry was the best covariance matrix for color analysis. Data were analyzed using the GLIMMIX procedure of SAS® 9.3 package (SAS Institute, Inc., USA). When significance ( $P \leq 0.05$ ) was detected for the fixed effect, means separations were performed by using the LSMEANS and DIFF functions. For contrasts, significance was detected by using the LSMESTIMATE function statements.

## Results

Under vacuum conditions, UV only, PAA+ASC, PAA, and PAA+UV did not decrease STEC loads. Applications of the MS1 bacteriophage cocktail (P) and combined treatments of P+PAA, P+UV, P+ASC, and ASC+UV provided optimal bacteria reduction when compared to UV, C, PAA+ASC, PAA, PAA+UV, and ASC ( $P < 0.0001$ , Figure 1). Samples treated with ASC+UV showed values statistically similar to treatments that provided the optimal reduction, but overall P only led to the highest numeric reduction (1.04 Log CFU/cm<sup>2</sup>). Contrasts were evaluated for: all treatments containing P (P, P + PAA, P+UV, and P+ASC) versus all other treatments (UV, PAA, ASC, PAA+ASC, PAA+UV, and ASC+UV), all treatments containing P versus control (C) and all other treatments versus control (C) (Table

1). Under vacuum, all treatments containing P determined the best reduction when compared to other treatments ( $P < 0.001$ ). As observed in Figure 1, the contrast analysis confirmed that treatments containing P significantly reduced STEC loads when compared to control ( $P = 0.0002$ ), however, when evaluating the effects of all other treatments versus C, no difference was observed ( $P = 0.3395$ ).

Under aerobic conditions, combined treatments of P+PAA and P+UV were significantly more efficient than all other treatments (Figure 2). Overall, P+UV provided the greatest STEC reduction ( $1.33 \text{ Log CFU/cm}^2$ ) whereas UV, ASC, and PAA+ASC did not reduce bacteria populations. When looking at the contrasts analyses (Table 1), results for treatments applied in aerobic conditions were similar than results reported for vacuum. Treatments containing P provided improved reduction on STEC loads when compared to all other treatments and control. When comparing the effects of all other treatments against C, no statistical difference was observed ( $P = 0.0781$ ).

The effects of UV on color attributes and lipid stability are shown in Table 2. No detrimental effect of UV light was observed for lipid stability. Treated samples showed lower MDA values when compared to control. Regarding color, no interaction between both fixed effects of time and treatment were observed. However, samples were yellower at 12 h when compared to 0 and 24h ( $P = 0.0083$ ) and less red (lower  $a^*$  values) ( $P = 0.0003$ ) and had lower Chroma values after 24 h ( $P = 0.0012$ ). Values of Hue angle gradually increased from 0 to 24 h ( $P < 0.0001$ ). Samples treated with UV were darker ( $P = 0.0330$ ), redder ( $P = 0.0011$ ), had a smaller hue angle ( $P = 0.0003$ ) and higher Chroma values ( $P = 0.0059$ ).

## Discussion

Overall, the 7-phage cocktail and its treatment combinations demonstrated optimal reduction of STEC O157:H7 and the “Big Six” serotypes on beef surface, under both aerobic and vacuum conditions. A bacteriophage cocktail comprising seven phages targeting individual STEC serotypes, to our knowledge has not been reported for food application. We have demonstrated previously that each phage is able to lyse each serotype of STEC in vitro by at least 96.2% (Shebs, Yeh-Parker, Torres, & de Mello, 2019) which can be in part due to the ability of the phages to bind to LPS or OmpC surface receptors that are likely similar among the *E. coli* serotypes or other *Enterobacteriaceae*, enabling them to attach to more than one host species (Montag, Hashemolhosseini, & Henning, 1990; Singh, Singh, Williams, Jones, & Abdullah, 1995; Singh, Upshaw, Abdullah, Singh, & Klebba, 1992; Yu & Mizushima, 1982). Thus, moving between different hosts would allow acquisition of different genetic material through transduction that could widen the phages’ host range to be effective against multiple foodborne pathogens and to discourage resistance against phage lysis (Fineran et al., 2009; Oechslin, 2018).

To our knowledge, the use of a phage cocktail solution that targets all seven adulterant STEC serotypes has not been demonstrated beyond six phages. Also, such cocktails have not been compared to commonly used antimicrobial solutions such as PAA or ASC. Processors segregate primals and subprimals that

are associated with STEC positive trim lots. The products are subsequently tested for STEC and are often treated with an antimicrobial such as an organic acid before being repackaged and tested for STEC later to ensure no adulterated products are released into commerce. Products that are not decontaminated must be further processed by being sold to a cooker for a reduced price or recalled which reduces the profitability for the company and industry. In the United States, weak acids and oxidizing agents are commonly used to decontaminate carcasses and subprimals. Peroxyacetic acid (PAA) and acidified sodium chlorite (ASC) are both oxidizing agents that are approved for use on beef, poultry parts, and carcasses while UV-C light has yet to be approved for use on meat and poultry products (USDA-FSIS, 2019). These antimicrobials have been reported to be efficacious against bacterial pathogens (Harris et al., 2006; Laury et al., 2009; Penney et al., 2007; Ransom et al., 2003), while others have found little or inconsistent efficacy (Ellebracht et al., 2005; Gill & Badoni, 2004; King et al., 2005; Shebs, Lukov, Giotto, Torres, & de Mello, 2020; Stivarius, Pohlman, Mcelyea, & Waldroup, 2002; Stopforth et al., 2004; Uyttendaele et al., 2001; Yeh et al., 2018). Weak acids such as lactic acid (LA), acetic acid, or citric acid act by diffusing across the cell membrane and dissociating in the cytoplasm therefore lowering the pH and causing a disruption in the bacterial membrane and inhibition of mechanisms that control pH homeostasis. This disruption causes the bacteria to synthesize acid stress proteins to avoid acid-induced damage and to maintain optimal cytoplasmic pH levels (Bearson, Bearson, & Foster, 1997; Desriac et al., 2013; Foster, 1995; Salmond, Kroll, & Booth, 1984; Nair, Upadhyaya, Amalaradjou, & Venkitanarayanan, 2017). PAA is the combination of acetic

acid and hydrogen peroxide that creates peracetic acid or Peroxyacetic acid in equilibrium. PAA releases active oxygen in the form of superoxide radicals that oxidizes susceptible proteins that disrupt the chemiosmotic functions of membrane transport by disrupting the cell wall (Baldry & Fraser, 1988; Block, 2001; Fraser, Godfree, & Jones, 1984; Liberti & Notarnicola, 1999; Nair et al., 2017) ASC is the combination of sodium chlorite and an acid, most commonly citric acid. This combination produces an active chlorine dioxide or oxy-chlorous compound that can oxidize cell membrane surfaces (Gordon, Kieffer, & Rosenblatt, 1972; Kross, 1984; Thiessen, Usborne, & Orr, 1984; Villarreal, Baker, & Regenstein, 1990). Chlorine dioxide has been reported to have a much better antimicrobial effect compared to both chlorine water and hypochlorous acid (Benarde et al., 1965; Lillard, 1979).

The efficacy of PAA has been evaluated against many pathogens in different foods. PAA (0.01%) has been shown to be effective against *E. coli* after 10 minutes yielding over 5 Log CFU/ml reduction in solution (Briñez, Roig-Sagués, Hernández Herrero, López-Pedemonte, & Guamis, 2006), however it has not demonstrated to be as effective when applied to meat (Purnell, James, James, Howell, & Corry, 2014; Shebs et al., 2020; Stopforth et al., 2004; Yeh et al., 2018). On fresh beef carcass surface, 0.02% PAA resulted in a 3.32 Log CFU/cm<sup>2</sup> reduction of *E. coli* O157:H7 after 10 h of spray chilling, but there was no reduction after 24 or 48 h (Stopforth et al., 2004). On fresh beef surface, immersion in 0.02% ASC or 0.02% PAA reduced a 5-strain *E. coli* cocktail by 1.9

and 1.4 Log CFU/cm<sup>2</sup>, respectively, although this was overshadowed by 0.5% cetylpyridinium chloride (CPC) which gave the optimal reduction of 4.8 Log CFU/cm<sup>2</sup> (Ransom et al., 2003). Bauermeister, Bowers, Townsend, & McKee (2008) demonstrated that a minimum PAA concentration of 0.0025% is an effective antimicrobial against *Salmonella typhimurium* on chicken, and higher concentrations of 0.015% and 0.02% showed an extended shelf life but had an off color and off odor. When Alonso-Hernando, Alonso-Calleja, & Capita (2013) compared the use of multiple antimicrobials against multiple pathogens on skinless chicken legs, they observed that ASC (1200 ppm) gave the best *E. coli* reduction (1.94 Log CFU/cm<sup>2</sup>) followed by 2% citric acid (1.47 Log CFU/cm<sup>2</sup>), while peroxyacids (220 ppm) did not result in any significant reduction. They also observed no significant decrease in *Salmonella enterica Enteritidis* with the use of peroxyacids. Similarly, King et al. (2005) did not observe any reduction in microbial counts when PAA was applied at 600 ppm or below at either 45°C or 55°C, but PAA of 1000 ppm reduced both *E. coli* O157:H7 and *Salmonella typhimurium* by up to 1.7 and 1.3 Log CFU/cm<sup>2</sup>, respectively. However, these reductions were overshadowed by 4% lactic acid (2.7 and 3.4 Log<sub>10</sub> CFU/ cm<sup>2</sup>, respectively). When tested in ground beef, PAA (0.02%) yielded the lowest *E. coli* counts after seven days of retail display, while ASC (0.1%) resulted in the lowest APC counts (Pohlman et al., 2007). PAA (180 ppm) has also been demonstrated by Brightwell, Clemens, Adam, Urlich, & Boerema (2009) to be efficacious against vacuum packaged beef microflora, extending the shelf life. In our study, we did not observe a significant ( $P > 0.05$ ) difference between samples treated with PAA and control samples under vacuum conditions, but we did observe a

significant ( $P < 0.05$ ) reduction under aerobic conditions. Under vacuum conditions, the only significant reduction by PAA was in combination with bacteriophage (1.09 Log CFU/cm<sup>2</sup>). Under aerobic conditions, PAA applied alone yielded a significant 0.48 Log CFU/cm<sup>2</sup> reduction and when combined with phage yielded a significant 1.07 Log CFU/cm<sup>2</sup> reduction.

Citric acid activated acidified sodium chlorite (ASC) has been well reported as an antimicrobial used against *E. coli* and *Salmonella* on meat with varying effectiveness. Ransom et al. (2003) immersed fresh beef contaminated with an *E. coli* O157:H7 cocktail into 0.02% ASC for 30 seconds and observed a significant 1.9 Log CFU/cm<sup>2</sup> reduction. Similarly, Kalchayanand et al. (2012) observed significant ( $P < 0.05$ ) reductions of an STEC cocktail on fresh beef flanks when they sprayed 1000 ppm ASC (0.4-2.0 Log CFU/cm<sup>2</sup>), but hot water (85°C) gave greater reductions (2.4-4.2 Log CFU/cm<sup>2</sup>). Stopforth et al. (2004) applied ASC (0.12%) onto pre-chilled carcasses against acid habituated and non-acid habituated *E. coli* O157:H7 cells and observed a 2.1 and 2.4 Log CFU/cm<sup>2</sup> decrease after 48 hours of spray chilling, respectively, although CPC application yielded undetectable levels of *E. coli* cells. When Visvalingam & Holley (2018) evaluated multiple concentrations of ASC on contaminated beef surface prior to grinding, they observed no significant reduction of *E. coli* at 30 and 100 ppm, and only 200 and 400 ppm maintained a significant decrease over 4 days in vacuumed ground beef. Similarly, ASC (1000 ppm) applied onto beef trim prior to grinding and refrigerated storage yielded a 0.5 Log CFU/g reduction for *E. coli*

after 6 h with no change at 24 h (Harris et al., 2012). Pohlman et al. (2007) evaluated *E. coli* reduction in ground beef over 7 days and observed ASC (0.1%) yielded the best reduction of APC after 7 days storage (2.26 Log CFU/g), but PAA gave the highest reduction of *E. coli* (1.39 Log CFU/g). In our study, we did not observe a significant ( $P > 0.05$ ) difference between control samples and ASC treated samples under both aerobic and vacuum conditions. When ASC was applied in combination with either UV-C light or with bacteriophage, then we observed a significant reduction ( $P < 0.05$ ) compared to the control under both packaging conditions.

The application of UV light has been approved for use by the Food and Drug Administration (FDA) to decontaminate food, food contact surfaces, juices, and water because of its non-thermal and environmentally friendly nature (Bintsis et al., 2000; FDA, 2000; Sommers et al., 2010). UV light applied at 250-270 nm is considered germicidal UV-C with 254 nm the most common wavelength used for disinfection (Bintsis et al., 2000; Guerrero-Beltrán & Barbosa-Cánovas, 2004). The antimicrobial ability of UV-C light is based on the ability for the light to be absorbed by the DNA of microorganisms to cause thiamine and cytosine cross-linking which cannot be replicated, leading to cell death (Escalona et al., 2010; Guerrero-Beltrán & Barbosa-Cánovas, 2004; Hijnen et al., 2006; Sastry et al., 2000; Snowball & Hornsey, 1988). When applied onto alfalfa seeds, Sharma & Demirci (2003) reported a 4.89 Log CFU/g reduction of *E. coli* O157:H7 with pulsed UV light was applied for 1 min at 8 cm distance from the emitters. Similarly, Ozer & Demirci (2006) applied UV light for 1 min at 8 cm distance onto salmon fillets, but reported a maximum 1.09 Log CFU/g reduction of *E.*

*coli* O157:H7 and a maximum 0.86 Log CFU/g reduction of *Listeria monocytogenes* after 30 sec at 5 cm distance. Furthermore, the quality of the salmon fillets was visually altered by the UV light at close distances, indicating the ability of UV light to affect quality attributes of fresh meats and the importance of using a larger distance such as 8 cm to not affect quality attributes of products. When applied on goat meat, a combination of 1% lemongrass oil and UV-C light (10 cm distance for 2 min) reduced *E. coli* K12 to undetectable levels (6.66 Log CFU/mL reduction) compared to a maximum 1.18 Log CFU/ml reduction by UV-C light alone (Degala, Mahapatra, Demirci, & Kannan, 2018). When broad spectrum light was applied onto chicken breast against *Salmonella enterica* and *Listeria monocytogenes*, pulsed light (200-1000 nm) applied for 200 sec was shown to reduce these pathogens by 2 and 2.4 Log CFU/ml, respectively (Paskeviciute, Buchovec, & Luksiene, 2011). Similarly, chicken breast treated with a combination of UV-C light (40 min exposure time) and a bacteriophage solution ( $1 \times 10^8$  PFU/g) reduced a *Listeria monocytogenes* cocktail by 2.04 Log CFU/g within 1 h of treatment compared to a maximum 0.84 Log CFU/g and 1.58 CFU/g reductions with phage or UV-C light alone, respectively (Yang, Sadekuzzaman, & Ha, 2017). It has been demonstrated that a longer exposure time to UV-C light can result in a higher reduction of pathogens. As previously discussed, there is potential for UV-C light to affect quality attributes of meat products such as color and lipid oxidation, although careful application of UV-C light can circumvent such negative effects while offering efficient, non-thermal,

and non-chemical pathogen reduction. In our study, we observed a significant ( $P < 0.05$ ) reduction when combining phage and UV-C light treatments which gave 0.83 and 1.46 Log CFU/cm<sup>2</sup> reductions for vacuum and aerobic conditions, respectively. Overall, control samples had better color stability. Meat color deterioration is typically caused by oxidation of myoglobin leads to higher L\* and lower a\*, b\*, and Chroma values (King, Shackelford, Rodriguez, & Wheeler, 2011). When we evaluated the effects of UV light exposure on color and lipid oxidation, we observed no effects on lipid oxidation. Effect of time on color was expected since oxygenation of myoglobin is responsible for final color. Application of UV is known for causing photo-oxidation of myoglobin, however when applied in lower doses no detrimental effects on color or lipid stability are observed. In our study, samples treated with UV were darker, redder, had a smaller hue angle and higher Chroma values. Hue angle and Chroma values express color nuances and saturation whereas large angles and lower chroma values are associated to discoloration and browning. In this study, UV application improved color parameters. Although UV light exposure is associated to mechanisms that compromise color, results presented in this research showed no detrimental effects on color caused by UV light.

In this study, all antimicrobial combinations with phage gave significant ( $P < 0.0001$ ) STEC reductions under both aerobic and vacuum conditions compared to all other antimicrobial treatments without phage. The efficacy of adding phage to other antimicrobial treatments has been previously reported (Leverentz et al., 2003; Sukumaran, Nannapaneni, Kiess, & Sharma, 2015; Yeh et al., 2018). Yeh et al. (2018) combined a bacteriophage preparation with lactic acid (LA), PAA, and tumbling UV light

and observed that the combination of bacteriophage and UV light yielded the greatest reduction (1.97 Log CFU/g) of *Salmonella* in ground beef. In addition, the combined application of phage with chemical antimicrobials (Chlorine, PAA, CPC) has shown to give significant ( $P < 0.05$ ) reductions of *Salmonella* up to 2.5 Log CFU/cm<sup>2</sup> on chicken skin after 24 hours (Sukumaran et al., 2015). When comparing chemical antimicrobials to phage applications, Hungaro, Mendonça, Gouvêa, Vanetti, & Pinto (2013) demonstrated a 5-bacteriophage cocktail reduced *Salmonella enteritidis* on chicken skin with comparable efficacy to chemical antimicrobials, allowing them to be alternative interventions. Shebs et al. (2020) is the only study to our knowledge that evaluates the efficacy of phages under vacuum conditions compared to chemical antimicrobials, 5% LA and 400 ppm PAA. They observed significant reduction of *E. coli* O157:H7 under aerobic conditions and vacuum conditions with phage treatment, and no significant reduction with LA or PAA application under vacuum conditions.

This study simulates rework procedures to decontaminate STEC contaminated beef subprimals using a novel 7-bacteriophage cocktail, PAA, ASC, and UV-C light alone and in combinations. We observed the greatest reduction of STEC under vacuum conditions by P and P+PAA combined treatments, and under aerobic conditions by combined Phage and UV-C light treatments, while no significant reduction was observed in samples treated with ASC, UV, UV+PAA, PAA+ASC. Recently, commonly used chemical antimicrobials have been losing their efficacy that could be due to the development of acid resistance (Berry &

Cutter, 2000; Brackett, Hao, & Doyle, 1994; Diez-Gonzalez, Callaway, Kizoulis, & Russell, 1998; Hovde, Austin, Cloud, Williams, & Hunt, 1999). When bacteria are exposed to mild or sublethal levels of acidic solutions (5-6 pH), they are able to survive and synthesize acid resistance proteins that increases their survivability when re-exposed to lower pH solutions (2-4 pH) (Brudzinski & Harrison, 1998; Foster & Hall, 1990; Garren et al., 1998; Goodson & Rowbury, 1989; Leyer et al., 1995; Lin et al., 1996). The development of acid tolerant bacteria has become a food safety concern due to the most common antimicrobial interventions being acidic, requiring new and alternative antimicrobial solutions.

Bacteriophages have been studied for use as alternative food safety interventions because of their advantages including being non-thermal, non-oxidizing, and self-perpetuating (Goodridge & Abedon, 2003; Kazi & Annapure, 2016; Loc-Carrillo & Abedon, 2011; Oliveira et al., 2012). These characteristics are what make phages applicable for many applications, including HEP scenarios. Under such vacuum packaging conditions, the bacteria likely convert to anaerobic fermentation metabolism for energy production due to the reduction of O<sub>2</sub>. Although vacuum packaging does not remove all traces of O<sub>2</sub> (Kelly, Cruz-Romero, Kerry, & Papkovsky, 2018), this shift to a less energy efficient metabolism could interfere with the phage lytic cycle, although the reductions of phage only treatments observed in this study were similar for vacuum (1.02 Log cfu/cm<sup>2</sup>) and aerobic (0.95 Log cfu/cm<sup>2</sup>) packaging. Under vacuum conditions, minimal oxygen is left within the package causing the *E. coli* to use fermentation as a way to adapt and survive, which greatly reduces the metabolic activity (James & Keevil,

1999; Stolper, Revsbech, & Canfield, 2010; Uden, Becker, Bongaerts, Schirawski, & Six, 1994) although it does not inhibit growth in all cases (Dykes, Moorhead, & Roberts, 2001) meaning that the metabolism of the bacteria is able to allow phage replication and cell lysis under vacuum conditions.

Optimal reduction of a 7-strain *E. coli* cocktail under both vacuum and aerobic packaging conditions was achieved by treatments containing our novel 7-bacteriophage cocktail. This could be due to the unique mechanism of the lytic cycle compared to mechanisms of chemical antimicrobials previously mentioned. The lytic cycle involves the hijacking of host replication mechanisms including ribosomes that begin to translate viral mRNA into protein virion structures. Lytic bacteriophages depend on the metabolism of the bacterial host to replicate and subsequently lyse the host cell via endolysin and hydrolase activity (Salmond & Fineran, 2015). At refrigeration temperatures, the bacterial metabolism slows and reduces the rate of protein synthesis and cellular growth (Berry & Foegeding, 1997; Broeze, Solomon, & Pope, 1978; Das & Goldstein, 1968; Farewell & Neidhardt, 1998). Although there have been studies that demonstrated better phage killing efficiency at higher temperatures (Liu et al., 2015; Tomat et al., 2018) there are reports of phage treatments that give efficacious foodborne pathogen reduction at low and refrigeration temperatures (Guenther et al., 2012; Hooton et al., 2011; Hungaro et al., 2013; Sukumaran et al., 2015; Tomat et al., 2018). Many studies have demonstrated bacteriophage applications on meats (Bigwood, Hudson, Billington, Carey-Smith, & Heinemann,

2008; Duc, Son, Honjoh, & Miyamoto, 2018; Hudson et al., 2013; Hudson, Billington, Wilson, & On, 2015; Hungaro et al., 2013; Liu et al., 2015; Shebs et al., 2020; Stratakos & Grant, 2018; Sukumaran et al., 2015; Thung et al., 2017; Tomat et al., 2018; Yeh et al., 2018, 2017; Zinno, Devirgiliis, Ercolini, Ongeng, & Mauriello, 2014), but few have demonstrated efficiency under vacuum conditions apart from this study or Shebs et al. (2020). Additionally, there are no studies to our knowledge that evaluate a phage cocktail that targets each USDA adulterant STEC serotype in combination with other antimicrobials. Results of this research suggest that bacteriophage cocktails that target the seven adulterant STEC serotypes in meats have higher pathogen killing efficacy than organic acids or UV-C light alone and should be further evaluated for their application in both aerobic and vacuum packaging conditions.

## **Conclusion**

Bacteriophage application yielded the greatest reduction of all seven adulterant STEC on beef surface alone and in combinations with other antimicrobial treatments under simulated High Event Period protocols. Peroxyacetic acid and ASC treatments provided a limited decrease under both aerobic and vacuum conditions. The combination of PAA, ASC, and UV antimicrobial interventions with a 7-bacteriophage cocktail led to significant reductions of STEC under both vacuum and aerobic atmospheres compared to individual applications. During High Event Periods, reworking beef primals and subprimals to reduce STEC loads under vacuum packaged conditions

could become more effective with the addition of bacteriophages that target *E. coli* O157:H7 and the 'Big Six' to provide an additional barrier in robust food safety systems.

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Table 1. Significance of contrasts analysis for treatments applied under vacuum and aerobic conditions

<b>Contrasts</b>	<b>P - value</b>	
	<b>Vacuum</b>	<b>Aerobic</b>
All treatments containing P <sup>1</sup> versus all other treatments <sup>2</sup>	<0.0001	0.0001
All treatments containing P <sup>1</sup> versus C <sup>3</sup>	0.0002	<0.0001
All other treatments <sup>2</sup> versus C <sup>3</sup>	0.3395	0.0781

<sup>1</sup>Bacteriophage (P), P + Peroxyacetic acid (PAA), P + Ultraviolet light (UV), and P + Acidified Sodium Chlorite (ASC).

<sup>2</sup>UV, PAA, ASC, PAA+ASC, PAA+UV, and ASC+UV.

<sup>3</sup>Control (C) treated with Buffered Petpone water.

Table 2. Color and lipid stability of beef samples treated with UV light

Lipid stability									
Mg MDA/kg	Treatment <sup>1</sup>				SEM <sup>2</sup>	P-value			
	Control	UV	SEM <sup>2</sup>	P-value					
	1.42 <sup>A</sup>	0.99 <sup>B</sup>				0.037			

Color Parameters										
	Time (h)				Treatment <sup>1</sup>		SEM <sup>2</sup>	P - value		
	0	1	12	24	Control	UV		Time	Treat	Time*Treat
L*	49.51	47.17	46.97	47.91	49.04 <sup>A</sup>	46.74 <sup>B</sup>	0.7265	0.3048	0.0330	0.9775
a*	14.44 <sup>a</sup>	15.06 <sup>a</sup>	14.39 <sup>a</sup>	11.55 <sup>b</sup>	12.89 <sup>B</sup>	14.83 <sup>A</sup>	0.3784	0.0003	0.0011	0.5641
b*	6.95 <sup>b</sup>	7.73 <sup>ab</sup>	8.37 <sup>a</sup>	7.42 <sup>b</sup>	7.65	7.59	0.3124	0.0083	0.8267	0.1423
Hue	25.72 <sup>c</sup>	27.06 <sup>c</sup>	30.29 <sup>b</sup>	33.54 <sup>a</sup>	31.12 <sup>A</sup>	27.19 <sup>B</sup>	0.9624	<0.0001	0.0003	0.2065
Chroma	16.05 <sup>a</sup>	16.94 <sup>a</sup>	16.68 <sup>a</sup>	13.77 <sup>b</sup>	15.04 <sup>B</sup>	16.68 <sup>A</sup>	0.4213	0.0012	0.0059	0.4494

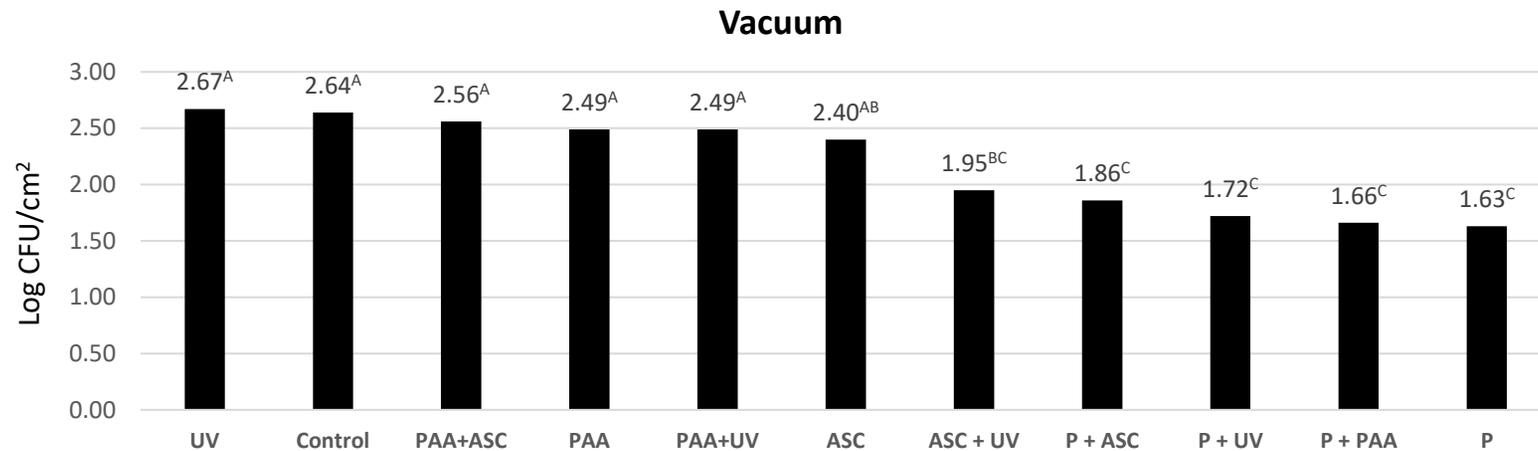
<sup>1</sup>Control=Not treated with UV light, UV= 254 nm, 23°C for 30 sec.

<sup>2</sup>Standard error for the mean for treatments.

<sup>A,B</sup>Different superscripts within treatment are significantly different.

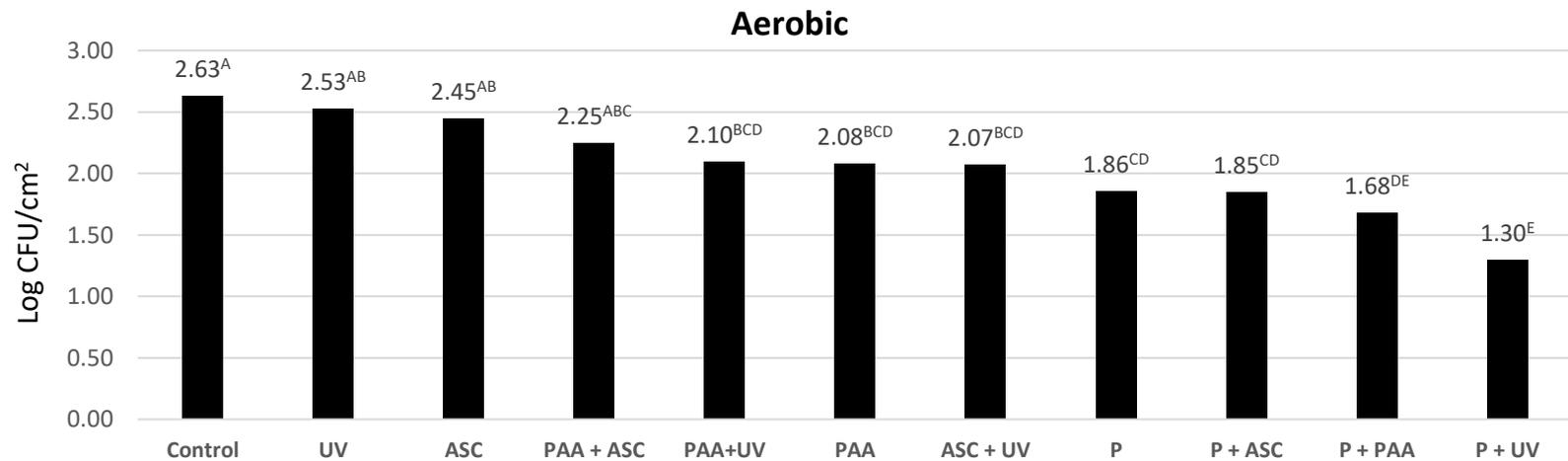
<sup>a,b,c</sup>Different superscripts within time are significantly different.

Figure 1. Least square means of STEC populations in beef samples (*m. Cutaneous trunci*) kept under vacuum and treated with BPW (C), Peroxyacetic acid (PAA, 400 ppm), Acidified Sodium Chlorite (ASC, 1200 ppm), Ultraviolet light - C (UV, 254 nm, 23°C for 30 s), MS1 Bacteriophages (P,  $10^8$  PFU/ml), PAA+ASC, PAA+UV, ASC+UV, P+PAA, P+ASC, and P+UV. Standard error of the mean = 0.2108.



<sup>A,B,C</sup> Means with different superscripts are significantly different at  $P < 0.001$ .

Figure 2. Least square means of STEC populations in beef samples (*m. Cutaneous trunci*) kept under aerobic conditions and treated with BPW (C), Peroxyacetic acid (PAA, 400 ppm), Acidified Sodium Chlorite (ASC, 1200 ppm), Ultraviolet light - C (UV, 254 nm, 23°C for 30 s), MS1 Bacteriophages (P, 10<sup>8</sup> PFU/ml), PAA+ASC, PAA+UV, ASC+UV, P+PAA, P+ASC, and P+UV. Standard error of the mean = 0.1919.



<sup>A,B,C,D,E</sup> Means with different superscripts are significantly different at P<0.001.

## Appendix VI. Phage amplification and concentration protocol

1. Inoculate 5x LB broth with fresh overnight culture at the ratio 1 mL culture to 5 mL 5x LB broth
2. Incubate at 37°C with shaking (230 RPM) for 1-3 hours
3. Inoculate 100  $\mu$ L of high titer ( $\geq 10^9$  PFU/mL) phage or 500  $\mu$ L of low titer phage ( $\leq 10^8$  PFU/mL)
4. Incubate at 37°C with shaking (230 RPM) overnight ( $18 \pm 6$  hours)
5. Centrifuge for 20 minutes at 5,000 x g
6. Filter liquid lysate with 0.2  $\mu$ M syringe filter into a new labeled conical tube
  - a. If propagating a large volume (>15 mL), filter supernatant with 0.45  $\mu$ M syringe filter followed by 0.2  $\mu$ M syringe filter
7. Transfer 12 mL into the upper reservoir of the 100 kDa Amicon filter (MilliporeSigma™ UFC910008)
8. Centrifuge at 4,000 x g for 9 min.
  - a. Do not spin the filter dry, centrifugation times will vary based on concentration
  - b. Do not exceed 12 mL if using a fixed angle rotor centrifuge
9. Remove the filtrate and add more phage lysate to upper reservoir and continue filtering through centrifugation until the phage lysate in the upper reservoir is less than 5 mL
10. Add 10 mL of SM Buffer into the upper reservoir to wash at 4,000 x g for 9 min. or until less than 10 mL of phage lysate
11. Perform the double agar assay as described by Adams (1959) and modified by Yeh et al. (2017) (Appendix IV)