Optimization of Spore Coat and Exosporium extracts removal from *Clostridioides difficile* Spores.

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Running Title: Spore surface layer extraction

Key Words: *C. difficile* spores, spore adherence, spore entry, aging, intestinal ligated loop, ileal loop, colonic loop, intestinal mucosa.

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Abstract

Analysis of the protein composition that make up *C. difficile* spores is essential in understanding

the protein composition of its surface layers however the spore surface is composed of protein

complexes hindering protein analysis. Here we compare commonly used protein extraction

methods used to extract proteins of *C. difficile* spores. Our results provide a systematic analysis of

how two commonly used spore coat / exosporium extraction methods compare in efficiency of

protein removal and spore structural analysis.

Introduction

Clostridioides difficile (*C. difficile*) is a Gram-positive anaerobe that has become a leading cause of antibiotic associated diarrhea in developed countries[1,2]. Treatment against *C. difficile* infections (CDI) resolve 95 % of primary cases; however, ~15 – 30 % of recovered CDI patients develop subsequent recurrent episodes of CDI, which result in mortality rates increasing up to 30%[3]. Although two major clostridial toxins, TcdA and TcdB, are essential for disease manifestation[4], the production of *C. difficile* spores during disease is essential for disease recurrence[5,6].

C. difficile spores are metabolically dormant and naturally resistant to antibiotics, heat, ethanol, and various cleaning agents[6,7]. *C. difficile* spores are made by a series of concentric layers that assemble under the control of the mother cell developmental program, specifically under the RNA polymerase sporulation-specific sigma factors, SigE and SigK [6,7]. The outermost exosporium layer of *C. difficile* spores play important roles in infection recurrence and interaction with host molecules [8-12]. Importantly, during late stages of sporulation development, *C. difficile* forms two distinctive spores that differ in the thickness of the exosporium morphotype, while the underlying spore coat remains identical [6,13,14].

Both, the spore coat and exosporium of *C. difficile* spores are proteinaceous layers made of > 50 and ~ 10 - 20 proteins, respectively[15,16]. Both layers reassemble a highly crosslinked protein matrix, and while the exosporium layer can be easily removed by protease digestions (i.e., trypsin or proteinase-k) while the spore coats remain intact[17]. Despite the easiness to proteolytically remove the exosporium layer, it is unclear whether the spore coat and exosporium layer are efficiently removed through chemical methods, commonly employed in *C. difficile* spore-research. Therefore,

in this work, our aim is to provide a systematic analysis of how two chemical methods (i.e., USD

and EBB) that are commonly employed to remove the spore coat /exosporium extracts for

immunoblotting analysis, remove these layers and impact the functional aspects of *C. difficile*

spores. For this, we utilized two exosporium markers (i.e, CdeC and CdeM)[10, 18, 19], and the

peptidoglycan-spore coat interface specific spore cortex lytic enzyme SleC[20, 21], to monitor the

efficiency of extraction of these layers through immunoblotting and transmission electron

microscopy.

Methods

Spore Preparation and Purification.

C. difficile strain R20291 was grown in Brain Heart Infusion broth (BHIS) (Difco) supplemented

with 0.5% yeast extract and 0.1% cysteine anaerobically at 37˚C. Overnight cultures were grown

in BHIS broth with 0.1% sodium taurocholate and 0.2% d-fructose. Cultures were diluted to an

OD_{600} of 0.5 and 250 uL was plated on 70:30 sporulation media under anaerobic conditions for 5

days at 37˚C. Sporulating cultures were collected from plates, resuspended in ice-cold distilled

water, and left in 4˚C overnight to allow for lysis of vegetative cells. After incubation, cells were

centrifuged and pellets washed with ice-cold distilled water 5 times followed by repeated

centrifugation and resuspension to remove vegetative cell debris. Spores were separated using a

60% sucrose gradient and centrifuged for 20 min at 14,000 rpm spores were then washed with ice-

cold water to remove residual sucrose. Washing and sucrose gradient repeated until spores reached

99% purity. Once pure, spores were quantified using a NeuBauer chamber and stored at -80˚C

until use at a $5x10^9$ spores/mL concentration.

Spore Extraction Methods

1.5x10^7 spores pelleted down and resuspended in 50 uL of buffer. Samples with EBB (EBB 8M

Urea, 2M Thiourea, 4% w/v SDS, 2% w/v β-mercaptoethanol) were boiled 10 minutes and spun

down at 15k rpm for 5 minutes. Supernatant containing spore extract was saved and spore pellet

washed three times to remove residual spore extract. Spore was then retreated with EBB for a total

of three extraction with supernatant saved between each step. USD (8M Urea, 1% w/v SDS, 50

mM DTT, 50 mM Tris-HCl (pH 8)samples were treated for 90 minutes at 37˚C and pelleted down.

Supernatant was saved and spore pellet washed. 1˚ USD or 1˚ treated spores were incubated with

lysozyme for 37˚C overnight for 24 hours.

Lysozyme Treatment

C. difficile spores were treated with lysozyme to analyze the effects of the extraction buffers on

the integrity of the spore coat. Spores ($1x10^7$ spores) were treated with 10mg/mL lysozyme in 25

mM phosphate buffer (pH 7.4) at 37˚C for 2 hours. Samples were pelleted and washed with

distilled water and analyzed under phase contrast microscopy.

Transmission Electron Microscopy

For transmission electron microscopy, *C. difficile* spores (10^{10}) were fixed overnight with 3%

glutaraldehyde, 0.1 M cacodylate buffer (pH 7.2) at 4˚C. Samples were centrifuged at 14k rpm for

5 minutes and supernatant discarded. They were then stained with1% osmium tetroxide in 0.05M

HEPES buffer (pH 7.4) overnight at 4°C. Treated samples were washed 5x with distilled water.

The spores were dehydrated stepwise in 30%, 50%, 70%, 90% for 15 minutes respectively,

followed by dehydration with 100% acetone 3 times for 30 minutes at each step. A small amount

of acetone is left covering the sample to prevent rehydration. Spore samples were embedded in

modified Spurr's resin (Quetol ERL 4221 resin; EMS; RT 14300) in a Pelco Biowave processor

(Ted Pella, Inc.). Initially, 1:1 acetone-resin for 10 min at 200W-with no vacuum, 1:1 acetone-

resin for 5 min at 200W-vacuum 20"Hg, followed by 100% resin 4 times at 200W for 5 minutes-

vacuum 20"Hg. The resin was removed, and sample pellet was transferred to BEEM conical-tip

capsule and filled with 100% fresh modified Spurr's resin. Sample was left to reach bottom of

capsule and subsequently left to polymerize for 48 hours in a 65°C oven followed by 24 hours at

room temperature. Ultrathin section ~100nm (silver-gold color) were obtained using Leica UC7

Ultramicrotome and placed on glow-discharged carbon coated 300-mesh Cu grids. Grids were

double lead stained with 2% uranyl acetate for 5 minutes and washed with filter sterilized (0.2uM

filter) distilled water   followed by 5 minute staining with Reynold's lead citrate and subsequent

washing as described. All ultrathin TEM sections were imaged on a JEOL 1200 EX TEM (JEOL,

Ltd.) at 100 kV, and images were recorded on an SIA-15C charge-coupled device (CCD)

(Scientific Instruments and Applications) camera at the resolution of 2,721 by 3,233 pixels using

MaxImDL software (Diffraction Limited). All equipment used is located at the Texas A and M

University Microscopy and Imaging Center Core Facility (RRID: SCR_022128).

Western Blot Analysis

Protein samples were resuspended in 2x SDS-PAGE loading buffer (BioRad) with 5% b-

mercaptoethanol and boiled for 10 minutes. Samples were run in a 12% acrylamide SDS-PAGE

gels. Proteins were transferred to nitrocellulose membrane and blocked at 4°C overnight in 3%

bovine serum albumin (BSA) in Tris-buffer saline (TBS) (pH7.4) with 0.1% TWEEN20 (TBS-T).

Western blots probed with 1:10,000 of primary antibody of mouse anti-CdeC anti-CdeM, and anti-

SleC in 1% BSA in TTBS for 1h at room temperature. Membrane was washed 3 times for 5

minutes with TBS-T and incubated in secondary antibody 1:10,000 dilution goat anti-mouse HRP

(Sigma).

Results and discussions

Experimental design of spore coat and exosporium extraction.

To test efficiency of extractions we used three spore markers. CdeC and CdeM for efficiency of exosporium extraction since they have been identified as key morphogenic exosporium proteins. For efficient extraction of spore cortex constituents SleC was used as a marker. Initially as shown in Figure 1, spores were treated with either EBB or USD. EBB treated samples were boiled 10 minutes and then spun down, supernatant containing spore surface layer extracts was saved and remaining spore pellet was washed to remove residual extraction buffer. USD treated spores were incubated at 37°C for 90 minutes and after incubation spun down. Supernatant was saved and spore pellet was washed. Initial spores treated with USD or EBB were then treated with 10 mg/mL of lysozyme which would degrade cortex. For all treatments extracts of treated spores were run through SDS PAGE and immunoblotted for western blot analysis.

Impact of EBB and USD in spore coat and exosporium extracts.

Efficiency of extractions using EBB and USD were tested through SDS-PAGE and western blot analysis. Extracts of initial EBB treated spores (1°) indicate a substantial amount of protein is removed from the spore. A variety of proteins are extracted as seen through the SDS-PAGE protein banding. A second subsequent EBB extraction (2°) extracts residual protein likely not extracted from the initial treatment (1°). Protein bands are evident at ~10, ~15, and ~25 kDa. A third EBB extraction (3°) most of the surface proteins are extracted, no protein is seen in SDS-PAGE stained with Coomassie blue. The spores were run through SDS-PAGE to identify protein left on spore

surface after each treatment. Initial 1° spores have protein present within the SDS-PAGE indicating

not all protein was removed with the initial extraction from the spore surface consistent with the

protein that was able to be extracted following the 2° EBB extraction. Spores of 2° extraction

display protein bands indicating some protein was left on spore surface. However, spores of 3°

treatment no protein is present in SDS-PAGE indicating most if not all protein is removed with 2

total treatments. Laemmli control efficiently extracts surface proteins of however the amount is

less compared to EBB. Similarly, one treatment with Laemmli does not completely remove spore

surface proteins as seen through residual protein in the spore.

USD spore extracts as seen in Figure 2A reveal a large amount of protein is removed from the

spore surface with an initial extraction (1°). Following a subsequent extraction (2°) there is

additional protein that is able to be removed as seen in the SDS-PAGE. This indicates that not all

the protein is able to be removed from an initial treatment and substantial amount is left on the

spore surface. Similar to EBB treatment, there is little to no protein seen in spore USD extracts of

3°. Spores treated with USD (1°) have residual protein on spore surface after initial USD extraction

and a protein band is clearly seen ~10 kDA likely whole spore running through SDS PAGE.

Residual protein is seen on spore in both 2° and 3° indicating spore surface still contains protein.

Effect of EBB and USD in exosporium markers.

Western blot analysis using anti-CdeM and anti-CdeC antibodies were used as markers for efficient

extraction of exosporium proteins. CdeM is located on the outermost layer of the exosporium

whereas CdeC is found within the exosporium with interaction with spore coat. EBB extracts of

1° using anti-CdeM antibodies display immunoreactive bands at ~70 kDa, ~55 kDa, and 27 kDa

indicating formation of higher order oligomeric species as well as monomer (~27 kDa) (Figure

2B). Immunoblotting using anti-CdeM antibodies of 1° USD spore extracts display

immunoreactive bands at ~70 kDa and 55 kDa however no monomer is seen ~27kDa. Laemmli

extractions display immunoreactive bands at ~70 kDa and ~27 kDa but lacks the 55 kDa band seen

within the USD and EBB extracts. This suggest usage of traditional Laemmli may overlook certain

oligomeric bands as we've seen in USD and EBB. EBB spore extracts of 2° and 3° lack

immunoreactive bands indicating a majority of CdeM protein is removed with an initial extraction.

Likely, bands seen within SDS-PAGE in Figure 2A may be unknown spore surface constituents.

No CdeM immunoreactive bands were seen on 1° and 2° treated EBB spores. In the case of USD,

like EBB there were no immunoreactive bands seen in the treated spores as well as the subsequent

2° and 3° extractions indicating a majority of the CdeM protein is efficiently removed with a single

extraction. Laemmli is able to efficiently extract CdeM as no protein is seen in 1° Laemmli treated

spores.

Use of anti-CdeC antibodies, EBB extracts displayed immunoreactive band at ~55 kDa

corresponding to the monomeric form of CdeC (Figure 2C). Although predicted to be ~44.4 kDa

CdeC present immunoreactive bands at 55 kDa likely due to interactions with other spore surface

constituents and or formation of oligomers. No immunoreactive bands are seen in EBB extracts of

2° and 3° or in extracted spores indicating that EBB efficiently extracted most of CdeC within

spore surface. USD spore extracts 1°, western blot indicated immunoreactive bands at ~55 kDa as

well as ~130 kDa indicating formation of oligomers in USD but not EBB. Likely, EBB is able to

more efficiently break down CdeC oligomers compared to USD. USD extracts of 2° indicate

immunoreactive band at 55 kDa suggesting USD failed to remove all CdeC protein from spore

surface with an initial extraction. Supporting this, immunoreactive band ~55 kDa is seen in 1°

treated spores again indicating not all protein is removed in an initial extraction and some CdeC

remains on spore surface as seen with 2˚ USD extraction removing residual protein that was not

removed with the initial extraction. EBB however is able to efficiently remove all CdeC within

spore surface whereas USD leaves residual protein and does not efficiently dissociate oligomeric

species compared to EBB. Laemmli extraction extracts a majority of CdeC from the exosporium

as seen in initial extraction and lack of immunoreactive band in treated spore.

Effect of EBB and USD in spore coat – peptidoglycan interface marker.

As a peptidoglycan marker SleC was used to detect efficient extraction of spore surface layers as

well as cortex proteins. During sporulation SleC is processed and appears as three immunoreactive

bands, full length ~55 kDa, pro-SleC ~37 kDa, and active SleC ~27 kDa.Initial EBB extraction 1˚

displayed immunoreactive bands for all three versions of SleC at the expected ~55 kDa, ~37 kDa,

and ~27 kDa. A second extraction with EBB however had a very faint band indicating most of the

protein is extracted with one EBB treatment. Treated EBB spores 1˚ displayed a very faint

immunoreactive band suggesting there is some residual SleC not extracted with the initial

treatment that is extracted with a second extraction 2˚ as seen in Figure 2D. However, no

immunoreactive bands are seen in the treated spores for 2˚ and 3˚. Initial extraction 1˚, with USD

extracted a majority of the SleC with expected immunoreactive bands similar to EBB at ~55 kDa,

~37 kDa, and ~27 kDa. However, USD failed to completely extract SleC as immunoreactive band

was present with a second USD extraction at ~37 kDa. A weak band is seen ~37 kDa in 1˚ treated

spores consistent with USD not completely removing SleC with the initial extraction resulting in

immunoreactive bands with a 2˚ extraction. No SleC is extracted with a third extraction or seen in

2˚or 3˚ treated spores. This indicates two extractions is sufficient to completely extract SleC from

the spore. Laemmli gave consistent immunoreactive bands for all forms of SleC but failed to

completely remove SleC as seen with a strong immunoreactive band within the treated spore at

~37 kDa. EBB consistently is able to extract more of the surface layer proteins CdeC and CdeM

as well as cortex marker SleC in comparison to USD.

Impact of EBB and USD in lysozyme-triggered germination

To assess effects of USD and EBB treatment on spore coat permeability and extraction efficiency,

spores were treated with 10mg/mL lysozyme overnight at 37 ˚C for 24 hours. Untreated spores

will appear phase bright whereas spores which have degraded cortex due to permeability to

lysozyme will appear phase dark. EBB treated spores appear phase bright before treatment

however turn phase grey after an initial treatment with EBB treatment likely due to extraction of

exosporium and spore coat proteins resulting in rehydration of the core and or DPA loss due to

boiling. USD spores in contrast appear phase bright after initial treatment similar to the untreated

control suggesting EBB is a harsher treatment compared to EBB. EBB spores were treated with

Lysozyme and appeared phase dark after 24 h our incubation at 37˚C indicating that EBB affected

spore coat permeability leading got cortex degradation. In contrast, USD spores which appeared

phase bright after initial extraction were phase dark after lysozyme treatment. This indicates that

USD is able to permeabilize the spore coat. Although USD was not as efficient at extraction of

spore surface layers in contrast to EBB is it still able to permeabilize the spore coat as seen through

phase dark spores in Figure 3.

Lysozyme treatment of EBB- and USD-treated spores.

EBB and USD are efficient at extracting the spore surface layers and permeabilizing the spore

coat, increasing its susceptibility to lysozyme turning phase dark as seen in Figure 3. To identify

effect of extraction on inner spore coat layers 1° treated spores (Lane 1) were incubated with

lysozyme for 24 hrs at 37 °C and subsequently treated with EBB or USD and run through SDS-

PAGE (Lane 2). In Figure 4A, treatment of 1° spores with lysozyme followed by EBB or USD

released increased extraction of protein from the spore. In contrast Lane 3 which was treated with

lysozyme and Laemmli had a extracted more protein than Lane 2. The lysozyme is denoted by an

arrow at the predicted ~14 kDa.

Western blot analysis using anti-CdeM antibodies revealed that use of lysozyme did not lead to

any further extraction of CdeM consistent with the data from Figure 2 where most if not all CdeM

protein is removed with an initial USD or EBB extraction from the spore surface. In the case of

anti-CdeC antibody immunoreactive bands are seen at ~70 kDa and ~55 kDa in USD treated spores

with lysozyme. As seen in Figure 2, one USD extraction was not sufficient for complete removal

of CdeC from the spore surface. CdeC immunoreactive bands are seen in samples treated with

lysozyme Lane 2 and Lane 3 and not Lane 1 indicating that lysozyme allowed for further extraction

of CdeC from the spore surface layers. CdeC impacts spore coat assembly likely release when

lysozyme degrades the cortex and spore coat is degraded. In the case of SleC a marker for cortex

SleC immunoreactive bands are present in EBB treated spores Lane 1, and no immunoreactive in

Lane 2 where Lane 1 spores were treated with lysozymes for 24 hrs and underwent EBB treatment.

However, lysozyme treated spores with Laemmli were able to extract SleC. In the case of USD,

SleC is found within 1° USD treated spores Lane 1, however no immunoreactive SleC bands are

seen in either lysozyme treatment Lane 2 and Lane 3. Similarly in the case of Laemmli treated

spores alone an initial Laemmli treatment removed most if not all SleC from the spore as treatment

with lysozyme did not release addition SleC protein.

Figure legends

Figure 1. Schematic of extraction treatment. Purified R20291 C. difficile spore (5x109 spores/mL) were pelleted down and resuspended in USD or EBB. USD treated spores were incubated at 37°C for 90' min. EBB treated spores were boiled for 10' min. Spores were pelleted down for 5'min at 14,000 rpm and supernatant 1°S saved for SDS-PAGE and Western Blot Analysis. Spore pellet (1°) was washed 3x with distilled H2O and centrifuged in between washes at 14,000rpm for 5'min. 1° treated spores were subsequently treated two more times with USD or EBB as described above denoted as 2° and 3°. Supernatant and a portion of spore pellet is saved between each step for analysis. 1° spores were treated with 10mg/mL for 24 hours at 37°C to identify effects of treatments on spore coat integrity.

Figure 2. Extraction of Spore Outer layers using EBB and USD. A) 15% SDS-PAGE of purified R20291 spores treated with USD, EBB, and 2x Laemmli. Spores were treated as described in Methods and Figure 1. 1°,2°, 3° indicate consecutive extraction treatment of spores, Laemmli 1° indicates one time Laemmli extraction. Western blot analysis of treated samples using: **B)** anti-CdeM antibodies, arrows indicate immunoreactive bands at ~70 kDa, 55 kDa, and 27 kDa respectively. **C)** anti-CdeC antibodies, arrows indicate immunoreactive bands indicating extraction of CdeC with bands at ~130 kDa, 55 kDa, and 37 kDa, respectively. **D)** anti-SleC antibodies indicate immunoreactive bands at 55kDa full length SleC, 37 kDa pro-SleC, and 27kDa active SleC. Higher olimeric bands likely oligomerization of the protein.

Figure 3. Effect of extraction on spore coat permeability to lysozyme. Effects of USD and EBB treatment on spore coat integrity and lysozyme resistance. R20291 *C. difficile* spores were treated

for 90 min at 37°C incubator with USD, washed and imaged under phase contrast. EBB treated

spores were boiled for 10 min, centrifuged for 5 min at 14,000rpm and washed. Spores were treated

with 10 mg/mL lysozyme for 24 hours at 37°C. Effects on spore coat integrity and degradation of

peptidoglycan indicated by phase dark spores under phase contrast.

Figure 4. Lysozyme treatment of surface layer extracted spores. A) 15% SDS-PAGE of

purified R20291 spores treated with USD, EBB, and 2x Laemmli. Spores were treated as described

in Methods and Figure 1**.** 1° and 2° spores treated with 10mg/mL lysozyme, 1°spores were not

treated with lysozyme after initial USD/EBB treatment. 1°Laemmli treated spores with and

without lysozyme also shown. ~14kDa band on SDS corresponding to lysozyme indicated. **B)**

Western blot analysis of treated samples using **i)** anti-CdeM antibodies, show a lack of

immunoreactive bands at ~70 kDa, 55 kDa, and 27 kDa respectively. **ii)** anti-CdeC antibodies,

arrows indicate immunoreactive bands indicating extraction of CdeC with bands at 55 kDa, and

37 kDa, respectively. **iii)** anti-SleC antibodies indicate immunoreactive bands at 55kDa full length

SleC and 37 kDa pro-SleC.

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Figure 1: Schematic of spore treatment

Figure 3: SDS Western Blots

Figure 3: Phase Contrast

Figure 4: SDS Western Blots

A)

SDS-PAGE

	Ladder	EBB			USD			Laemmli	
		1	2	3	1	2	3	1	2

Ladder: 180, 130, 100, 70, 55, 40, 35, 25, 15, 10

← Lysozyme

EBB	+	+	+	-	-	-	-	-
USD	-	-	-	+	+	+	-	-
Laemmli	-	-	+	-	-	+	+	+
Lysozyme	-	+	+	-	+	+	-	+

B)

Western Blots

i) α-CdeM

	EBB			USD			Laemmli	
	1	2	3	1	2	3	1	2

70, 55, 25

ii) α-CdeC

	EBB			USD			Laemmli	
	1	2	3	1	2	3	1	2

CdeC→, CdeC→

70, 55, 40, 35, 25

iii) α-SleC

	EBB			USD			Laemmli	
	1	2	3	1	2	3	1	2

FL SleC→, pro SleC→

70, 55, 35

EBB	+	+	+	-	-	-	-	-
USD	-	-	-	+	+	+	-	-
Laemmli	-	-	+	-	-	+	+	+
Lysozyme	-	+	+	-	+	+	-	+