

# Identification of *Escherichia coli* O157 by Using a Novel Colorimetric Detection Method with DNA Microarrays

Beatriz Quiñones,<sup>1</sup> Michelle S. Swimley,<sup>1</sup> Amber W. Taylor,<sup>2</sup> and Erica D. Dawson<sup>2</sup>

## Abstract

Shiga toxin-producing *Escherichia coli* O157 is a leading cause of foodborne illness worldwide. To evaluate better methods to rapidly detect and genotype *E. coli* O157 strains, the present study evaluated the use of ampliPHOX, a novel colorimetric detection method based on photopolymerization, for pathogen identification with DNA microarrays. A low-density DNA oligonucleotide microarray was designed to target *stx1* and *stx2* genes encoding Shiga toxin production, the *eae* gene coding for adherence membrane protein, and the *per* gene encoding the O157-antigen perosamine synthetase. Results from the validation experiments demonstrated that the use of ampliPHOX allowed the accurate genotyping of the tested *E. coli* strains, and positive hybridization signals were observed for only probes targeting virulence genes present in the reference strains. Quantification showed that the average signal-to-noise ratio values ranged from  $47.73 \pm 7.12$  to  $76.71 \pm 8.33$ , whereas average signal-to-noise ratio values below 2.5 were determined for probes where no polymer was formed due to lack of specific hybridization. Sensitivity tests demonstrated that the sensitivity threshold for *E. coli* O157 detection was 100–1000 CFU/mL. Thus, the use of DNA microarrays in combination with photopolymerization allowed the rapid and accurate genotyping of *E. coli* O157 strains.

## Introduction

SHIGA TOXIN-PRODUCING *Escherichia coli* O157:H7 (*E. coli* O157) is an enteric pathogen known to cause human gastrointestinal illnesses with clinical spectra, ranging from bloody diarrhea and hemorrhagic colitis to life-threatening hemolytic-uremic syndrome (Rangel *et al.*, 2005; Karmali, 2009). In recent years, most risk management efforts have focused on identifying the presence and distribution of *E. coli* O157 in foods as part of imperative constituents of food safety programs (Karmali, 2009; Buchanan and Appel, 2010). Thus, the O157 serotype is considered to be the most commonly reported serotype linked to outbreaks in North America.

The rise in foodborne-related outbreaks of *E. coli* O157 has heightened the importance of developing improved methods to rapidly detect and characterize virulent strains (Bettelheim and Beutin, 2003; Bettelheim, 2007; Buchanan and Appel, 2010). Established culturing methods for the identification of pathogenic *E. coli* can be labor intensive and time consuming and assess phenotypic and genotypic markers by screening only a small number of determinants. Molecular-based technologies, such as DNA microarrays, offer a viable alternative

to screen multiple markers simultaneously (Uttamchandani *et al.*, 2009; Call, 2005).

One challenge of using the DNA microarray platforms for pathogen detection has been developing cost effective as well as sensitive procedures that would indicate the positive signals on the array (Call, 2005; Kuck and Taylor, 2008; Vora *et al.*, 2008). Fluorescent assays are commonly used methods for microarray-based detection of pathogens. However, limitations of these assays are that labeling of target DNA can be inconsistent and highly variable, and they utilize expensive and nonportable scanners for data acquisition and analysis (Call, 2005; Kuck and Taylor, 2008; Vora *et al.*, 2008). Other nonfluorescent, colorimetric assays employ the use of unstable reagents that require temperature-controlled environments and that have variable development times, leading to an increase in nonspecific backgrounds (Kuck and Taylor, 2008). Improved procedures for pathogen surveillance are thus needed with sufficient sensitivity, cost effectiveness, and suitability for routine testing. To implement the use of better detection methods for categorizing *E. coli* O157 strains, the present study evaluated ampliPHOX, a novel and innovative colorimetric procedure that is based on light-initiated signal

<sup>1</sup>Produce Safety and Microbiology Research Unit, Western Regional Research Center, Agricultural Research Service, U.S. Department of Agriculture, Albany, California.

<sup>2</sup>InDevR, Inc., Boulder, Colorado.

amplification through polymerization (Kuck and Taylor, 2008; Sikes *et al.*, 2008; Dawson *et al.*, 2009). Our findings demonstrate that photopolymerization is a simple and rapid method for the accurate genotyping of *E. coli* O157 strains with DNA microarrays.

## Materials and Methods

### Bacterial strains and growth conditions

*E. coli* O157:H7 strains RM1600 (feedlot strain 8) and RM1697 (feedlot strain 42) were isolated from beef cattle feed yards in Kansas (Kimura *et al.*, 2000). *E. coli* O157:H7 strains RM4876 and RM6011 were isolated from watershed samples collected in the Salinas Valley in California (Cooley *et al.*, 2007; Quiñones *et al.*, 2009). *E. coli* O55:H7 strain RM3654 (CPK100) and *E. coli* O113:H21 strain RM3655 (CPK117) were obtained from the Canadian Pediatric Kidney Disease Research Center in Ottawa, Ontario, Canada. Nonpathogenic *E. coli* strain RM5034 (strain K12; ATCC-29425) was obtained from the American Type Culture Collection (Manassas, VA). The genotypic and phenotypic characteristics of the *E. coli* reference strains were described previously (Kimura *et al.*, 2000; Cooley *et al.*, 2007; Quiñones *et al.*, 2009). Bacterial cultures were propagated on Luria-Bertani (LB) agar (Difco, Detroit, MI) or were grown aerobically with constant shaking (200 rpm) in LB broth (Difco) at 37°C.

### Construction of the DNA microarray

The 30-mer oligonucleotide probes targeting the intimin adherence protein (*eae*) (Zhang *et al.*, 2002; Ramachandran *et al.*, 2003), O157 O-antigen perosamine synthetase (*per*) (Bilge *et al.*, 1996; Reeves *et al.*, 1996; Shimizu *et al.*, 1999), Shiga toxin 1 (*stx1*), and Shiga toxin 2 (*stx2*) (Nataro and Kaper, 1998; Bettelheim, 2007; Serna and Boedeker, 2008) in pathogenic *E. coli* are shown in Table 1. The *E. coli* O157-specific probes were designed by using PRIMER3 software (Rozen and Skaltsky, 2000) and were purchased from Eurofins MWG Operon (Huntsville, AL) with a 5'-amino-C6 modification for covalent binding to the slide surface. The probes were spotted in triplicate at a final concentration of 20 μM on aldehydesilane-coated glass slides (Nexterion® Slide AL; SCHOTT, Inc., Elmsford, NY). As a control for the photopolymerization detection process, a synthetic, 24-mer oligonucleotide probe with 5'-amino-C6 and 3'-biotin modifications (InDevR, Inc., Boulder, CO) was spotted at a final concentration of 500 nM. This biotinylated control oligonucleotide did not have any sequence homology to any *E. coli* genes. The microarrays were manufactured by Applied Microarrays, Inc. (Tempe, AZ) with a spot diameter size of 250 μm and a center-to-center spacing of 700 μm. After printing, an adhesive silicone well (0.5 mm thickness, 9 mm diameter; Grace Bio-Labs, Bend, OR) was placed in the center of the printed array, and the microarray slides were stored in a desiccator until further use.

### Polymerase chain reaction

Polymerase chain reaction (PCR) reagents were supplied by New England Biolabs (Ipswich, MA). PCR primers were purchased from Eurofins MWG Operon with a 5'-phosphorylated modification for the forward primers and a 5'-biotin modification for the reverse primers (Table 1). For preparing bacterial crude lysates, cultures of the reference strains were

TABLE 1. LIST OF DNA OLIGONUCLEOTIDES USED IN THE DNA MICROARRAY

Target gene	Primer sequences for PCR		Amplicon size (bp)	Accession no.	Reference
	Probe sequences (5' → 3')	Forward (5' → 3')			
<i>stx1A</i>	GTTCTTATGTAATGACTGCTGAAGATGTTG	CTGTGGCAAGAGCGATGTTA	163	AAG57228	This study
<i>stx2A</i>	GTTAATGGAGTTCAGTGGTAATACAATGAC	AATCAGTCGTCACCTACTGGT	193	AAG55587	This study
<i>per</i>	TAGGCTACAATTATAGGATGACAAATATCT	GGTGAAGGTGGAAATGGTTGTC	206	AAG57096	This study
<i>eae</i>	GTTAATCTGCAGATGGTAATAACTTTGAC	CCGCTCTTGGTATCGCTGGT	223	AAG58823	This study

PCR, polymerase chain reaction.

grown aerobically in LB broth for 24 h, and 100  $\mu$ L of the bacterial culture was collected by centrifugation at 2000 g for 5 min. Cell pellets were resuspended in 100  $\mu$ L of HyPure™ molecular biology-grade water (HyClone Laboratories, Inc., Logan, UT) and incubated at 100°C for 20 min. The lysates were centrifuged at 2000 g for 5 min, and the supernatants were collected and frozen until further use. For experiments that assessed the sensitivity threshold of the detection assay, cell lysates were prepared from tenfold dilutions of a culture from *E. coli* O157 strain RM1697. Multiplex PCR amplifications were performed in a 50  $\mu$ L reaction mixture, each containing 5  $\mu$ L of the bacterial crude lysate, 0.5  $\mu$ M of *eae* primers, 0.1  $\mu$ M (each) of *per*, *stx1*, and *stx2* primers, 1 $\times$ ThermoPol *Taq* buffer with 2 mM MgSO<sub>4</sub>, 200  $\mu$ M deoxynucleoside triphosphates, and 2.5 U of *Taq* polymerase. The reaction mixtures were placed in a Dyad Peltier Thermal Cycler (Bio-Rad Laboratories, Hercules, CA) with the following settings: 2 min at 94°C, followed by 30 cycles of 30 sec at 94°C, 1 min at 60°C, 45 sec at 72°C, and a final extension time of 5 min at 72°C.

#### Microarray hybridization

For each hybridization reaction, 45  $\mu$ L of PCR amplicons was purified by using the MinElute® PCR purification kit (Qiagen, Valencia, CA), and 20  $\mu$ L of the eluate was digested with 10 U of lambda exonuclease and 1 $\times$ lambda exonuclease reaction buffer (Epicenter Biotechnologies, Madison, WI) in a final volume of 23  $\mu$ L for 5 min at 37°C, followed by immediate addition of an equal volume of InDevR 2 $\times$ Hyb Buffer (InDevR, Inc.). The hybridization mixture was applied to each microarray that was previously rinsed with distilled water for 5 min, and the microarray slides were further incubated for 1 h at room temperature. After hybridization, the slides were transferred to a microarray wash bin (InDevR, Inc.) and were rinsed briefly with InDevR Wash Buffer D, Wash Buffer A for 1 min, Wash Buffer D briefly, Wash Buffer B for 5 min, and Wash Buffer C for 5 min and then dried.

#### Microarray labeling, signal amplification, and analysis

The specific colorimetric detection of signals on the microarray were detected with the ampliPHOX™ Detection System (InDevR, Inc.), a light-initiated signal amplification through polymerization (Kuck and Taylor, 2008; Sikes *et al.*, 2008; Dawson *et al.*, 2009). The hybridized microarrays were labeled subsequently with ampliTAG™ labeling solution (InDevR, Inc.) for 5 min at room temperature in a dark humidity chamber, and immediately after labeling, microarrays were rinsed briefly with Wash Buffer D, Wash Buffer C for 5 min, and then dried. For the detection of positive signals on the microarray, ampliTAG-labeled microarrays were covered with 40  $\mu$ L ampliPHY™ reagent and photoactivated for 220 sec by using the ampliPHOX Reader (InDevR, Inc.). Immediately after photoactivation, arrays were rinsed with water, stained with 2 drops of ampliRED™ for 5 min and finally rinsed again with water to observe polymer formation. Color digital images of the stained arrays were collected by using the ampliPHOX Reader and the associated ampliVIEW™ software 1.0 (InDevR, Inc.). For each spot, the quantification of signal and background intensities was quantified by using ampliVIEW software and expressed as signal-to-noise ratio (SNR). SNR values were calculated as the difference between the mean pixel intensities of the signal minus the

mean pixel intensities of the background divided by the standard deviation of the background (He and Zhou, 2008).

## Results

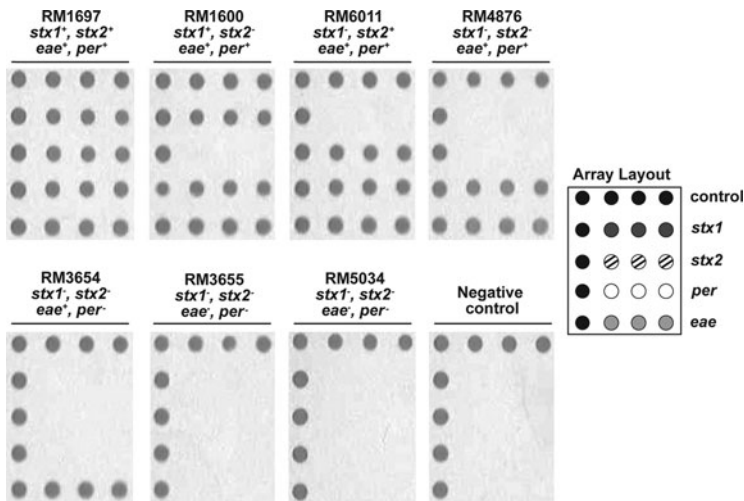
### *A novel colorimetric method for detection of pathogenic E. coli on DNA microarrays*

To implement the use of improved detection methods with DNA microarrays, the present study evaluated photopolymerization, a novel colorimetric detection method (Kuck and Taylor, 2008; Sikes *et al.*, 2008; Dawson *et al.*, 2009), for genotyping *E. coli* O157 strains. To provide a proof of concept for testing photopolymerization, a low-density 30-mer DNA oligonucleotide array was constructed to target the *E. coli* O157 virulence genes *eae*, *per*, *stx1*, and *stx2* (Table 1). To determine that a gene target was present in the sample, a multiplex PCR amplification was performed to generate biotinylated fragments, ranging from 163 to 223 base pairs (Table 1). The PCR-amplified fragments were subjected to lambda exonuclease digestion to produce single-stranded DNA targets for achieving a rapid microarray hybridization (Boissinot *et al.*, 2007). For photopolymerization to occur, a streptavidin-conjugated photoinitiator is used to specifically label the microarrays that have been hybridized with the biotin-labeled, single-stranded DNA targets (Kuck and Taylor, 2008). After irradiating at a wavelength absorbed by the photoinitiator ( $\lambda$  404 nm), polymer forms exclusively in only a few minutes where the probe sequences were spotted on the microarray (Kuck and Taylor, 2008; Dawson *et al.*, 2009); polymer formation can be observed after staining (Kuck and Taylor, 2008).

### *Specificity and sensitivity for pathogenic E. coli detection with photopolymerization*

To validate the specificity of using photopolymerization with a low-density microarray for detecting pathogenic *E. coli* O157, several reference strains with different genotypes were examined (see Materials and Methods section). Bacterial crude lysates, prepared from the tested strains, were subjected to a multiplex PCR amplification of *eae*, *per*, *stx1*, and *stx2* loci, and the amplified gene targets were hybridized to the DNA microarray. The labeling and subsequent microarray signal amplification was achieved using photopolymerization. The results demonstrated polymer formation for all spots targeting the genes represented on the microarray when *E. coli* O157 strain RM1697, a strain that harbors all four genes, was examined (Fig. 1). Polymer formation on microarray probes targeting *stx1*, *eae*, and *per* but not for *stx2* was observed for the *stx2*-negative *E. coli* O157 strain RM1600 (Fig. 1). Further, specificity in photopolymerization was observed also when analyzing the *E. coli* O157 strains RM6011 and RM4876 and the *E. coli* O55 strain RM3654 (Fig. 2). Polymer formation was only observed on control probes for the photopolymerization reaction for the *E. coli* O113 strain RM3655 (Fig. 1), a strain that tested negative for the genes represented on the microarray, as determined in a PCR assay (data not shown). Similar results to those obtained for strain RM3655 were also observed for the nonpathogenic *E. coli* strain RM5034 and for a negative control reaction lacking any DNA template (Fig. 1). These findings demonstrated a high level of probe specificity, and the patterns of photopolymerization correlated with the genotype of the strain.





**FIG. 1.** Specific identification of pathogenic *Escherichia coli*. *E. coli* reference strains with different genotypes were examined by DNA microarrays and photopolymerization (see Materials and Methods section). Each panel demonstrates polymer formation on probes spotted on the DNA microarrays. The array layout shows the construction of the microarray. Probes targeting *stx1* (dark gray), *stx2* (hatched), *per* (white), and *eae* (light gray) were spotted in triplicate, and positive controls for polymer formation (black) were spotted along the edges.

To quantitate the specificity of the photopolymerization reaction, the SNR values were measured for each probe on the microarray when testing the *E. coli* reference strains. Image quantification of *E. coli* O157 strain RM1697 showed that positive signals detected for all probes targeting virulence genes had SNR values, ranging from  $58.2 \pm 5.58$  to  $66.83 \pm 7.55$  (Fig. 2). Similarly, high SNR values, ranging from  $47.73 \pm 7.12$  to  $76.71 \pm 8.33$ , were measured after analysis of positive signals for the other *E. coli* O157 strains RM1600, RM6011, and RM4876 as well as for *E. coli* O55 strain RM3654 (Fig. 2). Probes that had no polymer formation due to a lack of specific hybridization had significantly lower SNR values, ranging from  $0.69 \pm 0.39$  to  $2.48 \pm 0.50$  (Fig. 2). The results from the specificity experiments indicated a clear difference between signal intensities and background noise for probes with polymer formation.

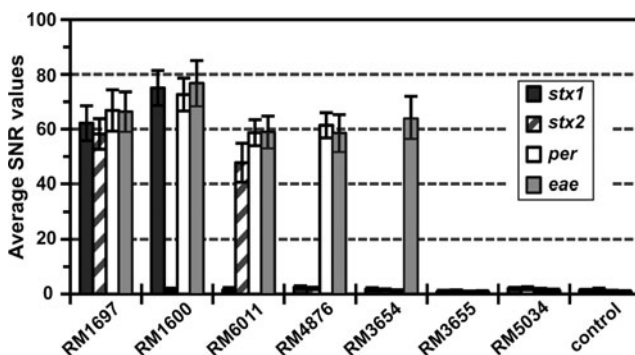
The sensitivity threshold for identifying pathogenic *E. coli* by using DNA microarrays with photopolymerization was further investigated. Various cell concentrations of *E. coli* O157 strain RM1697 were tested on the microarray, and the SNR values were quantified after performing the photo-

polymerization reaction. The limit of detection was defined by the findings from validation experiments for specificity (Fig. 2), demonstrating that spots on the array without specific formation of polymer yielded SNR values  $< 2.5$ . In addition, previous studies have shown that an SNR value of 3 is considered to be the minimum signal intensity for accurate quantification (Verdnick *et al.*, 2002; He and Zhou, 2008; Park *et al.*, 2008). Based on these observations an SNR value of 3 was established as our detection limit for positive signals (Fig. 3), and the results from the sensitivity experiments demonstrated that the threshold for detecting strain RM1697 by using DNA microarrays with photopolymerization was determined to be between 100 and 1000 CFU/mL (Fig. 3).

## Discussion

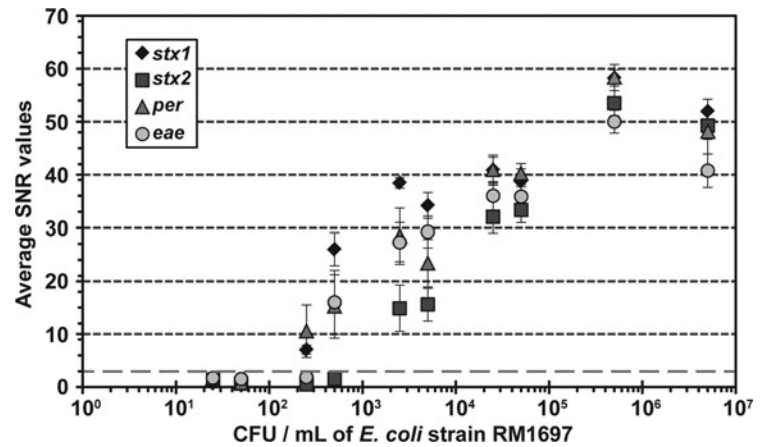
Outbreaks of foodborne disease caused by *E. coli* O157 continue to cause concern among consumers, regulatory agencies, and the food industry (Mandrell, 2009; Buchanan and Appel, 2010). To address this issue, risk management efforts have focused on identifying the presence and distribution of *E. coli* O157 in foods due to large outbreaks and severe illnesses associated with this particular serotype (Karmali, 2009; Buchanan and Appel, 2010). For use in surveillance and outbreak epidemiology, there is a need for the development of rapid, simple, and inexpensive assays for pathogen detection. Molecular-based technologies, such as DNA microarrays, offer an alternative to screen simultaneously for multiple virulence markers and to assist in the identification of the virulence types of foodborne pathogens (Call, 2005; Uttamchandani *et al.*, 2009).

For pathogen detection with DNA microarrays, procedures that enzymatically incorporate a fluorescent dye during DNA amplification are most commonly used due to the availability of reagents and instrumentation (Call, 2005; Vora *et al.*, 2008; Uttamchandani *et al.*, 2009). However, these fluorescent assays can lack consistency in the labeling of the DNA and may require procedures that are time consuming and use expensive scanners for data analysis (Kuck and Taylor, 2008; Vora *et al.*, 2008). Alternative methods that are colorimetric show more uniformity in DNA labeling. A limitation of these colorimetric assays, measuring alkaline phosphatase activity or silver precipitation, is that they employ posthybridization re-



**FIG. 2.** Quantification of signals on the DNA microarray. *E. coli* reference strains were tested by using DNA microarray and ampliPHOX detection (see Materials and Methods section). For each spot on the microarray, the signals for each spot on the microarray were quantified and expressed as signal-to-noise ratio (SNR). Each bar represents the average SNR values  $\pm$  standard deviations of three independent experiments with triplicate measurements for each probe.

**FIG. 3.** Sensitivity of *E. coli* O157 detection with DNA microarrays with ampliPHOX detection. Various cell concentrations of the reference *E. coli* O157 strain RM1697 were tested on the DNA microarray, and the SNR values were quantified after photopolymerization. The dashed line delineates the detection limit for positive signals, and each symbol represents the average SNR values  $\pm$  standard deviations from three independent experiments with triplicate measurements for each probe.



agents that result in high variability in signal detection (Kuck and Taylor, 2008). Given the limitations of current procedures, the present study evaluated photopolymerization for identification of pathogenic *E. coli*, using low-cost reagents and instrumentation that are amenable to high-throughput sampling and routine pathogen surveillance (Kuck and Taylor, 2008). Previously, photopolymerization was employed successfully for molecular diagnostic assays detecting *Streptococcus* species (Dawson *et al.*, 2009) and for genotyping influenza viruses (Kuck and Taylor, 2008). Real-time PCR methods have been developed for the identification of virulent strains of *E. coli* O157 (Kawasaki *et al.*, 2010; Suo *et al.*, 2010); however, the cost for the reagents and instrumentation required for these analyses with real-time PCR impede the use of this molecular method for routine strain characterization in smaller research facilities. To provide a proof of concept for testing photopolymerization for foodborne pathogen detection with DNA microarrays, the present study constructed a low-density DNA microarray targeting *stx1*, *stx2*, *eae*, and *per*, genes encoding virulence factors that are considered to be good indicators of pathogenic *E. coli* O157.

Validation experiments demonstrated that the use of DNA microarrays with photopolymerization for detection of *E. coli* O157 resulted in an accurate genotyping of the tested strains. Polymer formation was clearly evident only for those probes targeting virulence genes known to be present in the reference strains. Quantification of the specificity by measuring the SNR values for each probe on the array demonstrated that there was a clear distinction between signal intensities and background noise, an important aspect for data analysis and interpretation of results with DNA microarrays (Verdnick *et al.*, 2002; He and Zhou, 2008). These results were achieved by using a 30-mer oligonucleotide probe allowing a more specific detection of the target gene. Previous studies that assessed optimal conditions for fluorescent detection on DNA microarrays demonstrated that the use of short, 30-mer probes were equivalent to background noise, whereas optimal hybridization signals required 70-mer or 100-mer oligonucleotide probes (Letowski *et al.*, 2004). However, the use of longer probes resulted also in lower specificity due to cross-hybridization with nontarget genes in some species (Letowski *et al.*, 2004). In the present study, the use of 30-mer probes on the microarray in combination with photopolymerization facilitated the detection of specific target regions in the genes selected for identifying *E. coli* O157. Our quantification analysis showed that positive signal intensities

had SNR values that were at least 20 times higher than spots without any specific hybridization.

Further, our analyses demonstrated consistently that the sensitivity of *E. coli* O157 detection by using photopolymerization was found to be 100–1000 CFU/mL in all validation experiments measuring assay sensitivity. This level of sensitivity was similar to a previous report of *E. coli* O157 detection with fluorescent methods (Call *et al.*, 2001). In the present study, detection at cell concentrations <100 CFU/mL was variable (data not shown), due potentially to an effect of PCR inhibitors or to inefficient cell recovery during the assay, as reported previously (Call *et al.*, 2001). Thus, the results from the present study demonstrated that the sensitivity of *E. coli* O157 detection by using photopolymerization with DNA microarrays was similar to fluorescence. Previously, similar detection sensitivities for photopolymerization and fluorescence were reported for influenza A virus identification with a low-density microarray (Kuck and Taylor, 2008). Additionally, our study demonstrated that photopolymerization resulted in the rapid genotyping of *E. coli* O157 in only 2–4 h, a significant improvement when compared to our results with fluorescence detection methods that required several days for virulent strain identification (Quiñones *et al.*, 2007). Future work is aimed at further development of DNA microarrays with photopolymerization by adding pathogen-specific probes for identifying virulent strains of *E. coli* O157 and non-O157 from distinct geographical locations and sources. The pathogen-specific probes on the expanded DNA microarrays will target several non-O157 serovars implicated in severe human disease (Brooks *et al.*, 2005; Bettelheim, 2007; Karmali, 2009; Buchanan and Appel, 2010). Additional probes will be included to target virulence determinants that code for adhesins, proteases, cytotoxins, and effectors and that are considered to be necessary for *E. coli* strains to be pathogenic (Bettelheim, 2007; Karmali, 2009; Buchanan and Appel, 2010).

## Conclusions

Photopolymerization is a simple, rapid, and quantitative DNA microarray-based detection method for assessing the genetic composition of *E. coli* O157 strains by examining genes encoding Shiga toxin and selected virulence determinants that have been associated with pathogenic *E. coli*. This approach facilitates the evaluation of the potential virulence of *E. coli* O157 strains and provides relevant information to the food

industry for determining the prevalence and risks of pathogenic *E. coli* in food production environments.

### Acknowledgments

This work was supported by the U.S. Department of Agriculture, Agricultural Research Service CRIS project number 5325-42000-045, and by the National Institutes of Health/National Institute of Allergy and Infectious Diseases grant number U01A1070276. The authors would like to thank Edith Pierre-Jerome for excellent technical assistance and to Robert E. Mandrell and Michael Cooley (USDA/ARS/WRRRC, Albany, CA) for providing the bacterial strains used in this study.

### Disclosure Statement

B.Q. and M.S.S. declare no competing financial interests. E.D.D. and A.W.T. are both paid employees of InDevR, Inc., a for-profit entity.

### References

- Bettelheim KA. The non-O157 Shiga-toxigenic (verocytotoxigenic) *Escherichia coli*; under-rated pathogens. *Crit Rev Microbiol* 2007;33:67–87.
- Bettelheim KA and Beutin L. Rapid laboratory identification and characterization of verocytotoxigenic (Shiga toxin producing) *Escherichia coli* (VTEC/STEC). *J Appl Microbiol* 2003;95:205–217.
- Bilge SS, Vary JC Jr., Dowell SF, and Tarr PI. Role of the *Escherichia coli* O157:H7 O side chain in adherence and analysis of an *rfb* locus. *Infect Immun* 1996;64:4795–4801.
- Boissinot K, Huletsky A, Peytavi R, Turcotte S, Veillette V, Boissinot M, Picard FJ, Martel EA, and Bergeron MG. Rapid exonuclease digestion of PCR-amplified targets for improved microarray hybridization. *Clin Chem* 2007;53:2020–2023.
- Brooks JT, Sowers EG, Wells JG, Greene KD, Griffin PM, Hoekstra RM, and Strockbine NA. Non-O157 Shiga toxin-producing *Escherichia coli* infections in the United States, 1983–2002. *J Infect Dis* 2005;192:1422–1429.
- Buchanan RL and Appel B. Combining analysis tools and mathematical modeling to enhance and harmonize food safety and food defense regulatory requirements. *Int J Food Microbiol* 2010;139 Suppl 1:S48–S56.
- Call DR. Challenges and opportunities for pathogen detection using DNA microarrays. *Crit Rev Microbiol* 2005;31:91–99.
- Call DR, Brockman FJ, and Chandler DP. Detecting and genotyping *Escherichia coli* O157:H7 using multiplexed PCR and nucleic acid microarrays. *Int J Food Microbiol* 2001;67:71–80.
- Cooley M, Carychao D, Crawford-Miksza L, Jay MT, Myers C, Rose C, Keys C, Farrar J, and Mandrell RE. Incidence and tracking of *Escherichia coli* O157:H7 in a major produce production region in California. *PLoS One* 2007;2:e1159.
- Dawson ED, Taylor AW, Smagala JA, and Rowlen KL. Molecular detection of *Streptococcus pyogenes* and *Streptococcus dysgalactiae* subsp. *equisimilis*. *Mol Biotechnol* 2009;42:117–127.
- He Z and Zhou J. Empirical evaluation of a new method for calculating signal-to-noise ratio for microarray data analysis. *Appl Environ Microbiol* 2008;74:2957–2966.
- Karmali MA. Host and pathogen determinants of Verocytotoxin-producing *Escherichia coli*-associated hemolytic uremic syndrome. *Kidney Int Suppl* 2009;(112):S4–S7.
- Kawasaki S, Fratamico PM, Horikoshi N, Okada Y, Takeshita K, Sameshima T, and Kawamoto S. Multiplex real-time polymerase chain reaction assay for simultaneous detection and quantification of *Salmonella* species, *Listeria monocytogenes*, and *Escherichia coli* O157:H7 in ground pork samples. *Foodborne Pathog Dis* 2010;7:549–554.
- Kimura R, Mandrell RE, Galland JC, Hyatt D, and Riley LW. Restriction-site-specific PCR as a rapid test to detect enterohemorrhagic *Escherichia coli* O157:H7 strains in environmental samples. *Appl Environ Microbiol* 2000;66:2513–2519.
- Kuck LR and Taylor AW. Photopolymerization as an innovative detection technique for low-density microarrays. *Biotechniques* 2008;45:179–186.
- Letowski J, Brousseau R, and Masson L. Designing better probes: effect of probe size, mismatch position and number on hybridization in DNA oligonucleotide microarrays. *J Microbiol Methods* 2004;57:269–278.
- Mandrell RE. Enteric humans pathogens associated with fresh produce: sources, transport, and ecology. In: *Microbial Safety of Fresh Produce*. Fan X, Niemira B, Doona CJ, Feeherry F, Gravani RB (eds.). Ames, IA: IFT Press/Wiley-Blackwell Publishing, 2009, pp. 5–41.
- Nataro JP and Kaper JB. Diarrheagenic *Escherichia coli*. *Clin Microbiol Rev* 1998;11:142–201.
- Park SJ, Kang CH, Chae JC, and Rhee SK. Metagenome microarray for screening of fosmid clones containing specific genes. *FEMS Microbiol Lett* 2008;284:28–34.
- Quiñones B, Massey S, Friedman M, Swimley MS, and Teter K. Novel cell-based method to detect Shiga toxin 2 from *Escherichia coli* O157:H7 and inhibitors of toxin activity. *Appl Environ Microbiol* 2009;75:1410–1416.
- Quiñones B, Parker CT, Janda JM Jr., Miller WG, and Mandrell RE. Detection and genotyping of *Arcobacter* and *Campylobacter* isolates from retail chicken samples by use of DNA oligonucleotide arrays. *Appl Environ Microbiol* 2007;73:3645–3655.
- Ramachandran V, Brett K, Hornitzky MA, Downton M, Bettelheim KA, Walker MJ, and Djordjevic SP. Distribution of intimin subtypes among *Escherichia coli* isolates from ruminant and human sources. *J Clin Microbiol* 2003;41:5022–5032.
- Rangel JM, Sparling PH, Crowe C, Griffin PM, and Swerdlow DL. Epidemiology of *Escherichia coli* O157:H7 outbreaks, United States, 1982–2002. *Emerg Infect Dis* 2005;11:603–609.
- Reeves PR, Hobbs M, Valvano MA, Skurnik M, Whitfield C, Coplin D, Kido N, Klena J, Maskell D, Raetz CRH, and Rick PD. Bacterial polysaccharide synthesis and gene nomenclature. *Trends Microbiol* 1996;4:495–503.
- Rozen S and Skaletsky H. Primer3 on the WWW for general users and for biologist programmers. In: *Bioinformatics Methods and Protocols*, Volume 132. Misener S and Krawetz SA (eds.). Totowa, NJ: Humana Press, 2000, pp. 365–386.
- Serna A and Boedeker EC. Pathogenesis and treatment of Shiga toxin-producing *Escherichia coli* infections. *Curr Opin Gastroenterol* 2008;24:38–47.
- Shimizu T, Yamasaki S, Tsukamoto T, and Takeda Y. Analysis of the genes responsible for the O-antigen synthesis in enterohaemorrhagic *Escherichia coli* O157. *Microb Pathog* 1999;26:235–247.
- Sikes HD, Hansen RR, Johnson LM, Jenison R, Birks JW, Rowlen KL, and Bowman CN. Using polymeric materials to generate an amplified response to molecular recognition events. *Nat Mater* 2008;7:52–56.
- Suo B, He Y, Tu SI, and Shi X. A multiplex real-time polymerase chain reaction for simultaneous detection of *Salmonella* spp.,

- Escherichia coli* O157, and *Listeria monocytogenes* in meat products. *Foodborne Pathog Dis* 2010;7:619–628.
- Uttamchandani M, Neo JL, Ong BN, and Moochhala S. Applications of microarrays in pathogen detection and biodefence. *Trends Biotechnol* 2009;27:53–61.
- Verdnick D, Handran S, and Pickett S. Key considerations for accurate microarray scanning and image analysis. In: *DNA Image Analysis: Nuts and Bolts*. Kamberova G (ed.). Salem, MA: DNA Press, 2002, pp. 83–98.
- Vora GJ, Meador CE, Anderson GP, and Taitt CR. Comparison of detection and signal amplification methods for DNA microarrays. *Mol Cell Probes* 2008;22:294–300.
- Zhang WL, Kohler B, Oswald E, Beutin L, Karch H, Morabito S, Caprioli A, Suerbaum S, and Schmidt H. Genetic diversity of intimin genes of attaching and effacing *Escherichia coli* strains. *J Clin Microbiol* 2002;40:4486–4492.

Address correspondence to:  
Beatriz Quiñones, Ph.D.  
Produce Safety and Microbiology Research Unit  
Western Regional Research Center  
Agricultural Research Service  
U.S. Department of Agriculture  
800 Buchanan St., WAB214L  
Albany, CA 94710

E-mail: [beatriz.quinones@ars.usda.gov](mailto:beatriz.quinones@ars.usda.gov)

