

A New Lab Developed Real Time PCR Assay for Direct Detection of *C. Difficile* from Stool Sample without DNA Extraction

Brandon Li

Sino US Gene, Inc. 1452 #C, West Holt Ave. Pomona, CA 91768, USA

ABSTRACT

Clostridium difficile is a major cause of nosocomial antibiotic-associated infectious diarrhea and pseudomembranous colitis. Detection of *C. difficile* by anaerobic bacterial culture and/or cytotoxicity assays has been largely replaced by rapid enzyme immunoassays (EIA). However, due to the lack of sensitivity of stool EIA, we developed a multiplex real-time PCR assay targeting the *C. difficile* toxin genes *tcdB*. stool samples from hospitalized pediatric patients suspected of having *C. difficile*-associated disease were prospectively collected. Three testing modalities were evaluated, including enriched culture, cepheid Xpert and real-time Pcr (*tcdB*) on stool samples performed with *tcdB* gene-specific primers and hydrolysis probes. A total of 150 de-identified clinical specimen were analyzed. The sensitivities of stool real-time Pcr were 95% against cepheid Xpert *C. difficile* and 93% against enriched culture respectively, with a specificity of 97% and 94%. The lower limit of detection of the stool real-time PCR was 0.5 cFU/ml of per reaction for *tcdB*. Direct detection of *C. difficile* toxin genes in stool samples by real-time Pcr showed performance comparable to enriched culture. Real-time PCR of DNA from stool samples is a rapid and cost-effective diagnostic modality for patients that should facilitate appropriate patient management. (*Int J Biomed Sci* 2016; 12 (3): 83-88)

Keywords: *Clostridium difficile*; real-time PCR; enzyme immunoassays; *tcdB*

Corresponding author: Brandon Li, Sino US Gene, Inc. 1452 #C, West Holt Ave. Pomona, CA 91768, USA. Tel: 1-(626)-282-8661; E-mail: brandonl@usc.edu.

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INTRODUCTION

Clostridium difficile, a Gram-positive spore-forming bacillus, is the most common identifiable etiologic agent of antibiotic-associated diarrhea (13, 18). Initially described as a member of the commensal microbiota of neonates, *C. difficile* was identified as a causal agent of antibiotic-associated diarrhea in the 1970s (3, 11). The clinical presentation of *C. difficile*-associated disease (CDAD) can range from asymptomatic carriage in the gastrointestinal tract, mild diarrhea, and potentially fatal pseudomembranous colitis (13, 18). Symptoms occur secondary to the production of two exotoxins, toxin A and toxin B, which disrupt the integrity of the colonic mucosa (24).

Alarming changes in the epidemiology of CDAD, including an increase in both the incidence and severity of the disease, have highlighted concerns about patterns of *C. difficile* infection (17, 18, 19, 21). Analysis of U.S. hospital discharge data revealed that the national rates of CDAD doubled from 2000 to 2003 (17). In 2004, the Centers for Disease Control and Prevention reported that the mortality rate related to CDAD increased from 5.7 deaths per million individuals in 1999 to 23.7 deaths per million individuals (21). In addition to the profound morbidity and mortality, CDAD is also generating a substantial economic burden, with estimates ranging from \$1.3 million to more than \$3 billion annually (8, 16, 20). Due to the formidable impact of CDAD on the U.S. health care system, rapid and accurate diagnosis is essential for the timely enactment of infection control and treatment measures.

The changing epidemiology of *C. difficile* infections in the pediatric population is a serious concern. While benign neonatal colonization with toxigenic *C. difficile* is a well-documented phenomenon, recent studies have suggested an increased incidence of CDAD in children (2, 14, 22, 25). A large study encompassing data collected from 22 children's hospitals in the United States reported an increased prevalence of CDAD in children, including infants (increased by 53% from 2001 to 2006, with 26% of patients with CDAD ≤ 1 year of age) (14). Utilizing CDAD data from the Agency for Healthcare and Research Quality, a similar study noted that the highest number of CDAD hospitalizations occurred in patients ≤ 1 year of age (25).

Initial strategies to detect *C. difficile* consisted of anaerobic stool sample culture, usually with cycloserine-cefoxitin-fructose agar (CCFA) or a similar medium with or without a pretreatment alcohol shock step (7). Although this modality was quite sensitive and specific for detecting *C. difficile*, it took up to 5 days to confirm a negative

culture and it did not discriminate between toxigenic and nontoxigenic isolates without further testing strategies. Furthermore, colonies with indeterminate colony characteristics were tested with L-proline-aminopeptidase (PRO Disc) or other biochemical tests to ensure the accurate identification of *C. difficile* (9, 10). The development of the cell culture cytotoxicity assay circumvented stool sample culture by observing cytopathic effects of toxin B directly on cultured cells (4, 6). The cell culture cytotoxicity assay requires a neutralization step for specificity and maintenance of toxin-susceptible mammalian cell lines, and it takes 48 to 72 h to perform the assay (1, 5). Rapid antigen detection assays, consisting of common antigen testing (glutamate dehydrogenase) and toxin immunoassays, have largely replaced culture and the cytotoxic assay; however, neither type has the desired sensitivity or specificity to reliably confirm or rule out CDAD without the need for either serial testing or subsequent testing modalities. Therefore, real-time PCR is being investigated as the preferred diagnostic modality due to its rapid turnaround time and track record of superior sensitivity and specificity.

Toxigenic strains of *C. difficile* contain a 19.6-kb pathogenicity locus (PaLoc) that includes five contiguous chromosomal genes responsible for the development of CDAD—*tcdABCDE* (24). *tcdA* and *tcdB* encode exotoxins A (enterotoxin) and B (cytotoxin), respectively; *tcdC* and *tcdD* encode negative and positive regulators, respectively, that control the level of toxin production; and *tcdE* is purported to encode a holin-like protein thought to facilitate toxin release from the bacterial cell wall (24). Because toxins A and/or B are implicated in CDAD and genetic diversity of the PaLoc has been reported (23), we developed and clinically validated one hydrolysis probe real-time PCR assays targeting the *tcdB* genes (12, 15, 24). While the molecular methods utilized by this assay were not novel, the application of molecular testing for *C. difficile* infection is unique when the stool sample could be tested directly without nuclear acid extraction. This will greatly facilitate quick testing of *C. difficile* in clinical setting.

MATERIALS AND METHOD

***C. difficile* Strains:** The following strains were used for LoD study: *C. difficile* ATCC 43255 (ZeptoMetrix), *C. difficile* NAP1A (ZeptoMetrix).

Extraction, Real-time PCR Amplification and Detection: Lab developed *C. difficile* Direct Kit contains all reagents for on-board extraction and real-time PCR amplification.

Fifty µL of *C. difficile* Direct reaction mix was loaded into the reaction port and 50 µL of sample was directly loaded into the sample port on the Amplification cell. All testing was performed using real time PCR. Assay time is about 60 minutes.

Limit of Detection (LoD): The LoD for each *C. difficile* stock was determined as the lowest concentration with ≥95% detection in negative stool matrix.

Reproducibility: Thirty-six replicates of the following contrived panel in negative stool matrix were tested: *C. difficile* Low Positive (ATCC 43255), *C. difficile* Medium Positive (ATCC 43255), *C. difficile* Low Positive (NAP1A), *C. difficile* Medium Positive (NAP1A). Low Positive was defined as 1X LoD; medium positive was defined as 3X LoD.

Positive and Negative Agreement: A panel of 150 de-identified clinical specimens was evaluated using the Lab developed *C. difficile* Direct assay. Lab developed test results were compared to Cepheid Xpert *C. difficile* and enriched culture results.

Cross-Reactivity: The cross-reactivity panel of 126 different organisms consisted of industry equivalent 106 CFU/mL of bacteria or 105 TCID₅₀/mL of virus in negative stool matrix.

Inhibition/interference: The interference panel was contrived with the ATCC 43255 or NAP1A strain at 4-fold the LoD concentration. Each substance was spiked into the *C. difficile* contrived stool samples and tested using Lab developed *C. difficile* Direct.

RESULTS

Limit of Detection: *C. difficile* ATCC 43255 LoD was 0.5 CFU/mL. *C. difficile* NAP1A strain LoD was 1.6 CFU/mL in stool matrix (Table 1).

***C. difficile* Reproducibility:** For ATCC 43255 and NAP1A medium- and low-contrived panels, *C. difficile* strains were detected in 100% of replicates. Standard deviations were <0.99. Percent coefficients of variation were <3.3 (Table 2).

Table 1. *C. difficile* Limit of Detection

Bacterial Strain	(LoD) Concentration	Detection Rate	Average Ct	Maximum Ct	Minimum Ct
ATCC 43255	0.5 CFU/mL	95% (19/20)	39.1	41	38.5
NAP1A	1.3 CFU/mL	100% (20/20)	39.1	42.0	38.4

Table 2. Lab developed *C. difficile* Direct Quantitative Reproducibility

Channel/ Detector	Sample Name	N	Mean Ct	Between Instrument		Between Operator		Between Run		Within Run		Total	
				SD	%CV	SD	%CV	SD	%CV	SD	%CV	SD	%CV
C. diff (FAM)	Low Pos 43255	36	38.7	0	0	0	0	0	0	0.87	2.6	0.94	2.7
	Low Pos NAP1A	36	38.8	0	0	0	0	0.7	1	0.76	1.8	0.74	1.9
	Med Pos 43255	36	38	0.31	0.9	0	0	0	0	0.53	2	0.83	2.3
	Med Pos NAP1A	36	37.2	0.16	0.2	0	0	0	0	0.58	1.7	0.52	1.2
	Pos Control	36	31.2	0.27	0.5	0	0	0.1	0.3	0.37	1.6	0.38	1.6
IC (Q670)	Low Pos 43255	36	29.9	0	0	0	0	0	0	0.87	3.1	0.99	3.1
	Low Pos NAP1A	36	29.8	0.28	0.9	0	0	0	0	0.74	2.3	0.76	2.8
	Med Pos 43255	36	29.6	0.26	0.5	0	0	0	0	0.62	2.3	0.69	2.4
	Med Pos NAP1A	36	29.9	0.35	1.5	0	0	0	0	0.6	2.4	0.75	2.6
	Negative	36	30.1	0.12	0.4	0	0	0.22	0.7	0.79	2.8	0.81	2.5
	Pos Control	36	30.8	0.33	1	0	0	0	0	0.7	2.2	0.76	2.9

C. difficile Positive and Negative Agreement: Results from Lab developed C. difficile Direct and Cepheid Xpert C. difficile were in agreement for 95% of positive specimens and 97% of negative specimens (Tables 3 and 4).

Results from Lab developed C. difficile Direct and enriched culture were in agreement for 93% of positive spec-

imens and 94% of negative specimens (Tables 3 and 5).

Cross-Reactivity: No cross-reactivity was detected with the 126 pathogens tested (subset of representative strains listed in Table 6).

Substance Interference: No interference was detected with the substances tested (Table 7).

Table 3. Lab developed C. difficile Direct Positive and Negative Agreement

	Lab developed <i>C. difficile</i> Direct vs Cepheid Xpert <i>C. difficile</i> (n=150)	Lab developed <i>C. difficile</i> Direct vs Enriched Culture (n=110)
Positive Agreement (Sensitivity)	95%	93%
Negative Agreement (Specificity)	97%	94%

Table 4. Lab developed C. difficile Direct Agreement with Cepheid Xpert C. difficile

		Cepheid Xpert <i>C. difficile</i>		Total
		Positive	Negative	
Lab developed <i>C. difficile</i> Direct	Positive	32	3	35
	Negative	2	113	115
	Total	34	116	150

Table 5. Lab developed C. difficile Direct Agreement with Enriched Toxigenic Culture

		<i>C. difficile</i> Enriched Toxigenic Culture		Total
		Positive	Negative	
Lab developed <i>C. difficile</i> Direct	Positive	41	6	47
	Negative	3	100	103
	Total	44	106	150

Table 6. C. difficile Cross-Reactivity Pathogens Tested in Stool Matrix (Representative Strains)

<i>Abiotrophia defectivae</i>	<i>Candida catenulate</i>	<i>Clostridium septicum</i>
<i>Acinetobacter baumannii</i>	<i>Clostridium bifermentans</i>	<i>Clostridium tetani</i>
<i>Acinetobacter Iwoffii</i>	<i>Clostridium bolteae</i>	<i>Clostridium difficile</i> (non-toxigenic ATCC43593)
<i>Aeromonas hydrophila</i>	<i>Clostridium butyricum</i>	<i>Desulfovibrio piger</i>
<i>Alcaligenes faecalis</i> subsp. <i>Faecalis</i>	<i>Clostridium chauvoei</i>	<i>Edwardsiella tarda</i>
<i>Bifidobacterium longum</i>	<i>Clostridium fallax</i>	<i>Eggerthella lenta</i>
<i>Campylobacter coli</i>	<i>Clostridium ramosum</i>	<i>Enterobacter aerogenes</i>
<i>Campylobacter jejuni</i> sub sp. <i>jejuni</i>	<i>Clostridium scindens</i>	<i>Enterobacter cloacae</i>
<i>Candida albicans</i>		

Table 7. Interferents Tested in Stool Matrix

Substance	Active Ingredient	Starting Interferent Concentration	Final Sample Concentration ^a
Antacid and anti-gas generic	Aluminum hydroxide, magnesium hydroxide	1 mg/mL	0.1 mg/mL
Milk of Magnesia (liquid)	Magnesium hydroxide	2 mg/mL	0.2 mg/mL
Antacid generic	Calcium carbonate	1 mg/mL	0.1 mg/mL
Metronidazole	Metronidazole	140 mg/mL	14 mg/mL
Vancomycin	Vancomycin	14 mg/mL	1.4 mg/mL
Stearic acid	Stearic acid	40 mg/mL	4 mg/mL
Palmitic acid	Palmitic acid	20 mg/mL	2 mg/mL
Barium sulfate	Barium sulfate	50 mg/mL	5 mg/mL
Loperamide hydrochloride generic	Loperamide	0.05 mg/mL	0.005 mg/mL
Pepto-Bismol (liquid)	Bismuth subsalicylate	1.75 mg/mL	0.175 mg/mL
Preparation H	Phenylephrine	20% (w/v)	2% (w/v)
Trojan with nonoxynol-9	Nonoxynol-9	14 mg/mL	1.4 mg/mL
1% hydrocortisone cream generic	Hydrocortisone	20% (w/v)	2% (w/v)
Fleet	Mineral oil	20% (w/v)	2% (v/v)
Laxative generic	Sennosides	1 mg/mL	0.1 mg/mL
Moist towelettes generic	Benzalkonium chloride	100% (v/v)	10% (v/v)
KY Jelly	Glycerin	20% (w/v)	2% (w/v)
Nystatin (6,000 USP units/mg)	Nystatin	100,000 USP/mL	10,000 USP units/mL
Naproxen sodium generic	Naproxen	140 mg/mL	14 mg/mL
Mucin	Mucin	30 mg/mL	3 mg/mL
Whole blood (donor: RH)	Whole blood	50% (v/v)	5% (v/v)
TE (spiked baseline)	None	N/A	N/A

^aFor each substance, 100 µL of interferent at starting interferent concentration was added to 900 µL of stool matrix.

CONCLUSION

Lab developed C. difficile Direct can provide an option for simplified C. difficile testing on the real time PCR. This test was also comparable to both Xpert C. difficile and enriched toxigenic culture for identifying C. difficile.

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