

# Inactivation of Human Pathogens during Phase II Composting of Manure-Based Mushroom Growth Substrate

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## ABSTRACT

Commercial production of white button mushrooms (*Agaricus bisporus*) requires a specialized growth substrate prepared from composted agricultural by-products. Because horse and poultry manures are widely used in substrate formulations, there is a need to determine the extent to which the composting process is capable of eliminating human pathogens. In this study, partially composted substrate was inoculated with a pathogen cocktail (log 10<sup>6</sup> to 10<sup>8</sup> CFU/g) containing *Listeria monocytogenes*, *Escherichia coli* O157:H7, and *Salmonella*. Pathogen and indicator-organism reductions were followed at temperatures that typically occurred during a standard 6-day phase II pasteurization and conditioning procedure. Controlled-temperature water bath studies at 48.8, 54.4, and 60°C demonstrated complete destruction of the three pathogens after 36.0, 8.0, and 0.5 h, respectively. Destruction of *L. monocytogenes* and *E. coli* O157:H7 at 54.4°C occurred more slowly than *E. coli*, total coliforms, *Enterobacteriaceae*, and *Salmonella*. Microbial reductions that occurred during a standard 6-day phase II pasteurization and conditioning treatment were studied in a small-scale mushroom production research facility. After phase II composting, *E. coli*, coliforms, and *Enterobacteriaceae* were below detectable levels, and inoculated pathogens were not detected by direct plating or by enrichment. The results of this study show that a phase II composting process can be an effective control measure for eliminating risks associated with the use of composted animal manures during mushroom production. Growers are encouraged to validate and verify their own composting processes through periodic microbial testing for pathogens and to conduct studies to assure uniform distribution of substrate temperatures during phase II.

Increases in the number of foodborne disease outbreaks associated with fresh produce has been attributed, in part, to a shift in consumer preference toward fresh fruits and vegetables that are not subjected to high-temperature blanch or retort treatments (44). As this trend continues, there exists a need to reevaluate farm-to-table contamination risks under commercial growing and handling conditions.

Consumption statistics for mushrooms reflect overall fruit and vegetable trends. In 1970, only 28% of white button mushrooms (*Agaricus bisporus*) grown in the United States were marketed fresh; the remaining were sold as processed products. In contrast, 86% of the 2009 to 2010 mushroom crop was sold as fresh whole or sliced products (30). Production of *A. bisporus* mushrooms is highly centralized in the state of Pennsylvania, where in 2009 to 2010, 64% of the U.S. population consumed *Agaricus* mushrooms were grown.

There have been no reports to date of foodborne illness attributed to consumption of fresh commercially grown mushrooms. However, microbial surveys of fresh produce have provided evidence that mushrooms can become contaminated with *Listeria* spp. (18, 25, 38, 45, 55),

*Salmonella* spp. (14, 17, 35, 36, 38), enterohemorrhagic *Escherichia coli* (38), and *Campylobacter jejuni* (14). Several recalls and consumer advisories have been issued by government agencies after detectable levels of pathogens were found on mushrooms. In 2001, government authorities in Ireland advised consumers to cook fresh mushrooms that tested positive for *Salmonella* Kedougou (12, 17). In the United States, voluntary recalls of sliced mushrooms were issued by state health agencies in Georgia (11) and Ohio (48) after *Listeria monocytogenes* was detected. More recently, the Canadian Food Inspection Agency ordered recalls of Canadian-grown mushrooms after detectable levels of *L. monocytogenes* were found (6–9).

Commercial mushroom growing is a unique form of agriculture, in that an organic medium—known in the industry as growth substrate—must be prepared prior to *A. bisporus* inoculation. Specialized composting procedures promote the growth of thermophilic microorganisms that break down complex carbon and nitrogen compounds into forms that are more nutritionally available to mushrooms. Most mushroom growers in the United States have adopted a two-step substrate preparation process known as phase I and phase II composting (1). Phase I begins as agricultural raw materials (e.g., horse and poultry manure, straw, corn cobs, brewer's grain, cotton seed hulls, cocoa bean hulls, gypsum) are thoroughly mixed, wetted, and formed into

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long rows on outdoor concrete wharfs where they are turned periodically, watered, and reformed. Alternatively, the mixture can be moved into more efficient, enclosed bunker structures in which floor vents continuously force air through the pile. With both systems, rapid thermophile growth causes core pile temperatures to reach 70 to 80°C within 24 to 72 h. Phase I composting times range from 6 to 25 days, depending on the composition of the raw materials used and the efficiency of the equipment and facilities (37).

Phase II is a more controlled, indoor composting process designed to continue thermophilic breakdown of organic matter, eliminate any surviving pests and fungal pathogens, and disperse toxic ammonia gas that decrease crop yields and postharvest quality. Phase II traditionally begins when phase I substrate is moved into mushroom growing houses where the crop will eventually be grown; although more efficient enclosed tunnel structures are increasingly used for bulk substrate production (3). For both systems, the industry standard for phase II composting is to include an initial pasteurization treatment, wherein both substrate and air temperatures reach 60.0°C for at least 2 h, followed by a conditioning period when gradual cooling to 48.9 to 55.4°C takes place for the remaining 6 to 12 days (39). Room air temperature monitoring is included in the phase II protocol to assure uniform distribution of heat throughout the substrate. Because the chemical and physical characteristics of the substrate are critical to achieving a good crop, substrate temperatures are monitored continuously and controlled by careful introduction of steam or cool, outside air. The end result is a pest-free growth substrate that contains a diverse population of beneficial thermophilic bacteria and fungi (2, 22, 23, 31, 57).

After a series of food recalls and outbreaks were linked to on-farm contamination of fresh produce, the use of animal manures as agricultural soil supplements has come under increased scrutiny. Government and private agencies throughout the world have issued regulations and guidelines on hygienic composting management practices (15). The U.S. Food and Drug Administration (FDA) has issued general and commodity-specific farm food safety guidelines, known as Good Agriculture Practices (GAPs) (47, 50–53). FDA GAP guidance documents include handling and application practices to prevent raw and composted animal manures from becoming a source of produce contamination. California and Arizona lettuce and spinach growers have established farm food safety standards that are emerging as a GAP program model for other commodities (34). The Leafy Greens Marketing Agreement (5) established stringent standards for the use of composted animal manure including microbial limits of 1,000 MPN/g fecal coliforms and nondetectable levels, by enrichment methods, for *Salmonella* spp. and *E. coli* O157:H7. The recently enacted Food Safety Modernization Act of 2011 (P.L. 111-353) grants the FDA authority to establish similar microbial limits for composted animal manures used on produce farms.

In response to produce safety concerns and emerging regulations, the mushroom industry, in collaboration with Penn State University, proactively developed Mushroom Good Agricultural Practices (MGAP), a set of voluntary

standards for safe growing, harvesting, and handling of mushrooms (29). Among the MGAP standards required of growers who choose to participate in this program, is that substrate preparation procedures must be based on scientific principles that reduce potentially harmful microorganisms to acceptable levels. It can be expected that an FDA zero tolerance for human pathogens in composted animal manures will also apply to growth substrate prepared by mushroom growers.

Horse and poultry manure used in mushroom substrate formulations are known to be potential sources of *L. monocytogenes* (59) and *Salmonella* spp. (19–21), and horse manure is an infrequent source of pathogenic *E. coli* (26). In a recently reported microbial survey of *Listeria* spp. at a small-scale mushroom farm on the Penn State campus, *L. monocytogenes* was recovered from the phase I raw-material storage and composting environment (56). Although studies have been conducted on pathogen survival during animal manure composting (15, 60), there has been no research on the survival of human pathogens during commercial mushroom growth substrate composting. The goal of this study was therefore to determine survival characteristics of *L. monocytogenes*, *Salmonella*, and *E. coli* O157:H7 and selected fecal indicator organisms in mushroom growth substrate heated under temperature-controlled water bath conditions and in a small-scale phase II composting facility. The phase II pasteurization and conditioning step is the focus of this study, because it is the last point in the composting process during which a controlled and continuously monitored heat treatment is applied.

## MATERIALS AND METHODS

**Raw materials.** Phase I substrate used for inoculation studies was obtained from the Penn State University Mushroom Test Demonstration Facility (MTDF), a small-scale mushroom research farm. The ingredients used and the proportion added by weight in the standard MTDF mixture were wheat straw-bedded horse manure (57.1%), switch-grass straw (5.2%), kiln-dried poultry manure (2.0%), distiller's grain (2.6%), gypsum (1.6%), and water (31.5%). The ingredients (10.9 × 10<sup>3</sup> kg) were mechanically mixed and transferred to a bulk phase I bunker for a 6-day composting treatment. Core phase I pile temperatures were monitored with thermocouples embedded in the pile and controlled by continuous flow of air through a series of forced air vents on the bunker floor.

**Pathogen strains and preparation of inoculum.** Chicken-litter isolates of *Salmonella* serotypes Typhimurium (03-8970) and Heidelberg (03-8976) were obtained from the National Veterinary Services Laboratory (Ames, IA). *L. monocytogenes* Scott A (serotype 4b) and FSL R2-499 (serotype 1/2a, human epidemic origin) were obtained from the Penn State Food Microbiology Laboratory Collection. *E. coli* O157:H7 (ATCC strain 43889, American Type Culture Collection, Manassas, VA), was isolated from the feces of a patient with hemolytic uremic syndrome. Strain 93-0133 was obtained from the Penn State Food Microbiology Laboratory Collection.

Working cultures of each of the pathogens were maintained on tryptic soy agar supplemented with 0.6% yeast extract (TSAYE; Difco, BD, Sparks, MD). An isolated colony was added to 10 ml of TSBYE and incubated 18 to 22 h at 35°C. A 0.1-ml aliquot was

then transferred into 100 ml of TSBYE and incubated for 18 to 22 h at 35°C to produce stationary-phase populations. Each suspension was then centrifuged for 10 min at 9,000 rpm (Beckman model J2-21 centrifuge), and the pellet was resuspended in 100 ml of 0.85% NaCl solution. The six suspensions were combined as a composite culture (cocktail) containing approximately  $10^8$  CFU/ml each pathogen.

#### **Inoculation and treatment application: water bath study.**

Thermal inactivation studies on inoculated phase I substrate were conducted in a Department of Food Science BSL-2 microbiology laboratory. Phase I substrate (2.5 kg), obtained from the MTFD, was combined with the previously described pathogen cocktail (200 ml) and thoroughly mixed in a sterilized 6-liter stainless steel beaker with sterile forceps for 2 min. Preliminary tests confirmed that the inoculum was uniformly distributed throughout the substrate. Substrate aliquots (75 g) were weighed into polypropylene bags (UltraSource, Kansas City, MO) and vacuum sealed. The dimensions of the filled and sealed bags were 12.7 by 22.9 by 0.5 cm. The bags were heat treated by fully immersing them for up to 120 min in a temperature-controlled water bath (Thelco Model 84, Precision Scientific Co., Buffalo, NY) adjusted to 48.8, 54.4, or 60°C. Temperature come-up and come-down times were determined by inserting a temperature probe (Precision 0.01 Degree, Control Co., Friendswood, TX) in the geometric center of an uninoculated filled bag. At the end of each temperature-time treatment, duplicate bags were removed and cooled in an ice water bath until the contents reached room temperature. The contents were enumerated immediately for pathogens as described below. Uninoculated bags of phase I substrate were used to follow changes in microbial indicator populations at each temperature and time. The experiment was repeated for each temperature. After inoculation experiments were completed, all substrate samples were autoclaved at 121°C and properly disposed.

**Inoculation and treatment application: phase II composting study.** Microbial inoculation and treatment protocols were reviewed by the Penn State Institutional Biosafety Committee (IBC) prior to conducting experiments (IBC no. 15454). Because the MTFD produces mushrooms for campus food service outlets, it was not possible to conduct phase II inoculation studies at this facility. Instead, finished phase I substrate was transported to the Penn State Mushroom Research Center (MRC) and into a pilot-scale phase II room (7.9 m<sup>2</sup>) wherein precise temperature control could be maintained. Contamination risks were minimized by the fact that no mushrooms at the MRC are sold to the public, and inoculation procedures took place off-site in a Department of Food Science biosafety level II laboratory. All bags were wiped with 70% ethanol prior to leaving the laboratory.

Phase I substrate (2.50 kg), obtained from the MTFD, and the pathogen cocktail (200 ml) were filled simultaneously into three polypropylene bags (4 liter; Unicorn Imp. and Mfg., Garland, TX) that are normally used for cultivating non-*Agaricus* specialty mushrooms. Each bag had two perforated filter patches (3 by 6 cm, 0.2 µl) on the sides to facilitate gas exchange between the inside of the bag and the external environment. The contents of each bag were mixed thoroughly by manually inverting and massaging the bags for 2 min. Preliminary tests confirmed that this technique was sufficient to uniformly distribute the inoculum. Two hundred milliliters of 0.85% NaCl was added to a fourth bag to serve as an uninoculated control. After mixing, a 250-g sample was aseptically removed from inoculated and uninoculated bags for determination of initial populations of pathogens and indicator microorganisms, respectively. A wireless temperature probe (HOBO Water Temp

Pro, Onset Computers, Bourne, MA) was placed at the approximate center of each of the bags, and a metal twist was applied to loosely seal the contents. The bags were then transported to the MRC.

At the MRC, each of the four bags was placed in the center of a standard wooden mushroom growing tray (61 by 61 by 30 cm). Uninoculated phase I substrate was filled around and above each bag until tray capacity was reached. The trays were stacked vertically and evenly spaced between eight additional trays of uninoculated substrate. Another stack of uninoculated trays of substrate was added to fill the room in order to facilitate consistent and controllable air temperatures. Substrate temperatures were continuously monitored by placing eight thermocouples in uninoculated substrate. Room air temperature was monitored by a single thermocouple placed in the center of the room. During phase II, as thermophilic decomposition reoccurred, a programmable temperature controller (SamStat, Landenberg, PA) was used to raise or lower substrate temperature by automatically venting steam or cool air, respectively, into the room. Pasteurization was considered complete by the MRC operator when the temperature of the air and the lowest reading probe in the substrate reached 60°C for 2 h. Conditioning continued until the end of the 6-day process, after which the bags were removed from the trays and returned to the Food Science Department for enumeration of pathogens and indicator microorganisms. Levels for moisture, total carbon, and total nitrogen in uninoculated substrate were determined at the Penn State Agricultural Analytical Services Laboratory (University Park, PA) by using U.S. Department of Agriculture (USDA) standard test methods for examination of compost. Uninoculated substrate remaining in trays was disposed of on-site so that it could not be reused for other MRC experiments.

**Microbiological analysis.** For water bath experiments, 25 g of substrate was combined with 225 ml of buffered peptone water (Difco, BD) in a variable-speed stainless steel laboratory blender (model 38BL54, Waring Laboratory, Torrington, CT) and blended for 30 s at medium speed. For the phase II composting study, 250 g of substrate was combined with 2,250 ml of buffered peptone water in a sterilized 6-liter stainless steel beaker. The mixture was stirred by hand for 2.0 min with sterile forceps. Infusions obtained by both methods were serially diluted in buffered peptone water for microbial enumeration.

Aerobic plate count, *E. coli*, total coliforms, and *Enterobacteriaceae* populations in uninoculated substrate were enumerated with Petrifilm methods from 3M (St. Paul, MN) with incubation at 37°C for 24 h. *L. monocytogenes*, *Salmonella* spp., and *E. coli* O157:H7 populations were determined by spread plating 0.1 ml on modified Oxford agar (Difco, BD), xylose lysine deoxycholate (XLD) agar (Difco, BD), or MacConkey sorbitol agar (Difco, BD), respectively, followed by incubation for 24 to 48 h at 35°C. Black colonies on modified Oxford agar or XLD agar were considered presumptive positive for *Listeria* spp. or *Salmonella* spp., respectively, and were confirmed positive by using a *Listeria* visual immunoassay (TECRA International, Frenchs, Australia) or a latex agglutination test (Oxoid, Thermo Scientific, Waltham, MA). Presumptive-positive *E. coli* O157:H7 colonies on MacConkey sorbitol agar were colorless and were confirmed as positive for the O157 antigen by using an *E. coli* O157 latex agglutination test (Oxoid).

When pathogens could not be detected by direct plating, standard enrichment procedures for *Listeria* spp. (46), *Salmonella* spp. (49), and *E. coli* O157:H7 (54) were used to confirm complete thermal destruction. *L. monocytogenes* was enriched by transfer-

TABLE 1. Destruction of *Listeria monocytogenes*, *Escherichia coli* O157:H7, and *Salmonella* in mushroom growth substrate held at 48.8, 54.4, and 60.0°C

Temp (°C)	Time (h)	Population (log CFU/g) <sup>a</sup>		
		<i>Listeria monocytogenes</i>	<i>Escherichia coli</i> O157:H7	<i>Salmonella</i>
48.8	0	7.96 ± 0.03	8.09 ± 0.08	7.69 ± 0.02
	0.5	7.58 ± 0.02	7.93 ± 0.03	7.26 ± 0.03
	1.0	7.30 ± 0.21	7.75 ± 0.09	7.07 ± 0.12
	2.0	6.79 ± 0.17	7.35 ± 0.08	6.15 ± 0.02
	4.0	5.45 ± 0.33	6.61 ± 0.12	4.66 ± 0.41
	6.0	5.04 ± 0.26	6.35 ± 0.18	3.93 ± 0.03
	12	3.85 ± 0.00	4.74 ± 0.51	2.50 ± 0.05
	24	1.00 ± 0.00 <sup>b</sup>	1.00 ± 0.00	ND <sup>c</sup>
	36	ND	ND	ND
54.4	0	7.94 ± 0.15	8.02 ± 0.24	7.44 ± 0.28
	0.25	6.70 ± 0.10	6.26 ± 0.18	5.33 ± 0.23
	0.5	6.20 ± 0.03	5.27 ± 0.13	3.65 ± 0.35
	1.0	5.35 ± 0.05	4.28 ± 0.72	1.00 ± 0.00
	1.5	4.95 ± 0.00	2.59 ± 1.59	1.00 ± 0.00
	2.0	3.24 ± 0.24	1.65 ± 0.65	1.00 ± 0.00
	4.0	1.00 ± 0.00	1.00 ± 0.00	1.00 ± 0.00
	6.0	1.00 ± 0.00	1.00 ± 0.00	1.00 ± 0.00
	8.0	ND	ND	ND
60.0	0	7.92 ± 0.03	8.39 ± 0.11	7.90 ± 0.07
	0.05	6.20 ± 0.08	2.50 ± 1.50	1.00 ± 0.00
	0.10	5.40 ± 0.32	ND	ND
	0.15	4.30 ± 0.35	ND	ND
	0.20	1.00 ± 0.00	ND	ND
	0.25	1.00 ± 0.00	ND	ND
	0.50	ND	ND	ND
	1.0	ND	ND	ND
	2.0	ND	ND	ND

<sup>a</sup> Mean values obtained from two replicate experiments ± standard deviations.

<sup>b</sup> 1.00 ± 0.00, negative by direct plating method (<1.0 log CFU/g), positive by enrichment method (>1 cell per 10 g).

<sup>c</sup> ND, not detected. Negative by direct plating method (<1.0 log CFU/g), negative by enrichment method (<1 cell per 10 g).

ring 10 g of substrate to 90 ml of UVM formulation broth (Difco, BD) and incubating at 35°C for 24 h. One milliliter of the infusion was transferred to 10 ml of Fraser broth (Difco, BD) and incubated for an additional 24 h at 35°C. Loopfuls were streaked onto modified Oxford agar and incubated for 24 h at 35°C. Presumptive-positive colonies appeared black and were confirmed positive for *Listeria* spp. by using *Listeria* visual immunoassay. *Salmonella* was enriched by adding 10 g of substrate to 90 ml of lactose broth (Difco, BD) and incubating for 24 h at 35°C. One milliliter was added to 10 ml each of selenite-cysteine broth (Difco, BD) and tetrathionate broth (Difco, BD), followed by incubation for 24 h at 35°C. Each broth was streaked onto XLD agar (Difco, BD). Colonies presumptively positive for *Salmonella* were black on XLD agar and were confirmed with latex agglutination (Oxoid). Pathogenic *E. coli* were enriched by transferring 10 g of compost to 90 ml of brain heart infusion broth (Difco, BD) and incubating at 35°C for 3 h. The broth was then decanted into 100 ml of double-strength tryptose phosphate broth (Difco, BD) and incubated 20 h at 44°C. After incubation, loopfuls were streaked onto MacConkey sorbitol agar and incubated 24 h at 35°C. Presumptive-positive colonies appeared colorless and were confirmed with latex agglutination (Oxoid).

**Statistical analysis.** Bacterial populations were converted to log CFU per gram. Analysis of variance and mean separation

analysis by using Tukey's honestly significant difference at the  $\alpha = 0.05$  level were used to compare bacteria levels obtained from the mean of two replicate water bath or MRC experiments. Regression analysis was performed to determine death rates for microorganisms.

## RESULTS

**Water bath study.** Substrate inoculated with *L. monocytogenes*, *E. coli* O157:H7, and *Salmonella* was held in vacuum-sealed polypropylene bags to prevent moisture loss during extended heating and to allow precise control of temperature and time treatments. Temperature come-up times for substrate held in a water bath at 48.8, 54.4, and 60.0°C were 4.5, 5.0, and 5.0 min, respectively. Initial levels of the three pathogens were between 7.4 and 8.4 log CFU/g. Population levels for each of the pathogens during heating at each of the three temperatures are shown in Table 1. Thermal destruction of each of the pathogens was affected significantly ( $P \leq 0.05$ ) by heating temperature and time. *Salmonella* and *E. coli* O157:H7 appeared to be less heat resistant than *L. monocytogenes*; however, there were not enough data points at 60°C to make statistically valid comparisons of thermal activation rates for all temperatures.

TABLE 2. Decimal reduction times (D-values) at 54.4°C for destruction of *Listeria monocytogenes*, *Escherichia coli* O157:H7, and *Salmonella* in inoculated substrate and indicator organisms in uninoculated substrate

Microorganism	$D_{54.4}$ -value (min)
<i>Listeria monocytogenes</i>	27.0 A <sup>a</sup>
<i>Escherichia coli</i> O157:H7	22.1 A
<i>E. coli</i>	13.9 B
Total coliforms	13.9 B
<i>Enterobacteriaceae</i>	13.5 B
<i>Salmonella</i>	10.8 B

<sup>a</sup> Mean values obtained from two replicate experiments followed by the same letter are not significantly different ( $\alpha = 0.05$ ).

Minimum times at which all three of the pathogens were not detectable by plating and enrichment were 36.0, 8.0, and 0.5 h at 48.8, 54.4, and 60°C, respectively.

Destruction of indicator microorganisms and pathogens were followed at 54.4°C so that enough data points could be generated to make statistically valid rate comparisons. Pretreatment populations of *E. coli*, total coliforms, and *Enterobacteriaceae* were 5.4 to 6.0 log CFU/g. Therefore, initial pathogen inoculum levels were adjusted to a similar range of 5.0 to 6.0 log CFU/g (individual data not shown).

Linear log reductions for each of the microbial indicators and pathogens at 54.4°C for up to 2 h were observed, as evidenced by greater than 0.9  $R^2$  values for each type or species. Microbial heat resistance was therefore compared by calculating decimal reduction times at 54.4°C ( $D_{54.4}$ -values).  $D_{54.4}$ -values for *L. monocytogenes* and *E. coli* O157:H7 were significantly ( $P \leq 0.05$ ) higher than for *Salmonella* and for *E. coli*, total coliforms, and *Enterobacteriaceae* indicator microorganisms (Table 2).

**Phase II composting study.** A representative plot of temperature changes that occurred within bags of substrate during the 6-day (144 h) phase II MRC process is shown in Figure 1. Remote probe readings from within inoculated bags were consistent with process control measurements taken directly from trays of uninoculated substrate. The pasteurization interval, during which the operator verified that substrate and room air temperatures increased to at least 60°C for 2 h, began approximately 24 h after phase II began. During the conditioning interval that followed, introduction of cool air into the room lowered substrate temperatures to between 46 and 53°C. After phase II composting, the substrate contained 77.2% moisture, 36.2% total carbon (dry weight basis), and 1.8% total nitrogen (dry weight basis) (C:N = 20.1).

Data for the destruction of naturally occurring indicator microorganisms and inoculated pathogens during phase II composting are shown in Table 3. Microbial populations were only determined immediately before and after composting, because entry into the phase II room was not permitted once it was sealed. No pathogens were found in any of the uninoculated samples. Initial levels of *L. monocytogenes*, *Salmonella*, and *E. coli* O157:H7 after

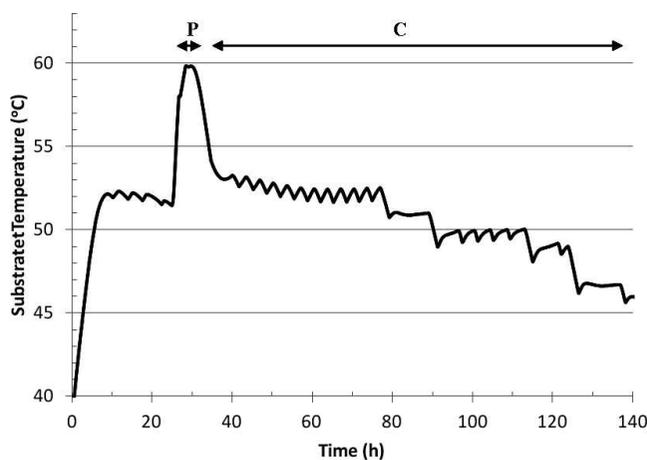


FIGURE 1. Representative plot of substrate temperatures taken during phase II composting. Plot line represents the average temperature reading taken from four bags every 0.1 h. Pooled standard deviation = 1.4°C. P, pasteurization; C, conditioning.

inoculation ranged from 7.2 to 8.1 log CFU/g. After phase II composting, none of the pathogens were detected by direct plating or by enrichment.

Despite phase I core pile temperatures of up to 80°C, populations of *E. coli*, coliforms, and *Enterobacteriaceae* prior to phase II composting were between 4 and 6 log CFU/g (Table 3). Destruction of the three microbial indicators followed that of the three pathogens. All were below detectable levels after phase II composting. In contrast, aerobic plate count populations, initially at nearly 9 log CFU/g, decreased by less than 2 log CFU/g during phase II composting.

## DISCUSSION

The  $D_{54.4}$ -values determined in the water bath study are within ranges reported in several studies on the destruction of the same pathogens in meat and poultry heated at 54 to 56°C (13, 28, 32, 33). The  $D_{54.4}$ -value for destruction of *E. coli* O157:H7 in mushroom substrate is comparable to the  $D_{55}$ -value of 35.4 min reported for composted dairy manure by Jiang et al. (24). The more rapid destruction of *E. coli*, total coliforms, and *Enterobacteriaceae* compared with *L. monocytogenes* and *E. coli* O157:H7 at 54.4°C indicate that fecal indicators are not appropriate for predicting pathogen destruction during phase II composting.

The high levels of *E. coli*, total coliforms, and *Enterobacteriaceae* in substrate used for phase II inoculation studies confirms our assumption that phase I composting is not an adequate microbial hurdle for eliminating pathogens associated with horse and poultry manure. It is likely that during phase I, uneven distribution of heat within the pile occurred, which resulted in areas where temperatures were low enough to permit survival of mesophilic bacteria.

The absence of recoverable pathogens and less than detectable levels of microbial indicators after phase II composting is a reasonable result, given the temperature history data from which Figure 1 was derived. The time during MRC composting experiments at which all substrate temperature measurements were at or above 60°C was 1.9 h,

TABLE 3. Destruction of *Listeria monocytogenes*, *Escherichia coli* O157:H7, and *Salmonella* in inoculated substrate and indicator microorganisms in uninoculated substrate during phase II composting

	Microbial population (log CFU/g) <sup>a</sup>	
	After phase I	After phase II
Inoculated pathogens		
<i>Listeria monocytogenes</i>	7.95 ± 0.21	ND <sup>b,c,d</sup>
<i>Escherichia coli</i> O157:H7	7.95 ± 0.21	ND <sup>b,c</sup>
<i>Salmonella</i>	7.40 ± 0.28	ND <sup>b,c</sup>
Indicator organisms		
<i>E. coli</i>	4.70 ± 0.14	ND <sup>b</sup>
Total coliforms	5.10 ± 0.28	ND <sup>b</sup>
<i>Enterobacteriaceae</i>	5.80 ± 0.28	ND <sup>b</sup>
Aerobic plate count	8.75 ± 0.35	6.90 ± 0.43

<sup>a</sup> Mean ± standard deviation of the mean population of bacteria. Populations were significantly different before and after phase II composting for all microorganisms ( $\alpha = 0.05$ ).

<sup>b</sup> Negative by the direct plating method (<1.0 log CFU/g).

<sup>c</sup> Negative by the enrichment method (<1 cell per 10 g).

<sup>d</sup> ND, not detected.

while the water bath study (Table 1) showed that only 0.5 h at 60°C was required to achieve complete destruction of all pathogens. Moreover, the contribution of the conditioning period to the total lethality of the phase II process should be considered. The time substrate was exposed to temperatures ranges of 48.8 to 54.3 and 54.4 to 59.9°C was 21 and 5 h, respectively. According to Table 1, each of these temperature and time conditions is individually severe enough to achieve complete pathogen destruction. Thus, the total lethality achieved during MRC phase II composting was far more severe than required to achieve complete destruction of each of the inoculated pathogens. Absence of detectable levels of *E. coli*, total coliforms, and *Enterobacteriaceae* after phase II is also not unexpected given the more rapid destruction of these fecal indicators compared with *L. monocytogenes* and *E. coli* O157:H7 (Table 2). High aerobic plate count levels before and after phase II composting can be attributed to selective survival of nonpathogenic thermophilic bacteria and fungi during prolonged exposure to phase II temperatures (22, 23, 31, 57).

Results from the water bath and MRC phase II composting studies demonstrate that on-farm verification of phase II composting as an effective food safety control method can only be achieved by testing for the most heat-resistant pathogens in substrate, which this study has shown are *L. monocytogenes* and/or *E. coli* O157:H7.

Although the phase II method used in this study is the standard procedure used in the mushroom industry (2), actual microbial reductions achieved can differ among farms. In general, differences in microbial heat resistance have been attributed to chemical and physical characteristics of the matrix tested (13). During composting, pathogen

destruction rates are influenced by the C:N ratio of the ingredient mixture; the amount of volatile acids, free ammonia, and moisture present; microflora levels; and temperature gradients within piles (15, 60). Differences in the types of raw materials and their proportion in mushroom growth substrate can affect chemical, physical, and microbiological characteristics of formulations and thus the amount of heat generated during composting. The moisture content of the phase II substrate was higher and the C:N ratio was lower than optimal levels for pathogen reduction reported in studies of other types of agricultural composting systems (42, 43). However, these levels are within the expected range that occurs on commercial mushroom farms (3, 39). Further experiments on the kinetics of pathogen inactivation as affected by moisture and C:N ratio were not conducted, because the mushroom composting process has been optimized for *A. bisporus* crop yield and quality, and any deviation from well-established protocols would likely be unacceptable to the industry.

It should be noted that the pathogen inoculum used for the experiments in this study, was prepared from cultures grown in the laboratory at 35°C. On commercial farms, any pathogenic bacteria that survive phase I composting would likely to have been exposed to high, but sublethal, temperatures. Several model system and food studies have demonstrated increased thermotolerance of *L. monocytogenes*, *E. coli* O157:H7, and *Salmonella* spp. initially exposed to sublethal temperatures between 40 and 52°C (4, 10, 16, 27, 40). Singh et al. (41) reported that thermal resistance of *L. monocytogenes*, *E. coli* O157:H7, and *Salmonella* spp. in dairy compost was higher if the inoculum was first heat shocked at 47.5°C for 1 h. Therefore, it is possible that any pathogens present in phase I ingredients are more resistant to phase II composting temperatures than those used in the inoculum for this study. The growth stage of pathogens used to prepare the inoculum might also affect thermotolerance (58). Further studies could be conducted with pathogen cultures grown at different life cycle stages and that have been heat adapted at sublethal temperatures that occur during phase I composting prior to inoculation. Varying inoculation levels would also be useful to compare the efficacy of phase II composting over a wide range of initial pathogen populations.

Nevertheless, the observed  $\geq 7$ -log reductions in *L. monocytogenes*, *E. coli* O157:H7, and *Salmonella* show that a phase II composting process can be an effective food safety control measure that meets stringent microbial standards for composted animal manures, including those established by the Leafy Greens Marketing Agreement (5) and FDA composting regulations.

Although safety considerations preclude pathogen inoculation studies on commercial farms, growers should collect additional data to validate their own phase II composting process. Growers should consider that substrate temperature gradients can exist in growing rooms or in tunnels, and studies should be conducted to detect any cold spots during phase II composting. Temperature monitoring probes should be placed in these areas to assure that lethal temperatures reach every part of the substrate.

Growers can take steps to verify that a phase II composting process is consistently applied by regularly reviewing temperature profile records and by testing substrate for *L. monocytogenes* at least once per year or anytime major changes are made to the type and proportion of raw materials used. Because temperature monitoring and control equipment are important for obtaining consistent phase II heat treatments, they should be regularly maintained and calibrated to ensure that readings are accurate.

Where phase I composting is conducted at the same location as growing operations, MGAP standards for preventing cross-contamination should be adhered to. These include keeping areas where horse and poultry manure are received, stored, and handled physically separated from mushroom growing areas and controlling the movement of equipment and workers between these areas.

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