Outer Membrane Vesicles Secreted by *Bacteroides fragilis* Inhibit CFTR Chloride

Secretion by Human Colon Organoids

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Abstract. The goals of this study were to develop a model to study host pathogen interactions

in primary human colon organoids and to test the hypothesis that *Bacteroides fragilis* toxin

(BFT-2) secreted in outer membrane vesicles (OMVs) modulates mucosal immunity and CFTR

Cl⁻ secretion. Since Bacteroides species reside in mucus, OMVs are likely to represent a

mechanism of communication between Bacteroides and the host. Two strains of Bacteroides

were studied, Enterotoxigenic *Bacteroides fragilis* (ETBF), which produces BFT-2, and the non-

toxigenic *Bacteroides fragilis* strain NCTC 9343 (NTBF) that does not produce BFT-2. We also

utilized two additional strains of *Bacteroides fragilis:* one in which *bft*-2 was knocked out (ETBF

Δ*bft*), and one that was engineered to contain *bft-2* (NTBF+*bft)*. We report that *Bacteroides*

fragilis OMVs reduced CFTR Cl⁻ secretion but had no effect on tight junction or cell adhesion

proteins, transepithelial resistance (TER) or cytokine secretion by primary human colon

organoids. NTBF OMVs containing BFT-2 were more effective in reducing CFTR Cl⁻ secretion

than NTBF lacking BFT-2. We conclude that OMVs secreted by Bacteroides can be an

important mechanism of host pathogen interactions in the colon by reducing CFTR Cl⁻ secretion.

Keywords: *Bacteroides fragilis*, outer membrane vesicles, CFTR, transepithelial resistance,

colon organoids

Introduction. The human gastrointestinal tract is colonized by ~100 trillion bacteria, including

several members of the genus Bacteroides, which comprise approximately 30% of the

microbiota [1-5]. Although *Bacteroides fragilis* (*B. fragilis*) is one of the least abundant species,

Enterotoxigenic *B. fragilis* (ETBF) is a pathogenic bacterium that induces diarrhea, colitis and

causes tumor formation through the secretion of Bacteroides fragilis toxin (BFT-2), a

metalloprotease [3,6-8]. Many, but not all, studies on colon cell lines have shown that recombinant

BFT-2 rapidly (~1-3 hrs.) causes cell damage and disrupts cell adhesion by cleaving E-

cadherin, a cell adhesion protein, leading to an increase in epithelial paracellular permeability,

IL-8 secretion [3,9-11,12], fluid secretion driven by CFTR Cl$^-$ channels [13,14] and diarrhea [15-17].

However, in some studies on colon cell lines, E-cadherin remained unchanged after exposure to

ETBF supernatant [9]. The effect of NTBF and ETBF OMVs on IL-8 secretion and E-cadherin in

colon cell lines is dose, strain and time dependent [9,12,18]. It has been suggested that the BFT-2-

induced increase in permeability leads to paracellular movement of bacteria and bacterial

products from the lumen of the colon to the submucosal layer and blood, an effect that can train

the immune system [2]. Recombinant BFT-2 applied to the apical side of HT 29/C1, HT-29, T84,

CMT93 or MDCK cell lines decreases transepithelial resistance (TER) and cleaves E-cadherin

[9,11,15,17,19,20]. One study on the T84 colon cell line reported no effect of BFT-2 on short circuit

current but did report that BFT-2 reduced TER [20].

Many Gram-negative bacteria that reside in the mucus lining of the colon secrete outer

membrane vesicles (OMVs) [12,21-26] that stimulate inflammation and decrease ion transport,

including CFTR Cl$^-$ secretion by epithelial cell lines [21-23,27]. Therefore, it is likely that

communication between *B. fragilis* and human colon cells is also mediated by the secretion of

OMVs and other soluble factors including MUC-2, antimicrobial peptides, and anti-inflammatory

cytokines [7,28,29]. However, nothing is known about the effects of BFT-2-containing OMVs

secreted by *Bacteroides* on primary human colon organoids.

In contrast to ETBF, the Nontoxigenic *B. fragilis* (NTBF) strain does not express *bft-2* and it

promotes gut homeostasis through the production of short-chain fatty acids and polysaccharide

A, stimulates barrier-protective molecules including ZO-1 and MUC-2, as well as the secretion

of antimicrobial peptides, and anti-inflammatory cytokines [1,7,30].

A recent review questioned the relevance of cell lines as models to elucidate the physiology and

pathophysiology of the colon since cell lines are rarely normal, and contain several mutations

and chromosomal defects that enable their persistence [31]. In particular, it has been reported that

the gene expression pattern of the Caco-2 cell line, which is studied as a surrogate of the colon,

was similar to that of epithelial cells of the small intestine [32]. Since, as noted above, colon cell

lines do not produce consistent responses to recombinant BFT-2 [3,9,10], one of the most

important challenges involves the assessment of BFT-2 secreted in OMVs on primary human

colon organoids [1]. To elucidate the biological activity of BFT-2 secreted in OMVs, we obtained

colon biopsies from multiple healthy human donors to develop differentiated two-dimensional

monolayers of primary colon organoids. Primary colon organoids contain the multiple cell

populations found in native tissue, compared to cell lines, which do not fully differentiate [31].

Accordingly, the goals of this study were to develop a model to study host pathogen interactions

in primary human colon organoids, and to test the hypothesis that BFT-2 secreted in OMVs

modulates mucosal immunity, TER and CFTR Cl^- secretion, a major driver of fluid secretion in

the colon [14]. We report that OMVs had no effect on cell adhesion or tight junction proteins,

notably E-cadherin and claudin-2, TER, or sodium reabsorption, but did inhibit forskolin-

stimulated CFTR Cl^- secretion, an effect predicted to decrease fluid secretion by the colon.

NTBF OMVs containing BFT-2 had a more inhibitory effect on CFTR Cl^- secretion by primary

human colonoids than NTBF OMVs lacking BFT-2. In addition, OMVs were not pro-

inflammatory as determined by measuring the secretion of 48 cytokines. OMVs also reduced

CFTR Cl^- secretion by T84 cells, a colon cell line, but did not alter TER. These results

demonstrate that OMVs and BFT-2 secreted by *B. fragilis* have very different biological effects

on primary human colon organoids compared to colon cell lines exposed to recombinant BFT-2.

Results

Isolation and characterization of OMVs secreted by Bacteroides. To test the hypothesis

that BFT-2 in OMVs leads to a disruption in the integrity of epithelial barriers and modulates

CFTR Cl⁻ secretion, we utilized four strains of *B. fragilis*, including ETBF and ETBF in which the

bft-2 gene was deleted (ETBF Δ*bft*), NTBF, a strain that does not express *bft-2,* and NTBF in

which the *bft*-2 gene was expressed (NTBF+*bft*). Nanoparticle tracking analysis (NTA), which

measures the size of vesicles by dynamic light scattering, revealed that the size of OMVs

isolated from bacterial cultures was similar for most strains, except that ETBF OMVs were

significantly smaller than NTBF OMVs (**Figure 1**).

Figure 1: Bacteroides OMV particle size determined by NTA. Individual measurements are

shown as gray circles, with group means (black diamonds) and error bars (±SD). Treatments:

ETBF, ETBF Δ*bft*, NTBF, and NTBF+*bft*. Letters above bars indicate statistical groupings based

on one-way ANOVA with Tukey's HSD post-hoc test; groups sharing the same letter are not

significantly different (P > 0.05), while groups with different letters are significantly different (P <

0.05).

According to NTA analysis, the number of OMVs in each bacterial culture was: 1.4×10^{11}

particles/ml for ETBF, 2.8×10^{11} particles/ml for ETBF Δbft, 2.0×10^{11} particles/ml for NTBF and

2.2×10^{11} particles/ml for NTBF+*bft*. To confirm that we isolated OMVs, transmission electron

microscopy (TEM) was utilized to image OMVs (**Figure 2**). OMVs were not observed in process

control (PC, bacterial culture media not exposed to bacteria and run through the OMV isolation

process); however, OMVs were observed in media isolated from ETBF, NTBF, ETBF Δ*bft* and

NTBF+*bft* (**Figure 2**). The mean diameter (±SEM) of OMVs as measured by TEM were similar

in all groups (P>0.77): ETBF [(59.0 nm ± 2.0 nm, (n=31)], ETBF Δ*bft* [(63.2 ± 3.6 nm (n=21)],

NTBF [(60.2 ± 4.0 nm, (n=14)] and NTBF+*bft* [(58.0 ± 5.0 nm (n=11)]. Most Bacteroides OMVs

had typical morphology, but we also observed some collapsed vesicular forms and the presence

of some irregularly shaped vesicles, similar to images published recently for Bacteroides OMVs

secreted by ETBF and NTBF [33]. As expected, the size of OMVs was assay dependent: dynamic

light scattering tends to overestimate size, whereas TEM tends to underestimate size [34,35].

Figure 2. Representative transmission electron microscopy images of negatively stained

Bacteroides OMVs. (A) Process control. (B) ETBF. (C) ETBF Δ*bft*. D. NTBF. E. NTBF+*bft*.

Images taken by an observed blinded to the origin of the OMVs. Scale bars 100 nm.

Analysis of the proteome of OMVs was conducted to confirm that BFT-2 was present in ETBF

and NTBF+*bft* and absent in ETBF Δ*bft* and NTBF. Results of the proteomics analysis of OMVs

are presented in **Figure 3 and** Supplemental Tables 1-4. We detected 568 proteins in NTBF

OMVs and 437 proteins in ETBF OMVs, with 247 proteins being found in both strains. The

amount of BFT-2 was similar in ETBF OMVs and NTBF+*bft* OMVs as determined by intensity-

based quantification ($\log_2$ 12.43 for NTBF+*bft* and $\log_2$ 12.90 for ETBF). BFT-2 could not be

detected by proteomic analysis in NTBF and ETBF Δ*bft* OMVs.

Figure 3. ETBF and NTBF OMV proteomes. Differential abundance analysis reveals significant differences between the proteome of ETBF and NTBF OMVs. The list of all identified proteins by proteomics analysis are presented in the Supplemental Tables. Supplemental Table 1 includes proteins detected exclusively in NTBF OMVs. Supplemental Table 2 includes proteins detected exclusively in ETBF OMVs. Supplemental Table 3 includes proteins detected in both ETBF and NTBF OMVs. Finally, Supplemental Table 4 includes differential expression of

proteins in ETBF and NTBF OMVs, the results of which are depicted in this Figure. In

accordance with previous research, our results show several TonB dependent receptors are

more abundant in OMVs secreted by ETBF compared to NTBF strains [36].

Characterization of primary human colon organoids: Development of a model to study

host pathogen interactions in the colon. Biopsies of human proximal colon were obtained

during clinical colonoscopy after obtaining informed consent from healthy volunteers at

Dartmouth Health. As described in methods, crypt cells were dispersed and grown in Matrigel

until 3D colon organoids were formed. 3D colon organoids were then seeded on Transwell

filters and grown for one week to develop polarized and differentiated colon organoids. We used

an imaging approach to characterize the cellular composition of colon organoids *in vitro* as

monolayers. **Figures 4-5** reveal that the colon organoids contain cell types found in the proximal

colon including mucin (MUC2) producing cells, CFTR expressing cells, SLC26A3 expressing

cells, and enterochromaffin cells [14,37]. These observations indicate that the human primary colon

organoids are a well differentiated epithelium, have similar cell types to the colon *in vivo* and are

a new model to study host pathogen interactions [14,37].

Figure 4. Representative images of human colon organoids. (A) Cross section of a

confluent monolayer of colon organoids grown on Transwell filters (10 μm thick: H&E-stained

cells). (B) differential contrast (DIC) image of a 6 μm thick frozen cross-section of differentiated

colon organoids on Transwell filters. (C) Image of the same colon organoid as in B but

immunofluorescent image of CFTR. (D) DAPI nuclear labeling of the same colon organoid

monolayer as in B and C. (E) Composite of B-D showing apical membrane labeling of CFTR.

Scale bar in panel E is 50 μm.

Figure 5. Representative *en face* images of human colon organoids on Transwell filters.

(a) DAPI labeling of nuclei. (b) Chromogranin A immunofluorescence of enterochromaffin cells.

(c) Immunofluorescence of MUC2. (d) Composite of A-C revealing that MUC2 and

Chromogranin A are expressed in different cell types. (e) Immunofluorescence of SLC26A3

(DRA, a Cl^-/HCO_3^- exchanger). (f) DAPI labeling of nuclei. (g) Composite of e and f. Scale bar in

panel g is 50 μm.

OMVs inhibit CFTR Cl^- secretion by primary human colon organoids. Previous studies

have shown that recombinant BFT-2 stimulates CFTR Cl^- secretion, when added to the

basolateral side of T84 cells [20]. To examine the effect of OMVs secreted by *B. fragilis* on CFTR

Cl^- secretion by primary human colon organoids, OMVs were isolated and applied to the apical

side (the side of the colon exposed to Bacteroides *in vivo*) of colon organoids for 1 hr. and then

forskolin-stimulated CFTR Cl^- secretion was measured in Ussing chambers immediately after

the 1 hr. exposure, or 24 hrs. after washing the OMVs from the organoids. Forskolin increases

the activity of protein kinase A, which phosphorylates and activates CFTR Cl⁻ secretion [38].

Figure 6 demonstrates that OMVs secreted by ETBF, ETBF Δ*bft* and NTBF+*bft* reduced

forskolin stimulated CFTR Cl⁻ secretion after 1 hr. of exposure compared to process control

(PC). Notably, OMVs secreted by ETBF reduced Cl⁻ secretion significantly more than OMVs

secreted by NTBF that does not express BFT-2. Expression of *bft* in NTBF (NTBF+*bft*) reduced

CFTR Cl⁻ secretion compared to NTBF OMVs, which do not express BFT-2.

Twenty-four hrs. after a 1 hr. exposure to OMVs, CFTR Cl⁻ secretion was not significantly

reduced by ETBF, ETBF Δ*bft*, or NTBF when compared to PC (**Figure 6**). However, 24 hrs.

after exposure to NTBF+*bft* OMVs, CFTR Cl⁻ secretion was less that CFTR Cl⁻ secretion

compared to NTBF OMVs (**Figure 6**).

OMVs had no effect on the amiloride-sensitive short circuit Na⁺ currents (Isc, μA/cm^2), either

after a 1 hr. exposure to OMVs or after 24 hrs. following the 1 hr. exposure to OMVs (**Figure 7**).

Figure 6. CFTR Cl⁻ secretion by human primary colon organoids measured as short

circuit current (Isc, μA/cm^2). Left: 1 hr. after exposure to PC or OMVs (2×10^{10}/ml OMVs, a

concentration of OMVs similar to that measured in conditioned media and in biological fluids

[21,39]). Right: 24 hrs. after a 1 hr. exposure to PC or OMVs (2×10^{10}/ml). Individual donor

responses are shown as dashed colored lines, with each color representing a different donor (n

= 3 donors; Donor 108A is female, Donors 100A and 107A are male). Solid black lines connect

the mean current values across treatments. Data expressed as mean ± one standard deviation.

Many significant differences from PC were observed at 1 hr. (ETBF, ***P = 0.00014; ETBF Δ*bft*,

***P = 0.00014; and NTBF+*bft*, **P = 0.0030). Expression of *bft* in NTBF (NTBF+*bft)* OMVs

reduced CFTR Cl⁻ secretion at 1 hr. compared to NTBF OMVs, as indicated by the horizontal

bar (**P = 0.01200). At 24 hrs. only NTBF+*bft* OMVs reduced CFTR Cl⁻ secretion compared to

PC (P=0.008). The horizontal bar indictes that NTBF+*bft* OMVs significantly reduced CFTR Cl⁻

secretion (**P = 0.004) compared to NTBF OMVs. Statistical analysis was performed using

mixed-effects linear models with donor as a random effect for each time point separately.

Experiments done with three donors of colon cells, three technical replicates per donor and

three different preparations of OMVs. Technical replicates were averaged as a single data point.

Figure 7. OMVs did not modify amiloride-sensitive short circuit Na⁺ currents. Amiloride-sensitive Na⁺ currents in response to OMVs (2×10¹⁰/ml OMVs) compared to PC. The difference in amiloride-sensitive currents (ΔIsc) between PC and OMVs exposed colon organoids is plotted. Colon organoids were exposed to OMVs for 1 hr. or 24 hrs. Each measurement of amiloride-sensitive current for each donor was adjusted by subtracting the Process Control (PC) current from the same Ussing chamber session and the same donor. Data from each donor (100A = red, 107A = green, 108A = blue) is depicted with dashed lines. The Isc OMVs vs. PC) for all donors is shown with a solid black line with ± one standard deviation. The horizontal line at zero represents no change from PC. Statistical analysis was performed using mixed-effects linear models with donor as a random effect for each time point separately. Experiments done with three donors of colon cells, three technical replicates per donor and three different preparations of OMVs. Technical replicates were averaged as a single data point.

Experiments were also conducted to determine if OMVs had an effect on TER, since previous studies have shown that recombinant BFT-2 reduced TER in several colon cell lines [9,15,17,19,20].

However, none of the OMVs had a significant effect on TER (**Figure 8**) compared to process

control (PC) at 1 hr. and 24 hrs.

Figure 8. Transepithelial resistance (TER) of primary human colon organoids. TER was

measured in colon organoids in which CFTR Cl⁻ secretion was measured 1 hr. (left figure) or 24

hrs. (right figure) after exposure to PC or OMVs (2×10¹⁰/ml OMVs) secreted by ETBF, ETBF

Δ*bft*, NTBF and NTBF+*bft*. Individual donor responses are shown as dashed colored lines, with

each color representing a different donor. Solid black lines connect the mean TER values

across treatments. Data expressed as mean ± one standard deviation. Statistical analysis was

performed using mixed-effects linear models with donor as a random effect for each time point

separately. Experiments done with three donors of colon cells, three technical replicates per

donor and three different preparations of OMVs. Technical replicates were averaged as a single

data point. No significant differences compared to PC (1hr. and 24 hr. exposure) were observed

($P > 0.05$).

Bacteroides reduces CFTR Cl⁻ secretion by T84 cells but has no effect on TER. We also

conducted studies on T84 cells, a colon cell line that has been used to study the effects of

recombinant BFT-2 [40-42]. Previous studies have shown that recombinant BFT-2 reduces the TER

in several colon cell lines [9,15,17,19,20]. To determine if *B. fragilis* OMVs also reduced TER and

influenced CFTR Cl⁻ secretion, we conducted Ussing experiments on monolayers of T84 cells

under conditions identical to the previously described exposure of primary human colon

organoids. Compared to PC, CFTR Cl⁻ secretion was reduced by all OMVs (**Figure 9a**). ETBF

OMVs were not more effective than NTBF OMVs in reducing CFTR Cl⁻ secretion. Moreover,

deletion of *bft-2* in ETBF did not significantly reduce the effect of ETBF OMVs on CFTR Cl⁻

secretion. OMVs also had no effect on TER (**Figure 9b**). Thus, OMVs reduced CFTR Cl⁻

secretion by both primary human colon organoids and T84 cells, without altering TER.

Figure 9. CFTR Cl⁻ secretion and TER by T84 cells exposed to OMVs. The protocol was

identical to that performed on primary human colon organoids. T84 monolayers were exposed

to OMVs ($2×10^{10}$/ml) for 1 hr. and CFTR Cl⁻ secretion (short-circuit current: Isc) was measured.

Data represent individual experiments (dashed lines) grouped by experimental date (colors) with

means ± one standard deviation. (a) CFTR Cl⁻ secretion. Each line connects experiments

conducted on the same day on the same pass of T84 cells. Compared to PC, CFTR Cl⁻

secretion was significantly reduced by ETBF (**P = 0.006), ETBF Δ*bft* (**P = 0.005), and

NTBF (*P = 0.032) OMVs. (b) TER in the same set of T84 cells used for measurements of

CFTR Cl⁻ secretion. No significant differences in TER compared to PC were observed after 1 hr.

of exposure to OMVs. Statistical analysis was performed using mixed-effects linear models with

donor as a random effect for each time point separately. Experiments done with three different

preparations of OMVs. Technical replicates were averaged as a single data point.

OMVs do not alter tight junction or cell adhesion protein abundance. Previous studies

have also shown that recombinant BFT-2 rapidly (<1 hr.) reduced the expression of the cell

adhesion protein E-cadherin in cell lines [9,15,17,19,20]. However, we observed no effect of

Bacteroides OMVs on the abundance of E-cadherin or on the tight junction protein claudin-2 in

human colon organoids as determined by immunofluorescence microscopy (**Figure 10**).

Figure 10. Representative composite images of human colon organoids. Organoids were

grown on Transwell filters and exposed to OMVs (2×10¹⁰/ml) for 1 hr. Claudin-2 (magenta) and

E-cadherin (green) were identified by immunofluorescence. DAPI (blue) identified nuclei. Each

image is a composite of colon organoids labeled for claudin-2, E-cadherin and nuclei (DAPI).

Confocal images were analyzed by an investigator who was blinded to the treatment. There was

no difference in claudin-2 or E-cadherin abundance using confocal microscopy and the Fiji

open-source image processing package to quantify fluorescence intensity (see **Figure 11** for

data).

Studies were also conducted to measure E-cadherin and claudin-2 fluorescence imaged by

confocal microscopy. None of the OMVs had a significant effect on E-cadherin or claudin-2

fluorescence (**Figure 11**) confirming the immunofluorescent images in Figure 10.

Figure 11. Measurements of E-cadherin (A) and claudin-2 (B) epifluorescence. Data were

generated by a co-author blinded to the experimental treatment. Five random images were

chosen for each treatment. There was no significant effect of OMVs $(2\times10^{10}$/ml) on E-cadherin

or claudin-2 abundance. AU, arbitrary fluorescent units. Data expressed as mean ± one

standard deviation.

Cytokine secretion by human primary colon organoids.

Previous studies on cell lines and human colon tissue reported that *B. fragilis* and recombinant

BFT-2 stimulates IL-8 secretion [9,15,17-20]. To examine the effect of OMVs on the innate immune

response, colon organoids were grown on Snapwell filters and exposed to PC or OMVs in the

apical media for 1 hr. or 24 hrs. after OMVs were removed by washing. **Figure 12** demonstrates

that OMVs had no significant effect on IL-8 secretion compared to PC. Moreover, there was no

significant effect of any OMVs compared to PC on the secretion of 48 cytokines measured using

the MILLIPLEX MAP Human Cytokine/Chemokine 48-Plex cytokine assay, although there were

some trends in the response. For example, compared to PC, 24 hrs. after exposure to ETBF

OMVs, and NTBF+*bf*t OMVs IL-8 increased compard to PC; however the changes were not

statistically significant.

Figure 12. Cytokine secretion following 1 hr. and 24 hrs. after exposure to OMVs secreted

by *Bacteroides fragilis* or to PC. There were no significant effects of OMVs (2×10^{10}/ml) versus

PC. Values represent means across all donors (n = 3 donors) for each treatment condition.

Color scale indicates standard deviations from the time point mean: red represents above-

average expression, blue represents below-average expression, and white represents average

expression for each cytokine at each time point. Only cytokines with complete data (values

above detection) across all conditions are shown. Data were log2-transformed prior to z-score

calculation.

Discussion

There were two primary goals of this study: (1) To develop a well-differentiated primary culture

of human colon organoids and (2) to elucidate the effect of endogenous BFT-2 in OMVs

secreted by *B. fragilis* on human primary colon organoids. We sucessfully developed a model of

colon organoids from three donors that displayed a differentiated epithelium containing cell

types found in the proximal colon including mucin (MUC2) producing cells, CFTR expressing

cells, SLC26A3 (DRA, a Cl^-/HCO^-_3 exchanger that plays a key role in fluid absorption and

enterocyte acid/base balance) expressing cells, and enterochromaffin cells that underlie

epithelial cells and secrete serotonin, which stimulates GI motility and fluid and electrolyte

secretion [14,37]. These observations indicate that the human primary colon organoids are a

representative model to study host pathogen interactions in the human colon [14,37]. Studies were

also conducted to examine the effect of OMVs on CFTR Cl^- secretion, amiloride-sensitive

sodium absorption, TER, and cytokine secretion. We found that OMVs secreted by most strains

of Bacteroides significantly reduced CFTR Cl^- secretion by colon organoids compared to PC

after 1 hr. exposure. Deletion of the *bft*-2 gene from ETBF reduced the inhibitory effect of ETBF

OMVs on CFTR Cl^- secretion. Moreover, NTBF+*bft* OMVs were significantly more inhibitory to

CFTR Cl^- secretion compared to OMVs secreted by NTBF. Both observations indicate that BFT-

2 inhibits CFTR Cl^- section. Twenty-four hrs. after exposure to OMVs, only NTBF+*bft* OMVs

reduced CFTR Cl^- secretion compared to NTBF OMVs and PC. Thus, BFT-2 in OMVs was, at

least in part, responsible for the decrease in CFTR Cl⁻ secretion by primary human colon

organoids. Since OMVs secreted by strains of *B. fragilis* that lacked *bft*-2 (BFT Δ*bft* and NTBF)

also reduced CFTR Cl⁻ secretion, there must be other factors in OMVs including, but not limited

to, proteins or miRNAs that inhibit forskolin-stimulated CFTR Cl⁻ secretion. By contrast to results

from human colon organoids all OMVs secreted by *B. fragilis* inhibited CFTR Cl⁻ secretion by

T84 cells to the same extent. The reason for this difference between the effect of OMVs on T84

cells and human colon organoids is unknown, but it may be related to several factors, including,

but not limited to, possible differences in the time dependent effects of BFT-2 as well as

differential expression of gene and signaling pathways between T84 tumor cells and primary

human colon organoids. Additional studies are required to identify these factors.

The observation that OMVs inhibited CFTR Cl⁻ secretion by primary human organoids is

interesting to note since infection with ETBF causes secretory diarrhea, which one might expect

to be caused by increased CFTR Cl⁻ secretion, a primary driver of fluid secretion by the

colon[13,14]. However, there is differential expression and regulation of ion transporter proteins

along the length of the colon, as well as differential responses to drugs, hormones, and immune

factors along the length of the colon [13,14]. Accordingly, our results are applicable to the proximal

human colon, and other segments of the colon may react differently to OMVs and BFT-2. It is

the composite effect of ETBF and BFT-2 on the gastrointestinal tract that determines the

response *in vivo* to ETBF.

Our studies also did not observe a significant effect of OMVs on cytokine secretion by human

colon organoids. Most notably, IL-8 secretion was not significantly increased by OMVs(**Figure**

12), in contrast to studies using recombinant BFT-2 in cells lines reporting that recombinant

BFT-2 increased IL-8 secretion [9,15,17-20]. The reason for this difference between the effect of

OMVs on colon cell lines and human colon organoids is unknown, but it may be related to

several factors, including, but not limited to, differences in the amount of recombinant versus

native BFT-2 in OMVs, as well as differences in gene signaling pathways between colon cell

lines and human primary colon organoids. Studies have also shown that the effect of

recombinant BFT-2 and OMVs on colon cell lines is dose dependent, and it is likely that the

concentration of BFT-2 in OMVs is less that the doses of recombinant BFT-2 utilized in studies

on cell lines [11,18]. We conclude that biologically relevant levels of OMVs[21,39] secreted by *B.*

fragilis containing BFT-2 inhibit CFTR Cl⁻ secretion, but have no effect on amiloride-sensitive

sodium reabsorption, claudin-2, E-cadherin or TER, on primary human colonoids.

Fragipain was not detected in any of the OMV samples (Supplemental Table 4). Fragipain is

known to cleave BFT-2 into its active form [43], however, studies suggest that this protein is not

required for cleavage of BFT-2 in antibiotic treated mice [44]. More specifically, fragipain is

unnecessary when recombinant BFT-2 is mixed with the colon mucus of pathogen free or germ-

free mice, suggesting the mucosal activation process may be more prevalent in gastrointestinal

conditions containing increased mucus, such as CF [44]. Notably, primary human colon organoids

also secrete mucin. PBS is also able to facilitate the cleavage of BFT-2 into its active form, and

further suggests that fragipain is not required for BFT-2 activation [44].

In accordance with previous research, our proteomics data revealed that several TonB

dependent receptors are more abundant in OMVs secreted by ETBF compared to NTBF strains

(Supplemental Table 4) [45]. TonB dependent receptors are one part of outer membrane

structures that are critical for the influx of nutrients via SusC and SusD outer membrane

proteins, which were elevated in ETBF OMVs compared to NTBF OMVs [36]. An increase in

abundance of these proteins suggests an enhanced need for ETBF to increase nutrient

acquisition, thereby improving its ability to compete for similar niches that NTBF does not

struggle to fill. Regardless, these results pose interesting questions for mechanistic follow up

experiments in the future.

Importantly, although several studies have shown that recombinant BFT-2 degrades E cadherin

and claudin-2 in cell lines[9,15,17,19,20] , none of the OMVs tested herein had any effect on the

abundance of the tight junction protein claudin-2, the cell adhesion protein E-cadherin, or on the

TER. Claudin-2 is an important component of the tight junction, and reductions in claudin-2

abundance decrease TER[46,47]. Several possible explanations may account for the difference in

the TER, E-cadherin, claudin-2, and cytokine response of colon tumor cell lines and primary

human colon organoids. First, as noted above, the effects of BFT-2 are dose dependent [18].

OMVs secreted by ETBF and NTBF+*bft* contain endogenous levels of BFT-2. It is possible that

the concentration of recombinant BFT-2 in previous studies was much higher than the amount

of native BFT-2 in OMVs, although a comparison of concentrations is not possible given

available data. Second, the duration of exposure to OMVs may be a factor. The literature

reports that the effects of recombinant BFT-2 on E-cadherin and claudin-2 in cell lines required

anywhere from 1 hr. to several hrs. of exposure[3,9,10]. Our study was limited to a one-hr.

exposure, which has been long enough to produce significant effects in many previous studies

with recombinant BFT-2 and colon cell lines [3,9,10].

There are a few other reports describing the development of monolayers of epithelial cells

derived from healthy colon organoids [31,32]. Sen et al. [32] reported that the gene expression profile

of human colon organoids was significantly different from the Caco-2 cell line, an immortalized

cell line of human colorectal adenocarcinoma cells, which had a gene expression profile similar

to the small intestine. They reported that the effect of *Bifidobacterium longum* on gene

expression was not conserved between Caco-2 and colon organoids. Taglieri et al [31] reviewed

the literature on human colon organoids and noted that air liquid interface culture increased

differentiation of colon organoids. However, none of the other studies on colon organoids

exposed them to OMVs or reported CFTR Cl⁻ secretion.

OMVs secreted by most strains of *B. fragilis* in the present study inhibited forskolin-stimulated

CFTR Cl⁻ secretion by the proximal colon and T84 cells, which would reduce salt and fluid

secretion by the colon. However, studies in animals report that recombinant BFT-2 induces

secretory diarrhea [3,6-8]. One possible reason for the different effects of recombinant BFT-2 and

OMVs containing BFT-2 is that the effect of BFT-2 is dose dependent, and that recombinant

BFT-2 levels in published studies may be higher than the abundance of BFT-2 in OMVs. As

demonstrated in Figure 3 and the Supplemental Tables, OMVs contain many proteins in

addition to BFT-2, which may influence CFTR Cl^- secretion by primary human colon organoids.

Finally, it is important to note that there are other pathways that mediate transepithelial salt and

water transport by the proximal colon, including electroneutral NaCl and fluid reabsorption

mediated by the parallel operation of Na^+/H^+ and Cl^-/HCO_3^- (SLC26A3) exchangers. Since

Ussing chamber studies only measure electrogenic transport, it is possible that recombinant

BFT-2 (and endogenous BFT-2) may inhibit SLC26A3 and/or the Na^+/H^+ exchanger *in vivo* and

thereby reduce electroneutral NaCl and fluid reabsorption by the colon. Such an effect may

contribute to the ETBF-induced secretory diarrhea, despite the OMV reduction in electrogenic

CFTR Cl^- secretion reported herein. Thus, extensive additional studies, beyond the goals of the

present study, are required to determine if BFT-2 and OMVs inhibit Na^+/H^+ and Cl^-/HCO_3^-

electrically neutral NaCl and fluid reabsorption in primary human colon organoids. Such studies

would require use of radioactive isotopes and chemical inhibitors of electrically neutral

transporters and/or antisense experiments to selectively reduce electroneutral transporter

expression.

We also did not observe an effect of any OMVs on the TER or epithelial tight junction or cell

adhesion structure. In many, but not all, previous studies in colon cell lines, recombinant BFT-2

quickly (< 1 hr.) reduced the abundance of E-cadherin and the TER [9,15,17,19,20]. Although the

reason for this difference is unknown, one possible explanation for the different effects of

recombinant BFT-2 and OMVs is that, as noted above, the effect of recombinant BFT-2 is dose

and time dependent, and that recombinant BFT-2 levels in published studies may be higher

than the abundance of endogenous BFT-2 in in OMVs.

In contrast to studies using cells lines and recombinant BFT-2 [10,48-50], we did not observe a

significant increase in IL-8 secretion in primary human colon organoids exposed to OMVs. The

lack of effect of BFT-2 and OMVs on IL-8 secretion in our study are consistent with the

observation that OMVs did not reduce E-cadherin, an effect reported to be required for BFT-2 to

stimulate IL-8 secretion [10,48-50]. OMVs did not affect the secretion of any of the 48 cytokines as

determined by the MILLIPLEX MAP Human Cytokine/Chemokine 48-Plex cytokine assay. The

reason for this difference is also unknown and requires additional studies.

There are a few limitations of our study. First, since there are many differences in the content of

OMVs secreted by ETBF and NTBF such as DNA, RNA, miRNA, proteins (this study) and other

virulence factors [24,51], in addition to BFT-2, it is possible that some of these factors may act

alone of in concert with BFT-2 to inhibit CFTR Cl⁻ secretion. Second, although we did not

complement the *bft-2* gene in ETBF Δ*bft*, due in part to the technical challenge of

complementation in our clinical strain, we have complementary lines of evidence to support our

conclusions that BFT-2 inhibits CFTR Cl⁻ secretion in colon organoids. First, we utilized two

strains o*f B. fragilis* that lack BFT-2 (ETBF Δ*bft* and NTBF) and two that secrete BFT-2 (ETBF

and NTBF+*bft*). Our results on primary human colon organoids are similar in ETBF Δ*bft* and

NTBF, as are the results using ETBF and NTBF+*bft*.

Our study has numerous strengths. First, our experiments were conducted on well differented

human colon organoids from three healthy donors, and a cell line (T84). OMVs inhibited CFTR

Cl⁻ secretion by colon organoids and T84 cells, but had no effect on TER, or the expression of

claudin-2 and E-cadherin. Second since *B. fragilis* resides in mucus overlying colon organoids,

OMVs containing BFT-2 are biologically relevant. Notably, the concentration of OMVs used in

the present study was similar to levels observed in conditioned cell culture media and biological

fluids[21,39], and the same number of OMVs was used in all experimental groups [21,39].

In conclusion, we have examined the effects of OMVs on well differentiated primary cultures of

human colon organoids secreted by *B. fragilis* containing *bft-2 (*ETBF and NTBF+*bft)*, a strain of

B. fragilis in which *bft-2* was knocked out (ETBF Δ*bft*), and an NTBF strain that does not

express *bft-2.* In all cases, BFT-2 containing OMVs secreted by *B. fragilis* reduced forskolin-

stimulated CFTR Cl⁻ secretion by colon organoids more than the same strain lacking *bft-2* but

had no effect on tight junction structure (E-cadherin and claudin-2), TER, amiloride-sensitive

sodium absorption, or IL-8 secretion. Similar experiments on the T84 cell line also revealed that

OMVs secreted by *B. fragilis* reduced CFTR Cl⁻ secretion and had no effect on TER. We

conclude that *B. fragilis* OMVs containing biologically significant levels of BFT-2 inhibit forskolin

stimulated CFTR Cl⁻ secretion by primary human colon organoids, without effecting TER,

amiloride-sensitive Na⁺ currents and cytokine secretion. Additional studies, beyond the scope of

the present study, are required to elucidate the cellular mechanisms whereby OMVs rapidly

inhibit CFTR Cl⁻ secretion. Although the results in this study contradict many studies using cell

lines and recombinant BFT-2, we anticipate that our results will stimulate a discussion of the

effects on BFT-2 on the colon.

METHODS

Bacteroides culture and genetics. *B. fragilis* NCTC 9343 was chosen as the non-toxigenic *B.*

fragilis (NTBF) strain, since it is used widely [52-54]. *B. fragilis* 86-5443-2-2 was chosen as the

enterotoxigenic *B. fragilis* (ETBF) strain for similar reasons, and because it expresses BFT-2,

which is the most virulent of the three BFT isotypes [52-54]. *B. fragilis* 86-5443-2-2 with a *bft*

deletion (ETBF Δ*bft*), provided by Dr. Cynthis Sears, was confirmed by PCR [55]. To construct the

NTBF+*bft* strain, we first PCR amplified *bft-2* from *B. fragilis* 86-5443-2-2 and cloned it via

Gibson Assembly into the SalI and NcoI sites of the *Bacteroides* expression plasmid pNBU2-

P1T-DP-A21 [56]. This plasmid integrates in single copy into the chromosomal att sites of

Bacteroides fragilis. The sequence of the resulting plasmid was confirmed by Sanger

sequencing before transformation into *E. coli* S17-1 cells and conjugation via overnight aerobic

solid agar co-culture mating into *B. fragilis* 9343 (NTBF). Following incubation, matings were

resuspended and plated on BHIS agar containing gentamycin and erythromycin. Att

chromosomal integration site was confirmed by PCR [57]. Bacteroides strains were grown on

supplemented brain heart infusion medium (BHIS) agar plates or liquid broth cultures

supplemented with hemin and 1µg/ml of vitamin K3 (Acros Organics/Fisher Scientific, Geel,

Belgium) under anaerobic conditions (10% CO_2, 10% H_2, 80% N_2) in a Whitley A55 anaerobic

chamber (Don Whitley Scientific, Victoria Works, UK) at 37°C. All BHIS agar and liquid media

contained 60 µg/ml of gentamycin sulfate. For OMV purification, *B. fragilis* strains were streaked

on BHIS agar plates and were grown in anaerobic conditions for 48 hrs. Single colonies were

used to inoculate 5mL of BHIS liquid medium cultures, which were grown anaerobically for 24

hrs. The cultures were then expanded into 200mL BHIS cultures using a 1:100 dilution. The final

cultures were grown for 24 hrs. prior to being utilized to isolate OMVs.

Outer membrane vesicle (OMV) isolation. OMVs were isolated as described by us and others

previously [23,58]. Briefly, all overnight culture supernatants were centrifuged for 1 hr. at 2800 g

and 4°C to pellet bacteria. The supernatant was filtered twice through 0.45 µm PVDF

membrane filters (Millipore, Billerica, MA, USA) to remove any contaminating bacteria and

concentrated with 30K Amicon filters (Millipore, Billerica, MA, USA) at 2800 g and 4°C to obtain

~200µL of concentrate. The concentrate was resuspended in OMV buffer (20 mM HEPES, 500

mM NaCl, pH 7.4) and subjected to ultracentrifugation using a TH-660 Swinging Bucket rotor

(Thermo Fisher) for 2 hrs. at 200,000 g and 4°C to pellet OMVs. OMV pellets were then re-

suspended in 60% OptiPrep Density Gradient Medium (Sigma-Aldrich, Cat. # D1556) and

layered with 35%, 30% and 20% OptiPrep diluted in OMV buffer. OMVs in OptiPrep were

centrifuged for 18 hrs. at 100,000 g and 4°C. 500µl fractions were taken from the top of the

gradient, with OMVs residing in fractions 5-8, as determined by proteomic analysis of OMVs.

Electron microscopy of OMVs. OMVs were visualized with negative staining transmission

electron microscopy as described previously [22]. The microscopist was blinded to the

experimental treatment to minimize observer bias. Briefly, an aliquot of each sample was

applied onto a freshly glow-discharged 200-mesh nickel grid and allowed to sit for two minutes.

After incubation, excess fluid was gently removed with filter paper, and the grids were

sequentially rinsed with Millipore-filtered distilled water. While the grids remained slightly damp,

they were placed onto a drop of Nano-W tungsten stain (Ted Pella, Inc.). Excess stain was

absorbed with filter paper, and the staining step was repeated on a second drop of Nano-W,

which was allowed to sit for one minute before being wicked away and left to dry. Imaging of the

stained grids was conducted at 80 kV on a JEOL 1400 transmission electron microscope (JEOL

USA Inc.), and images were captured using an AMT XR11 digital camera (AMT).

Proteomics characterization of OMVs. To confirm that BFT-2 was secreted by ETBF and

NTBF+*bft* in OMVs, but not by ETBF Δ*bft* and NTBF OMVs, OMVs were isolated as described

above and pelleted and lysed in buffer containing 0.5%SDS, 50mM ammonium bicarbonate,

50mM NaCl, and Halt Protease Inhibitor. Mass spectrometry analysis was performed by the

Mass Spectrometry Technology Access Center (MTAC) at Washington University School of

Medicine using previously published protocols [59]. Samples were purified by trichloroacetic acid

precipitation. After pelleting and washing with ice-cold acetone, the resulting protein pellet was

resuspended in 8M urea and 0.4M ammonium bicarbonate, reduced with 4mM dithiothreitol,

and alkylated with 18mM iodoacetamide. The solution was then diluted to <2M urea, and 1μg of

trypsin was added for overnight digestion at 37°C. The resulting peptides were desalted using

C18 solid-phase extraction spin columns, and eluates were dried under vacuum using a

SpeedVac concentrator. Peptides were analyzed by LC–MS/MS using a nanoElute2 coupled to

a timsTOF Pro2 mass spectrometer (Bruker Daltonics). Samples were separated on a C18

column (150μm × 25cm, 1.5μm; Bruker). The mobile phases consisted of 0.1% formic acid (FA)

in water (mobile phase A) and 0.1%FA in acetonitrile (mobile phase B). Peptides were

separated using a 54-min. linear gradient from 3% to 35% mobile phase B, with a total run time

of 60 mins. at a constant flow rate of 600nL/min. The timsTOF Pro2 was operated in dia-PASEF

mode. Spectra were acquired across an m/z range of 300-1200, using 28 mass windows per

cycle. The inverse reduced ion mobility ($1/K_0$) range was set to 0.65–1.35 V·s/cm². The trapped

ion mobility spectrometry (TIMS) analyzer operated at a 100% duty cycle with a ramp time of 75

ms., yielding an overall cycle time of approximately 1.20 seconds. Collision energy was linearly

ramped from 59 eV at $1/K_0$ = 1.6 to 20 eV at $1/K_0$ = 0.6. dia-PASEF data files were analyzed

using directDIA™ workflow embedded in Spectronaut 18 (Biognosys AG) against the

Bacteroides fragilis database. Search parameters included trypsin digestion with cleavage after

K or R, allowance for up to two missed cleavages, carbamidomethylation of cysteine (static

modification), and variable modifications including oxidized methionine and Protein N-terminal

acetylation. Peptide spectrum matches (PSMs), peptides, and protein groups were identified

using a false discovery rate (FDR) threshold of 0.01. For quantification, all fragment ions

passing the FDR threshold were used, and protein abundances were calculated as the area

under the curve of extracted ion chromatograms within the defined peak boundaries for each

targeted ion. The mass spectrometry proteomics data have been deposited to the

ProteomeXchange Consortium via the PRIDE partner repository[60] with the dataset identifier

PXD073462.

Colon organoid cell culture. Biopsies from the proximal colon obtained during colonoscopies

of three healthy de-identified donors were obtained after informed consent at Dartmouth-Health.

Biopsies in PBS (-/-) with EDTA were placed on a shaking rocker in +4°C for 1 hr. to free the

crypts from the biopsies. Then the biopsies were vigorously pipetted up and down several times

to release crypts. The supernatant containing crypts was collected, and additional PBS was

added to the biopsies followed by vigorous pipetting. Crypts in the supernatant were counted,

pelleted, and resuspended in 55% Matrigel (Corning Life Sciences) and 45% isolation medium.

Cells were added to 24-well plates and then Matrigel solidified cell isolation medium (IntestiCult

complete medium with primocin [StemCell Technologies] supplemented with 50ug/mL

gentamycin, 50ug/mL vancomycin and 10uM of RhoKi) was added to each well. Cells were

maintained in culture until spherical organoids were observed. To develop polarized monolayers

of organoids, 1.5×10^6 cells (as spherical organoids) were collected from the Matrigel and

seeded on Transwell permeable filters (#3450: Corning Life Sciences for 24-mm Transwell) or

12-mm Snapwell permeable filters (#3801; Corning Life Sciences). Filters were precoated with a

1:20 dilution of Matrigel in cold DPBS for 2 hrs. and then removed just before seeding cells.

IntestiCult media (StemCell Technologies) was added to both apical and basolateral sides of the

filters to support cell growth. Once a confluent monolayer was obtained, IntestiCult

differentiation media (StemCell Technologies) was used to differentiate the cells prior to

experimentation. Cells were differentiated for one week before experiments.

Exposure of primary human colon organoids to OMVs. Polarized monolayers of colon

organoids on 12-mm Snapwell filters were washed with PBS to remove excess mucus, and 1.5

mL of differentiation media was added to the basolateral side. 2×10^{10} OMVs from all strains of

B. fragilis, a concentration of OMVs similar to that measured in conditioned media and in

biological fluids [21,39], or the same volume of process control (PC) media were applied to the

apical side of cells for 1 hr. and then removed. At the end of the 1hr. exposure and then 24 hrs.

later the apical and basolateral media were collected for measurements of cytokines

[MILLIPLEX MAP Human Cytokine/Chemokine 48-Plex cytokine assay (MilliporeSigma)].

Ussing chamber analysis was used to quantify CFTR Cl⁻ secretion, sodium absorption and TER

as described [61,62].

T84 cell culture. T84 cells, a human colon adenocarcinoma cell line (American Type Culture

Collection, Manassas, VA) were maintained in DMEM/F-12 media (ATCC) containing l-

glutamine supplemented with 5% fetal bovine serum (FBS: Life Technology). For short-circuit

current and TER measurements cells were cultured on Transwell filters (Corning, Ithaca, NY).

Experiments were performed 21 days postseeding as previously described [40-42].

Ussing chamber measurements of sodium and CFTR Cl⁻ currents. As described in detail

elsewhere [63-66] cells on Snapwell or Transwell filters were mounted in Ussing chambers. Primary

human colon organoids and T84 cells were exposed to either process control (PC: BHIS growth

media processed through the OMV isolation procedure, as a control for possible effects on

colon organoids due to the media or the OMV isolation process as recommended by the

International Society of Extracellular Vesicles [67]) or OMVs secreted by ETBF, ETBF Δ*bft*, NTBF,

or NTBF+*bft* were added to the apical side of monolayers for 1 hr. then CFTR Cl⁻ currents,

amiloride-sensitive sodium currents (in colon organoids only since T84 cells do display

amiloride-sensitive sodium currents) and TER were measured. Some monolayers were exposed

to OMVs for 1 hr., followed by washing OMVs from the apical surface and 24 hrs. later CFTR

Cl⁻ currents and TER were measured. To measure TER, a 1 mV pulse was applied to each

monolayer and the change in short circuit current was measured and TER was calculated using

Ohms Law. Electrogenic sodium currents were reported as amiloride-sensitive currents, since

amiloride inhibits the epithelial sodium channel ENaC [68]. To measure CFTR Cl⁻ secretion,

amiloride (50 μM) was first added to the apical solution to inhibit and measure sodium

reabsorption. Thereafter, Cl⁻ secretion was stimulated with forskolin (10 μM; Sigma-Aldrich),

which increases protein kinase A that phosphorylates and activates CFTR Cl⁻ secretion [38].

Finally, thiazolidinone (CFTR$_{inh}$-172, 20 μM; Millipore, Billerica, MA), an inhibitor of CFTR Cl⁻

channels, was added to the apical solution [64]. Data are expressed as the forskolin-stimulated,

CFTR$_{inh}$-172 inhibited short circuit current (Isc), which is presented as μA/cm^2. Data were

collected and analyzed using the Data Acquisition Software Acquire and Analyze program

(Physiologic Instruments, San Diego, CA).

Cytokine measurements. Cytokine secretion (apical and basolateral solutions) from primary

human colon organoids was measured with the MILLIPLEX MAP Human Cytokine/Chemokine

48-Plex cytokine assay (MilliporeSigma) following the manufactures instructions.

Immunocytochemistry. Immunocytochemistry was performed to characterize cell types in the

colon organoids and to evaluate the effects of OMVs on E-cadherin and claudin-2. Fixation of

colon organoids was performed using 4% paraformaldehyde for 15 mins. at room temperature,

followed by permeabilization with 0.25% Triton X-100 for 15 mins., blocking using 10% normal

goat serum in PBS, and primary antibody incubation at 4°C overnight. After washing, secondary

antibodies were added for 1 hr. at room temperature in the dark. SLC26A3 was labeled with a

primary anti-mouse antibody (1:250, Santa Cruz #sc-376187), and a secondary goat anti-

mouse-488 antibody (1:500, Santa Cruz #sc-376187). CFTR was labeled with a primary anti-

mouse antibody (1:100, Millipore # 05-583) and a secondary anti-mouse-488 antibody (1:500).

MUC2 was labeled with a primary anti-mouse antibody (1:300), and a secondary anti-mouse-

488 antibody (1:500, Abcam ab11197). Chromogranin A was labeled with a primary goat anti

rabbit antibody (1:100), and secondary goat anti rabbit-568 antibody (1:1000, Abcam ab15760).

Tight junction labeling was performed by fixing cells in 100% cold methanol at -20°C for 5 mins.,

blocking with 10% goat serum in PBS for 1 hr., and incubating with primary antibodies at 4°C

overnight. The secondary antibody was added for 1 hr. at room temperature the following day.

Claudin-2 was labeled using a primary goat anti-rabbit antibody (1:100, Thermo Fisher #51-

6100), and a secondary goat anti-rabbit-568 antibody (1:500). E-cadherin was labeled using a

primary goat anti-mouse antibody (1:250, BD Transduction lab #610182), and a secondary goat

anti-mouse-488 antibody (1:500). All antibodies were diluted in PBS with 5% normal goat

serum. DAPI staining to label nuclei was performed in fluorshield anti-fade mounting medium.

To image the colon organoids, a Yokogawa W1 spinning disk unit coupled to a Nikon Ti-E

inverted microscope was used. To examine the effect of OMVs and PC on claudin-2 and E-

cadherin, all settings on the confocal microscope were identical for acquiring images for

analysis. Analysis of claudin-2 and E-cadherin abundance was performed by an investigator

blinded to the treatments. The membrane areas with positive fluorescence signal were

segmented using trainable WEKA machine learning plugin in ImageJ [69]. The segmented objects

were thinned to 1 pixel using ImageJ's "Skeletonize" morphological operation. The mean and

median fluorescence intensities were measured from areas defined by the skeleton and

widened to 5 pixels using ImageJ's "Dilate" morphological operation.

Statistics. Data were analyzed using the R software environment for statistical computing and

graphics version 4.1.0 [70] , Ingenuity Pathway Analysis and Prism (GraphPad). Statistical

significance was calculated using either a mixed effect linear model, or likelihood ratio tests on

gene-wise negative binomial generalized linear models, as indicated in the figure legends. Data

were visualized, and figures were created using Prism or ggplot.

Author contributions

PS, RB, AN, BAS, CR, TTH and BR designed the research studies. PS, RB, AN, CR, TTH and

ZS conducted experiments on human colon organoids and T84 cells, acquired data, and

analyzed data. TBG recruited human subjects, provided informed consent and collected clinical

biopsies. YAG and BKC performed the proteomics analysis of OMVs. Electron microscopy of

OMVs was performed by DJT and Brad Vietje at the Microscopy Imaging Center at the

University of Vermont (RRID# SCR_018821). BAS wrote the first and revised draft of the

manuscript. All authors contributed to the article and approved the submitted version.

Declaration of competing interests

All authors declare no competing interests.

IRB statement

The Dartmouth Committee for the Protection of Human Subjects has determined that using

colon biopsies in this study is not considered human subject research because they contain no

patient identifiers. All tissues used for isolation were obtained under informed consent and

conform to HIPAA regulations to protect the privacy of the donor's personally identifiable

information.

Data availability. The mass spectrometry proteomics data are available in the Supplemental

Tables and have been deposited to the ProteomeXchange Consortium via the PRIDE partner

repository[60] with the dataset identifier PXD073462. All other reasonable requests for additional

data are available from the corresponding author.

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SUPPLEMENTAL TABLES

Supplemental Table 1. Mass spectrometry quantification of proteins detected exclusively in *B.*

fragilis NTBF OMVs.

Supplemental Table 2. Mass spectrometry quantification of proteins detected exclusively in *B.*

fragilis ETBF OMVs.

Supplemental Table 3. Mass spectrometry quantification of proteins detected in both *B. fragilis*

NTBF and *B. fragilis* ETBF OMVs.

Supplemental Table 4. Differential expression results of proteins identified in mass

spectrometry for both *B. fragilis* ETBF and *B. fragilis* NTBF

References

1   Jasemi, S. *et al.* Biological Mechanisms of Enterotoxigenic Bacteroides fragilis Toxin:

Linking Inflammation, Colorectal Cancer, and Clinical Implications. *Toxins (Basel)* **17**

(2025). https://doi.org/10.3390/toxins17060305

2   Ferranti, E. P., Dunbar, S. B., Dunlop, A. L. & Corwin, E. J. 20 things you didn't know

about the human gut microbiome. *J Cardiovasc Nurs* **29**, 479-481 (2014).

https://doi.org/10.1097/JCN.0000000000000166

3   Valguarnera, E. & Wardenburg, J. B. Good Gone Bad: One Toxin Away From Disease

for Bacteroides fragilis. *J Mol Biol* **432**, 765-785 (2020).

https://doi.org/10.1016/j.jmb.2019.12.003

4   Human Microbiome Project, C. Structure, function and diversity of the healthy human

microbiome. *Nature* **486**, 207-214 (2012). https://doi.org/10.1038/nature11234

5   Young, V. B. The role of the microbiome in human health and disease: an introduction

for clinicians. *BMJ* **356**, j831 (2017). https://doi.org/10.1136/bmj.j831

6   Saha, K., Zhou, Y. & Turner, J. R. Tight junction regulation, intestinal permeability, and

mucosal immunity in gastrointestinal health and disease. *Curr Opin Gastroenterol* **41**,

46-53 (2025). https://doi.org/10.1097/MOG.0000000000001066

7   Hill, C. A. *et al.* Bacteroides fragilis toxin expression enables lamina propria niche

acquisition in the developing mouse gut. *Nat Microbiol* **9**, 85-94 (2024).

https://doi.org/10.1038/s41564-023-01559-9

8   Sears, C. L. *et al.* Association of enterotoxigenic Bacteroides fragilis infection with

inflammatory diarrhea. *Clin Infect Dis* **47**, 797-803 (2008).

https://doi.org/10.1086/591130

9   Kim, W. S. *et al.* Bacteroides fragilis Toxin Induces Sequential Proteolysis of E-Cadherin

and Inflammatory Response in Mouse Intestinal Epithelial Cell Line. *Microorganisms* **13**

(2025). https://doi.org/10.3390/microorganisms13040781

10   Hwang, S., Gwon, S. Y., Kim, M. S., Lee, S. & Rhee, K. J. Bacteroides fragilis Toxin

Induces IL-8 Secretion in HT29/C1 Cells through Disruption of E-cadherin Junctions.

Immune Netw **13**, 213-217 (2013). https://doi.org/10.4110/in.2013.13.5.213

11   Guo, Y. *et al.* Mechanistic diversity of Bacteroides fragilis toxins and neutralization with

single domain antibody. *Cell Chem Biol* **33**, 102-116 e106 (2026).

https://doi.org/10.1016/j.chembiol.2025.12.009

12   Zakharzhevskaya, N. B. *et al.* Interaction of Bacteroides fragilis Toxin with Outer

Membrane Vesicles Reveals New Mechanism of Its Secretion and Delivery. *Front Cell*

Infect Microbiol **7**, 2 (2017). https://doi.org/10.3389/fcimb.2017.00002

13   Rong, Y. *et al.* Partially differentiated ileal and distal-colonic human F508del-cystic

fibrosis-enteroids secrete fluid in response to forskolin and linaclotide. *iScience* **28**,

112453 (2025). https://doi.org/10.1016/j.isci.2025.112453

14   Negussie, A. B., Dell, A. C., Davis, B. A. & Geibel, J. P. Colonic Fluid and Electrolyte

Transport 2022: An Update. *Cells* **11** (2022). https://doi.org/10.3390/cells11101712

15   Riegler, M. *et al.* Bacteroides fragilis toxin 2 damages human colonic mucosa in vitro.

Gut **44**, 504-510 (1999). https://doi.org/10.1136/gut.44.4.504

16   Obiso, R. J., Jr., Azghani, A. O. & Wilkins, T. D. The Bacteroides fragilis toxin fragilysin

disrupts the paracellular barrier of epithelial cells. *Infect Immun* **65**, 1431-1439 (1997).

https://doi.org/10.1128/iai.65.4.1431-1439.1997

17   Wu, S., Rhee, K. J., Zhang, M., Franco, A. & Sears, C. L. Bacteroides fragilis toxin

stimulates intestinal epithelial cell shedding and gamma-secretase-dependent E-

cadherin cleavage. *J Cell Sci* **120**, 1944-1952 (2007). https://doi.org/10.1242/jcs.03455

18   Sheikh, A. *et al.* Outer membrane vesicles from Bacteroides fragilis contain coding and

non-coding small RNA species that modulate inflammatory signaling in intestinal

epithelial cells. *bioRxiv* (2025). https://doi.org/10.1101/2025.06.25.661399

19   Kharlampieva, D. D. *et al.* Purification and characterisation of recombinant Bacteroides

fragilis toxin-2. *Biochimie* **95**, 2123-2131 (2013).

https://doi.org/10.1016/j.biochi.2013.08.005

20   Chambers, F. G. *et al.* Bacteroides fragilis toxin exhibits polar activity on monolayers of

human intestinal epithelial cells (T84 cells) in vitro. *Infect Immun* **65**, 3561-3570 (1997).

https://doi.org/10.1128/iai.65.9.3561-3570.1997

21   Charpentier, L. A. *et al.* Bacterial Outer Membrane Vesicles and Immune Modulation of

the Host. *Membranes (Basel)* **13** (2023). https://doi.org/10.3390/membranes13090752

22 Li, Z. *et al.* P. aeruginosa tRNA-fMet halves secreted in outer membrane vesicles

suppress lung inflammation in cystic fibrosis. *Am J Physiol Lung Cell Mol Physiol* **326**,

L574-L588 (2024). https://doi.org/10.1152/ajplung.00018.2024

23 Koeppen, K. *et al.* A Novel Mechanism of Host-Pathogen Interaction through sRNA in

Bacterial Outer Membrane Vesicles. *PLoS Pathog* **12**, e1005672 (2016).

https://doi.org/10.1371/journal.ppat.1005672

24 Yang, Y. *et al.* Bacteroides Fragilis-Derived Outer Membrane Vesicles Deliver MiR-5119

and Alleviate Colitis by Targeting PD-L1 to Inhibit GSDMD-Mediated Neutrophil

Extracellular Trap Formation. *Adv Sci (Weinh)*, e00781 (2025).

https://doi.org/10.1002/advs.202500781

25 Shagaleeva, O. Y. *et al.* Bacteroides vesicles promote functional alterations in the gut

microbiota composition. *Microbiol Spectr* **12**, e0063624 (2024).

https://doi.org/10.1128/spectrum.00636-24

26 Patrick, S., McKenna, J. P., O'Hagan, S. & Dermott, E. A comparison of the

haemagglutinating and enzymic activities of Bacteroides fragilis whole cells and outer

membrane vesicles. *Microb Pathog* **20**, 191-202 (1996).

https://doi.org/10.1006/mpat.1996.0018

27 Sharpe, S. W., Kuehn, M. J. & Mason, K. M. Elicitation of epithelial cell-derived immune

effectors by outer membrane vesicles of nontypeable Haemophilus influenzae. *Infect*

Immun **79**, 4361-4369 (2011). https://doi.org/10.1128/IAI.05332-11

28 Round, J. L. *et al.* The Toll-like receptor 2 pathway establishes colonization by a

commensal of the human microbiota. *Science* **332**, 974-977 (2011).

https://doi.org/10.1126/science.1206095

29 Lee, S. M. *et al.* Bacterial colonization factors control specificity and stability of the gut

microbiota. *Nature* **501**, 426-429 (2013). https://doi.org/10.1038/nature12447

30 Sears, C. L., Geis, A. L. & Housseau, F. Bacteroides fragilis subverts mucosal biology:

from symbiont to colon carcinogenesis. *J Clin Invest* **124**, 4166-4172 (2014).

https://doi.org/10.1172/JCI72334

31 Taglieri, M. *et al.* Colorectal Organoids: Models, Imaging, Omics, Therapy, Immunology,

and Ethics. *Cells* **14** (2025). https://doi.org/10.3390/cells14060457

32 Sen, A. *et al.* Comprehensive gene expression analysis of organoid-derived healthy

human colonic epithelium and cancer cell line stimulated with live probiotic bacteria. *Sci*

Rep **15**, 22325 (2025). https://doi.org/10.1038/s41598-025-07391-x

33 Rikkert, L. G., Nieuwland, R., Terstappen, L. & Coumans, F. A. W. Quality of

extracellular vesicle images by transmission electron microscopy is operator and

protocol dependent. *J Extracell Vesicles* **8**, 1555419 (2019).

https://doi.org/10.1080/20013078.2018.1555419

34 Noble, J. M. *et al.* Direct comparison of optical and electron microscopy methods for

structural characterization of extracellular vesicles. *J Struct Biol* **210**, 107474 (2020).

https://doi.org/10.1016/j.jsb.2020.107474

35 Weiss, A. *et al.* Transmission Electron Microscopy-based characterization of

Extracellular Vesicles from plasma and serum from Parkinson s Disease patients. *Cell*

Commun Signal **23**, 395 (2025). https://doi.org/10.1186/s12964-025-02383-w

36 Zakharzhevskaya, N. B. *et al.* Outer membrane vesicles secreted by pathogenic and

nonpathogenic Bacteroides fragilis represent different metabolic activities. *Sci Rep* **7**,

5008 (2017). https://doi.org/10.1038/s41598-017-05264-6

37 Parikh, K. *et al.* Colonic epithelial cell diversity in health and inflammatory bowel disease.

Nature **567**, 49-55 (2019). https://doi.org/10.1038/s41586-019-0992-y

38 Ghigo, A., De Santi, C., Hart, M., Mitash, N. & Swiatecka-Urban, A. Cell signaling and

regulation of CFTR expression in cystic fibrosis cells in the era of high efficiency

modulator therapy. *J Cyst Fibros* **22 Suppl 1**, S12-S16 (2023).

https://doi.org/10.1016/j.jcf.2022.12.015

39   Stanton, B. A. Extracellular Vesicles and Host-Pathogen Interactions: A Review of Inter-

Kingdom Signaling by Small Noncoding RNA. *Genes (Basel)* **12** (2021).

https://doi.org/10.3390/genes12071010

40   Howe, K. L., Wang, A., Hunter, M. M., Stanton, B. A. & McKay, D. M. TGFbeta down-

regulation of the CFTR: a means to limit epithelial chloride secretion. *Exp Cell Res* **298**,

473-484 (2004). https://doi.org/10.1016/j.yexcr.2004.04.026

41   Maitra, R., Shaw, C. M., Stanton, B. A. & Hamilton, J. W. Increased functional cell

surface expression of CFTR and DeltaF508-CFTR by the anthracycline doxorubicin. *Am*

J Physiol Cell Physiol **280**, C1031-1037 (2001).

https://doi.org/10.1152/ajpcell.2001.280.5.C1031

42   Maitra, R., Shaw, C. M., Stanton, B. A. & Hamilton, J. W. Functional enhancement of

CFTR expression by mitomycin C. *Cell Physiol Biochem* **11**, 93-98 (2001).

https://doi.org/10.1159/000047796

43   Pierce, J. V., Fellows, J. D., Anderson, D. E. & Bernstein, H. D. A clostripain-like

protease plays a major role in generating the secretome of enterotoxigenic Bacteroides

fragilis. *Mol Microbiol* **115**, 290-304 (2021). https://doi.org/10.1111/mmi.14616

44   Choi, V. M. *et al.* Activation of Bacteroides fragilis toxin by a novel bacterial protease

contributes to anaerobic sepsis in mice. *Nat Med* **22**, 563-567 (2016).

https://doi.org/10.1038/nm.4077

45   Elhenawy, W., Debelyy, M. O. & Feldman, M. F. Preferential packing of acidic

glycosidases and proteases into Bacteroides outer membrane vesicles. *MBio* **5**, e00909-

00914 (2014). https://doi.org/10.1128/mBio.00909-14

46   France, M. M. & Turner, J. R. The mucosal barrier at a glance. *J Cell Sci* **130**, 307-314

(2017). https://doi.org/10.1242/jcs.193482

47   Citi, S. *et al.* A short guide to the tight junction. *J Cell Sci* **137** (2024).

https://doi.org/10.1242/jcs.261776

48   Han, Y. W. *et al.* Interactions between periodontal bacteria and human oral epithelial

cells: Fusobacterium nucleatum adheres to and invades epithelial cells. *Infect Immun* **68**,

3140-3146 (2000). https://doi.org/10.1128/IAI.68.6.3140-3146.2000

49   Antosca, K. M. *et al.* Altered Stool Microbiota of Infants with Cystic Fibrosis Shows a

Reduction in Genera Associated with Immune Programming from Birth. *J Bacteriol* **201**

(2019). https://doi.org/10.1128/JB.00274-19

50   Lee, C. G. *et al.* Bacteroides fragilis Toxin Induces Intestinal Epithelial Cell Secretion of

Interleukin-8 by the E-Cadherin/beta-Catenin/NF-kappaB Dependent Pathway.

Biomedicines **10** (2022). https://doi.org/10.3390/biomedicines10040827

51   Choi, J. W., Kim, S. C., Hong, S. H. & Lee, H. J. Secretable Small RNAs via Outer

Membrane Vesicles in Periodontal Pathogens. *J Dent Res* **96**, 458-466 (2017).

https://doi.org/10.1177/0022034516685071

52   Franco, A. A. The Bacteroides fragilis pathogenicity island is contained in a putative

novel conjugative transposon. *J Bacteriol* **186**, 6077-6092 (2004).

https://doi.org/10.1128/JB.186.18.6077-6092.2004

53   Wu, S. *et al.* A human colonic commensal promotes colon tumorigenesis via activation

of T helper type 17 T cell responses. *Nat Med* **15**, 1016-1022 (2009).

https://doi.org/10.1038/nm.2015

54   Chan, J. L. *et al.* Non-toxigenic Bacteroides fragilis (NTBF) administration reduces

bacteria-driven chronic colitis and tumor development independent of polysaccharide A.

Mucosal Immunol **12**, 164-177 (2019). https://doi.org/10.1038/s41385-018-0085-5

55   Chung, L. *et al.* Bacteroides fragilis Toxin Coordinates a Pro-carcinogenic Inflammatory

Cascade via Targeting of Colonic Epithelial Cells. *Cell Host Microbe* **23**, 203-214 e205

(2018). https://doi.org/10.1016/j.chom.2018.01.007

56   Lim, B., Zimmermann, M., Barry, N. A. & Goodman, A. L. Engineered Regulatory

Systems Modulate Gene Expression of Human Commensals in the Gut. *Cell* **169**, 547-

558 e515 (2017). https://doi.org/10.1016/j.cell.2017.03.045

57   Degnan, P. H., Barry, N. A., Mok, K. C., Taga, M. E. & Goodman, A. L. Human gut

microbes use multiple transporters to distinguish vitamin B(1)(2) analogs and compete in

the gut. *Cell Host Microbe* **15**, 47-57 (2014). https://doi.org/10.1016/j.chom.2013.12.007

58   Bauman, S. J. & Kuehn, M. J. Purification of outer membrane vesicles from

Pseudomonas aeruginosa and their activation of an IL-8 response. *Microbes Infect* **8**,

2400-2408 (2006). https://doi.org/10.1016/j.micinf.2006.05.001

59   Koeppen, K. *et al.* An rRNA fragment in extracellular vesicles secreted by human airway

epithelial cells increases the fluoroquinolone sensitivity of P. aeruginosa. *Am J Physiol*

Lung Cell Mol Physiol **325**, L54-L65 (2023). https://doi.org/10.1152/ajplung.00150.2022

60   Perez-Riverol, Y. *et al.* The PRIDE database at 20 years: 2025 update. *Nucleic Acids*

Res **53**, D543-D553 (2025). https://doi.org/10.1093/nar/gkae1011

61   Hampton, T. H., Koeppen, K., Bashor, L. & Stanton, B. A. Selection of reference genes

for quantitative PCR: identifying reference genes for airway epithelial cells exposed to

Pseudomonas aeruginosa. *Am J Physiol Lung Cell Mol Physiol* **319**, L256-L265 (2020).

https://doi.org/10.1152/ajplung.00158.2020

62   Hampton, T. H. *et al.* Gene expression responses of CF airway epithelial cells exposed

to elexacaftor/tezacaftor/ivacaftor (ETI) suggest benefits beyond improved CFTR

channel function. *Am J Physiol Lung Cell Mol Physiol.* **Dec 1;327(6):L905-L916.** (2024).

https://doi.org/ doi: 10.1152/ajplung.00272.2024.

63   Stanton, B. A., Coutermarsh, B., Barnaby, R. & Hogan, D. Pseudomonas aeruginosa

Reduces VX-809 Stimulated F508del-CFTR Chloride Secretion by Airway Epithelial

Cells. *PLoS One* **10**, e0127742 (2015). https://doi.org/10.1371/journal.pone.0127742

64   Barnaby, R., Koeppen, K. & Stanton, B. A. Cyclodextrins reduce the ability of

Pseudomonas aeruginosa outer-membrane vesicles to reduce CFTR Cl(-) secretion. *Am*

J Physiol Lung Cell Mol Physiol **316**, L206-L215 (2019).

https://doi.org/10.1152/ajplung.00316.2018

65   Talebian, L., Coutermarsh, B., Channon, J. Y. & Stanton, B. A. Corr4A and VRT325 do

not reduce the inflammatory response to P. aeruginosa in human cystic fibrosis airway

epithelial cells. *Cell Physiol Biochem* **23**, 199-204 (2009).

https://doi.org/10.1159/000204108

66   Stanton, B. A. Effects of Pseudomonas aeruginosa on CFTR chloride secretion and the

host immune response. *Am J Physiol Cell Physiol* **312**, C357-C366 (2017).

https://doi.org/10.1152/ajpcell.00373.2016

67   Welsh, J. A. *et al.* Minimal information for studies of extracellular vesicles (MISEV2023):

From basic to advanced approaches. *J Extracell Vesicles* **13**, e12404 (2024).

https://doi.org/10.1002/jev2.12404

68   Berdiev, B. K., Shlyonsky, V. G., Karlson, K. H., Stanton, B. A. & Ismailov, II. Gating of

amiloride-sensitive Na(+) channels: subunit-subunit interactions and inhibition by the

cystic fibrosis transmembrane conductance regulator. *Biophys J* **78**, 1881-1894 (2000).

https://doi.org/10.1016/S0006-3495(00)76737-3

69   Arganda-Carreras, I. *et al.* Trainable Weka Segmentation: a machine learning tool for

microscopy pixel classification. *Bioinformatics* **33**, 2424-2426 (2017).

https://doi.org/10.1093/bioinformatics/btx180

70   R Core Team (2017) R: A Language and Environment for Statistical Computing.

https://www.R-project.org/