

RESEARCH PAPER



Function of Multidrug Resistance Transporters is Disrupted by Infection Mimics in Human Brain Endothelial Cells

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ABSTRACT

P-glycoprotein (*P-gp/ABCB1*) and breast cancer resistance protein (*BCRP/ABCG2*) modulate the distribution of drugs and toxins across the blood-brain barrier (BBB). Animal studies reported that infection-induced disruption of these transporters in the developing BBB impairs fetal brain protection. However, the impact of infection mimics on *P-gp/BCRP* function in human brain endothelium is less well understood. We hypothesized that Toll-like receptor ligands mimicking bacterial and viral infection would modify the expression and function of *P-gp* and *BCRP* in human brain endothelial cells (BECs). Human cerebral microvascular endothelial cells (hCMEC/D3) were challenged with bacterial [Lipopolysaccharide (LPS)] and viral-mimics [polyinosinic:polycytidylic acid (PolyI:C) or single-stranded RNA (ssRNA)], or pro-inflammatory cytokines interleukin (IL)-6, tumor necrosis factor (TNF)- α and interferon gamma (IFN)- γ . *P-gp* and *BCRP* function was assessed after 4 or 24 h, using Calcein-AM and Chlorin-6 assays, respectively. Western blot and qPCR quantified *P-gp/ABCB1* and *BCRP/ABCG2* expression following treatments. Infection mimics are potent modulators of drug transporters in human BECs *in vitro*. LPS and PolyI:C increased, while ssRNA exposure reduced *P-gp* activity. In contrast, LPS and PolyI:C decreased, while ssRNA increased *BCRP* activity ($P < .05$). There was little correlation between drug transporter function, gene expression and total protein level. Altered plasma membrane *BCRP* may suggest modified intracellular trafficking induced by infection in human BECs. Bacterial and viral infection mimics modify *P-gp* and *BCRP* transport function in human BECs, *in vitro*. This knowledge may contribute and have important implications for human brain protection and possible altered biodistribution of drugs and xenobiotics in the brain following exposure to TLR agonists.

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Introduction


The blood-brain barrier (BBB) is a selective barrier formed by specialized brain endothelial cells (BECs) that protect the brain from invasive pathogens and penetration of toxic compounds from the bloodstream.¹ The multidrug-resistance transporters, *P-glycoprotein (P-gp; encoded by ABCB1)* and breast cancer resistance protein (*BCRP; ABCG2*), belong to the ATP-binding cassette (ABC) transporters superfamily of efflux transporters and are central protective components of the BBB.² They are present at the luminal surface of the cerebral microvascular endothelium, where they actively efflux endogenous (steroid hormones, cytokines, chemokines) and exogenous substrates (analgesics, antiretrovirals, antiepileptics, antibiotics, synthetic glucocorticoids, selective

serotonin reuptake inhibitor (SSRIs), agrochemicals), back into the bloodstream. As such, they act to control accumulation of their substrates in the central nervous system (CNS).²

The development of the mammalian brain is highly vulnerable to xenobiotics and environmental toxins.^{1,3} Using animal models, our group demonstrated that BBB drug transport function is regulated by early exposure to several factors, including the pro-inflammatory cytokines interleukin (IL)-1 β , IL-6, and tumor necrosis factor (TNF)- α . These cytokines inhibit *P-gp* function in BECs, while exposure to glucocorticoids and transforming growth factor- β 1 (TGF- β 1) up-regulates *P-gp* function.^{1,4-7} Therefore, *P-gp* activity in BECs (the major component of the BBB) may either be reduced or increased depending

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on the nature of stimuli, resulting in altered accumulation of xenobiotics in the brain.

No studies have investigated the impact of bacterial or viral infection on drug transport function in BECs in adulthood or during development in humans. In the present study, we sought to investigate the impact of infection mimics on *P*-gp and BCRP transport function using an *in vitro* human brain endothelium model. We hypothesized that bacterial and viral mimics alter *P*-gp and BCRP function in the human brain endothelium. We challenged human cerebral microvascular endothelial cells (hCMEC/D3) with bacterial- and viral-mimics and key infection-associated cytokines and demonstrated that control (i.e. inhibition or activation) of human BEC function is dependent on the nature of infective/inflammatory stimuli.

Material and methods

Cell culture and reagents

The human cerebral microvascular endothelial cell line (hCMEC/D3, Cedarlane Labs #CLU512, Burlington, ON, Canada) was selected as a model to investigate drug transporters function since it has been extensively used and characterized to have a brain endothelial phenotype and is, thus a widely used model to investigate human BBB function.^{8–10} The cells have similar morphology and protein expression levels to primary human cerebral microvascular endothelial cells, and they are well adapted for drug transport studies.¹¹ The cells were purchased between passages 25 and 26 (P25/26), and cultured at 37°C in 5% CO₂ with EndoGROTM-MV Complete Culture Media Kit® (Millipore, ON, Canada, #SCME004), consisting of EndoGRO Basal Medium and EndoGRO Supplement kit containing recombinant human epidermal growth factor (5 ng/mL), L-glutamine (10 mM), hydrocortisone hemisuccinate (1.0 µg/mL), heparin sulfate (0.75 U/mL), ascorbic acid (50 µg/mL) and fetal bovine serum (5%; FBS). The media was supplemented with human basic fibroblast growth factor (1 ng/mL; Sigma, #F0291) and penicillin-streptomycin (1%; 10000 units – 10000 µg/mL, Life Technologies, #15140-122). Cells were expanded through P29, frozen in 5% DMSO in FBS for 24 h (–80°C), and then stored in liquid nitrogen. All procedures, treatments, and analyses were performed

with hCMEC/D3 cells at P30. Lipopolysaccharide (LPS, #L4391), polyinosinic:polycytidylic acid (PolyI:C, #P9582), and calcein-acetoxymethyl ester (Ca-AM, #177831) were purchased from Sigma-Aldrich (Oakville, ON, Canada). Single-stranded RNA (ssRNA, #tlr-lrna40) and its vehicle Lyovec (#lyec-12) were purchased from InvivoGen (San Diego, California, USA), and the cytokines IL-6 (#PHC0065), TNF-α (#PHC3015) and interferon-gamma (IFN)-γ (#PHC4031) were purchased from Gibco (Grand Island, New York, USA).

Baseline immunohistochemical characterization

Immunohistochemical analysis was performed to confirm the phenotype of the human BEC cell line. Cells were rinsed twice with cold PBS, fixed with ice-cold Methanol for 10 min, and then permeabilized with 0.2% Triton X-100 (5 min; room temperature). Cells were then blocked with 0.1% BSA in PBS (1 h), and incubated with primary antibodies anti-*P*-gp (Santa Cruz Biotechnology, Dallas, Texas, USA, #SC-55510, 1:200), anti-BCRP (Millipore, #MAB146, 1:200), anti-Claudin-5 (Invitrogen #352588, 1:200), anti-PECAM-1 (Dako, Burloak, ON, Canada, #M0823, 1:200) and anti-GFAP (Cell Signaling Technology, Whitby, ON, Canada, #3670, 1:300) overnight at 4°C. The primary antibodies anti-rabbit IgG (Abcam, Toronto, ON, Canada, #ab171870), anti-mouse IgG1 (Dako, #X0931), anti-mouse IgG2a (Dako, #X0943), and anti-mouse IgG2b (Dako, #X0944) were used as isotype controls. Subsequently, cells were washed (3X) and incubated with fluorochrome-labeled secondary antibody Alexa Fluor 488 (Molecular Probes Inc., Eugene, OR, USA, 1:300) and counterstained with DAPI (1 mg/mL, 60 min). The area and intensity of antibody staining were analyzed, and representative images were obtained using a spinning disc confocal fluorescent microscope (Leica DMI6000 B, Concord, ON, Canada) at 40X magnification.

Experimental design

hCMEC/D3 cells (P30) were plated in clear flat bottom 6-well or 96-well TC-treated polystyrene culture plates (Costar, Kennebunk, ME, USA, #3516; or Falcon, Mississauga, ON, Canada, #353072, respectively) at a density of 20000 cells/

cm² with EndoGROTM, and 5% charcoal-stripped (CS)-FBS in DMEM. After 24 h of incubation, cells were challenged with the Toll-like receptor (TLR)-4 ligand, LPS (10³⁻⁶ pg/mL bacterial mimic); the TLR-3 ligand PolyI:C (10³⁻⁶ pg/mL; double-stranded viral mimic) or the TLR-7/8 ligand ssRNA (10³⁻⁶ pg/mL; single-stranded viral mimic), to mimic bacterial and viral infection on human BECs, *in vitro*. The cytokines IL-6 (10¹⁻⁴ pg/mL), TNF- α (10¹⁻⁴ pg/mL), and IFN- γ (10¹⁻⁴ pg/mL) were used to determine the action of downstream mediators induced by infection/inflammation. Functional drug transport assays were undertaken or cells were harvested for total/plasma membrane protein or mRNA analysis 4- and 24 h after treatments. Dose and time of exposure regimens were selected based on previous studies from our group.^{4,6}

Gene expression analyses

Total RNA was extracted using the RNeasy Mini Kit (Qiagen #74104, Toronto, ON, Canada), according to the manufacturer's instructions. RNA concentration and purity were determined using a NanoDrop1000 Spectrophotometer (Thermo Scientific, Wilmington, DE, USA) and total RNA (1 μ g) was reverse transcribed to cDNA using iScriptTM Reverse Transcription Supermix (Bio-Rad #1708840, Mississauga, ON, Canada). The sequences of the primers used are listed in Table 1. SYBR Green reagent (Sigma #S9194) was used to run the qPCR, using the CFX 380 Real-Time system C1000 TM Thermal Cycler (Bio-Rad). The parameters used were: one cycle (95°C, 2 min), followed by 40 cycles (95°C, 5s) and 60°C for 20s. RNA (10 ng) was used per reaction and each sample was run in triplicate. Hypoxanthine-guanine

phosphoribosyltransferase (*HPRT*), TATA-box binding protein (*TBP*), and tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein zeta (*YWHAZ*) were used as reference genes and their geometric mean was used to normalize expression of our genes of interest. The relative mRNA expression was calculated by comparing the PCR cycle threshold (CT) between groups using the 2^{- $\Delta\Delta$ CT} Method,¹⁴ and the control group was normalized to 1.

P-gp and BCRP functional assays

P-gp activity was assessed as described previously by our group with adaptations.^{4-7,15} Briefly, charcoal-stripped (CS)-FBS (5%) was withdrawn and cells were washed twice with warm Tyrode salts' solution (Sigma, #T2145) supplemented with sodium bicarbonate (1 g/L; Sigma, #S6014). Cells were incubated with 10⁻⁶ M Ca-AM at 37°C and 5% CO₂ (1 h). Ca-AM is intracellularly cleaved by endogenous esterases, producing fluorescent calcein that can be used as an indirect measure of *P*-gp activity, since it is transported out of cells by *P*-gp. After incubation with Ca-AM, the plates were placed on ice and the cells were washed twice with ice-cold PBS, followed by cell lysis with 1% Triton X-100 (Sigma #X100) lysis buffer. The BCRP activity assay was adapted from a previous study¹⁶ using Chlorin 6 (Ce6, Santa Cruz Biotechnology, #SC-263067) as a specific BCRP substrate.¹⁷ Cells were incubated with 2 nM Ce6 (37°C; 5% CO₂; 1 h). Plates were placed on ice, cells were washed (2X) with cold Tyrode salt solution and lysis buffer added. Calcein and Ce6 accumulation were quantified using a fluorescent microplate reader, at excitation/emission wavelengths of 485/510 nm and 407/667 nm, respectively. The uptake of Ca-AM and Ce6 by *P*-gp and BCRP was confirmed under the

Table 1. Sequence of the primers used to run the qPCR in this study.

Primer	Forward	Reverse	Reference
<i>ABCB1</i>	agc aga ggc cgc cgc tgt tgc tt	cca ttc cga cct cgc gct cc	12
<i>ABCG2</i>	tgg aat cca gaa cag agct ggg gt	aga gtt cca cgg ctg aaa cac tgc	12
<i>HPRT</i>	tga cac tgg caa aac aat gca	ggg cct ttt cac cag caa gct	13
<i>IL-6</i>	tca atg agg aga ctt gcc tg	tgg ctt gtt cct cac tac tct	*
<i>YWHAZ</i>	act ttt ggt aca ttg tgg ctt caa	ccg cca gga caa acc agt at	12
<i>TLR-3</i>	tta cga aga ggc tgg aat gg	agg aac tcc ttt gcc ttg gt	12
<i>TLR-4</i>	aca gga aac ccc atc cag ag	att tgt ctc cac agc cac ca	12
<i>TLR-8</i>	tta tgt gtt cca gga act cag aga a	taa tac cca agt tga tag tgc ata agt ttg	*
<i>TBP</i>	tgc aca gga gcc aag agt gaa	cac atc aca gct ccc cac ca	13

*Primers designed using Primer-Blast tool at the National Center for Biotechnology Information website <<https://www.ncbi.nlm.nih.gov/tools/primer-blast/>>

exposure to their specific inhibitors, Verapamil and Ko143, respectively, to validate the assays (see Supplementary Figure 1). $N = 6-8$ was used per group; assays were repeated 2–4 times. Mean background readings were subtracted and *P*-gp and BCRP activity following treatment were expressed as a percentage of controls.

Protein expression

Cells were detached and lysed using lysis buffer (1 mol/L Tris-HCL pH 6.8, 2% SDS, 10% glycerol, containing protease and phosphatase inhibitor cocktail from Thermo Scientific (Mississauga, ON, Canada, #78440) and extracted by sonication. Protein concentration was determined using the Pierce BCA Protein assay kit (Thermo Scientific). Plasma membrane fractions were obtained using the Minute™ Plasma Membrane Protein Isolation and Cell Fractionation kit (Invent Biotechnologies, Plymouth, MN, USA, #SM-005), following the manufacturers' instructions. Total (30 μ g) or plasma membrane (10 μ g) proteins were separated by electrophoresis (100 V, 1 h) using SDS polyacrylamide gels (8%) and then transferred to polyvinylidene fluoride membranes using the Trans-Blot® Turbo™ Transfer System (Bio-Rad; 10 min). Membranes for BCRP analysis were blocked with skim milk (5%), while those for *P*-gp analysis were blocked with bovine serum albumin (BSA; 5%), in Tris-buffered saline containing 0.1% Tween (TBS-T; 1 h). Blots were incubated overnight (4°C) with the following primary antibodies: anti-*P*-gp (Abcam #ab170903, 1:1000), anti-BCRP (Abcam #ab108312, 1:2000), anti-ERK2 (Santa Cruz Biotechnology #SC-292838, 1:2000), and anti- Na^+/K^+ -ATPase (Abcam, #ab76020, 1:20000). Membranes were then washed (3X) with TBS-T, and incubated (1 h) with HRP-linked anti-rabbit secondary antibody (1:10000; GE Healthcare Bio-Science, Baie d'Urfe, QC, Canada). Protein-antibody complexes were detected by incubating the membranes with Laminate Crescendo Western HRP Substrate (5 mins; Millipore). Chemiluminescence was detected under UV using ChemiDoc™ MP Imaging system (Bio-Rad). BCRP and *P*-gp protein band intensity were quantified using Image Lab™ software and normalized against ERK2 or Na^+/K^+ -ATPase signals for total or plasma membrane protein levels assessment, respectively.

Statistical analyses

Statistical analysis was performed using Prism (version 7.0; GraphPad Software, Inc., San Diego, CA). Data were assessed for normal distribution using D'Agostino and Pearson or the Shapiro-Wilk test; any outliers were excluded using Grubbs' test. Gene and protein expression were analyzed using unpaired Student's *t*-test. Functional assays for *P*-gp and BCRP were analyzed using the Kruskal-Wallis test, followed by Dunn's multiple comparison test, comparing the different treatments to the control (vehicle) group. Number of samples (*n*) varied from 3 to 8 per group, depending on the experiment. Specific *n* values are identified in each table/figure legend. Differences were considered significant when *p*-value < 0.05. Data are presented as mean \pm standard error of the mean (SEM).

Results

Characterization of human BECs

Immunohistochemical analysis confirmed the presence of *P*-gp and BCRP in hCMEC/D3 cells [Figure 1](#). hCMEC/D3 cells also expressed claudin-5 and PECAM protein, consistent with a BEC phenotype. The astrocyte marker GFAP was not present. Mouse IgG1 was used as a negative control for GFAP and PECAM, and mouse IgG2a and 2b (insets) in [Figure 1](#) were used as negative controls for BCRP, *P*-gp and Claudin-5, respectively. No staining in negative controls was detected.

Bacterial- and viral-mimics and pro-inflammatory cytokine responses

Bacterial- and viral-mimics evoked a pro-inflammatory response in human BECs ([Supplementary Figure 2](#)). LPS induced a dramatic increase in *IL-6* mRNA expression, at 4 h and 24 h ($p < .001$ and $p < .05$). PolyI:C increased *IL-6* mRNA 24 h ($p < .05$) after exposure. ssRNA induced a near significant increase in *IL-6* mRNA at 4 h ($p = .061$) with a similar trend at 24 h. The responsiveness of *TLR* to their respective ligand, as well as potential cross-regulation, was investigated in human BEC 4 h after ligand stimulation ([Supplementary Figure 2](#)). LPS increased *TLR-4* ($p < .01$), with concomitant up-regulation

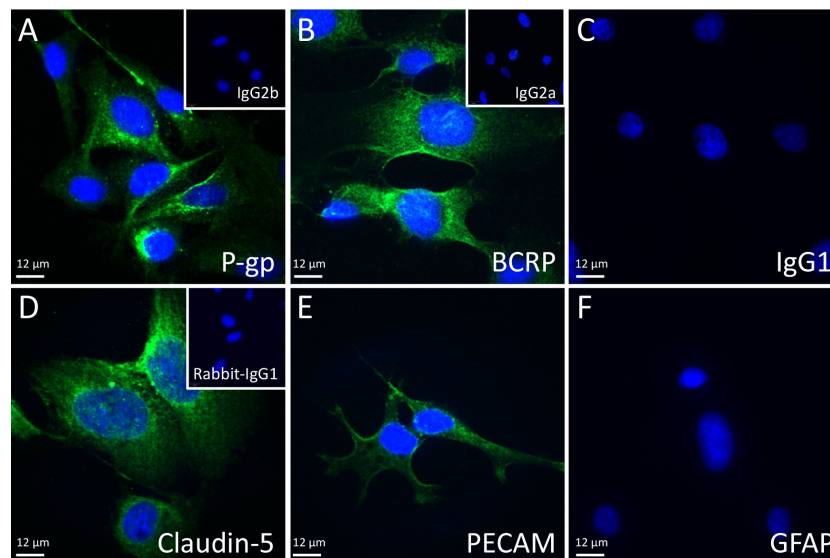


Figure 1. Representative fluorescent immunohistochemical images used for hCMEC/D3 characterization. Blue indicates DAPI staining, green represents specific antibody staining: A) *P-gp*, B) BCRP, C) IgG1, D) Claudin-5, E) PECAM, and F) GFAP. Insets show IgG2b (a), IgG2a (b) and rabbit IgG (d), 40X objective. Scale bars = 12 μm .

of *TLR-3* ($p < .001$) mRNA. PolyI:C increased *TLR-3* mRNA ($p < .05$), with no change in *TLR-4* or *TLR-8* mRNA. ssRNA exposure had no effect of *TLRs* mRNA levels.

The responses of cytoplasmic recognition receptors retinoid-acid-inducible gene I (*RIG-I*) and melanoma differentiation associated gene 5 (*MDA5*) to bacterial and viral mimics are presented in Supplementary Figure 3. *RIG-I* and *MDA5* recognize ssRNA and PolyI:C in the cytoplasm,¹⁸ activating an antiviral response mediated by nuclear factor- κB (NF- κB) and IFN regulatory factors (IRFs).¹⁹ LPS increased *RIG-I* ($p < .001$) and *MDA5* ($p < .05$ and $p < .001$) mRNA levels 4 h and 24 h after exposure. PolyI:C increased *MDA5* ($p < .05$) mRNA expression 24 h post-treatment, while ssRNA had no effect on *RIG-I* or *MDA5* mRNA.

***P-gp* and BCRP function**

As expected, exposure of hCMEC/D3 cells to Verapamil and Ko143 significantly increased Ca-AM and Ce6 accumulation, respectively. This confirms correct function of the activity assays (Supplementary Figure 1). LPS (10^5 pg/mL) treatment increased ($p < .01$) *P-gp* activity at 24 h, whereas BCRP activity was decreased in response to LPS (10^6 pg/mL, $p < .05$) at 24 h Figure 2. PolyI:C treatment (10^6 pg/mL) increased ($p < .05$) *P-gp* activity at 24 h, while PolyI:

C (10^4 and 10^5 pg/mL) decreased ($p < .05$, $p < .01$) BCRP activity 24 h after exposure. ssRNA treatment led to a marked decrease in *P-gp* function 4 h (10^4 pg/mL, $p < .01$; 10^5 pg/mL, $p < .05$) and 24 h (10^4 pg/mL, $p < .05$; 10^5 and 10^6 pg/mL, $p < .01$) after treatment. In contrast, ssRNA exposure resulted in a pronounced increase in BCRP function at 24 h (10^4 pg/mL, $p < .01$; 10^5 pg/mL, $p < .05$; Figure 2).

Expression of *P-gp/ABCB1* and BCRP/ABCG2

The effects of LPS on *ABCB1* and *ABCG2* mRNA levels and *P-gp* and BCRP total protein are illustrated in Table 2 and Figure 3, respectively. LPS decreased both *ABCB1* and *ABCG2* mRNA levels at 24 h ($p < .05$; Table 2), and reduced *P-gp* protein expression at 4 h ($p < .05$), however BCRP protein levels were increased by LPS 4 h post-treatment ($p < .05$). PolyI:C had no effect on *ABCB1* or *ABCG2* mRNA or *P-gp* protein at either time-point, but increased BCRP expression at 4 h of treatment ($p < .05$). ssRNA exposure up-regulated *ABCB1* mRNA at 24 h, and *ABCG2* at 4 h post-treatment, respectively ($p < .01$). While there was no effect of ssRNA on total *P-gp*, total BCRP protein levels were increased at 4 h ($p < .05$).

The inconsistency between total cellular *P-gp* and BCRP protein levels and *P-gp* and BCRP function following exposure to bacterial and viral mimics, led us to undertake analysis of *P-gp* and

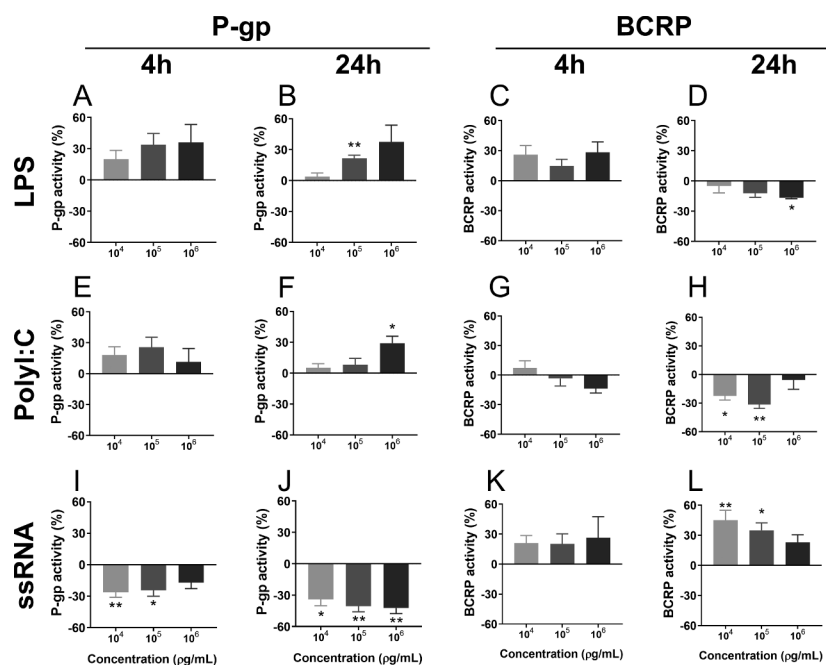


Figure 2. P-gp and BCRP function 4 h or 24 h after treatment with LPS (a-d), PolyI:C (e-h) or ssRNA (i-l), measured using the Ca-AM and Ce6 assays. Doses: 10^3 – 10^6 pg/mL. N = 6/group. Values are expressed as a percentage of control (vehicle) groups. Data are presented as mean \pm SEM. Statistical analysis: Kruskal-Wallis test, followed by Dunn's multiple comparison test. * p < .05, and ** p < .01.

Table 2. *ABCB1* and *ABCG2* mRNA levels following LPS, PolyI:C or ssRNA (10^6 pg/mL) treatment (4- or 24 h), measured using qPCR.

	ABCB1		ABCG2	
	4 h	24 h	4 h	24 h
Ctrl	1.00 \pm 0.05	1.00 \pm 0.05	1.00 \pm 0.22	1.00 \pm 0.12
LPS	0.88 \pm 0.04	0.77 \pm 0.02*	1.03 \pm 0.04	0.55 \pm 0.16*
Ctrl	1.00 \pm 0.21	1.00 \pm 0.11	1.00 \pm 0.17	1.00 \pm 0.12
PolyI:C	1.01 \pm 0.12	1.10 \pm 0.27	1.16 \pm 0.10	0.91 \pm 0.21
Ctrl	1.00 \pm 0.12	1.00 \pm 0.24	1.00 \pm 0.02	1.00 \pm 0.11
ssRNA	1.51 \pm 0.38	1.96 \pm 0.13**	1.45 \pm 0.19*	0.98 \pm 0.08

Statistical analysis: unpaired Student's *t*-test. N = 4–6/group. Mean \pm SEM. * p < 0.05 and ** p < 0.01 versus respective controls.

BCRP levels in the plasma membrane of human BEC cells following treatment [Figure 4](#). LPS had no effect on plasma membrane P-gp at 4 or 24 h; however, it reduced plasma membrane BCRP at 24 h post-treatment (p < .05). PolyI:C treatment increased P-gp levels in the plasma membrane at 4 h of exposure (p < .001), with no changes after 24 h. PolyI:C did not affect BCRP levels in the plasma membrane. ssRNA had no effect on P-gp in the plasma membrane but increased BCRP 4 h after treatment (p < .05).

Cytokine regulation of P-gp/ABCB1 and BCRP/ABCG2 mRNA expression and function P-gp and BCRP function and *ABCB1* and *ABCG2* mRNA levels were assessed in BECs following 4 h and 24 h exposure to key infection-associated pro-inflammatory cytokines, IL-6, TNF- α and IFN- γ [Figure 5](#) and [Table 3](#).

Exposure to IL-6 inhibited P-gp activity after 24 h at all concentrations investigated (10^{1-4} pg/mL; p < .05), though no effects were observed at 4 h. Similarly, IL-6 (10^3 pg/mL) down-regulated *ABCB1* mRNA levels (p < .001) 24 h after treatment. IL-6 also decreased BCRP function (10^3 and 10^4 pg/mL; p < .01) and *ABCG2* mRNA levels (10^3 pg/mL, p < .05) 24 h after treatment [Figure 5d](#) and [Table 3](#). There were no effects of IL-6 on BCRP function or *ABCG2* mRNA after 4 h of treatment. There were also no effects of TNF- α treatment on P-gp activity or *ABCB1* mRNA levels. However, TNF- α treatment decreased BCRP activity (10^3 and 10^4 pg/mL; p < .01) 4 h after treatment, though there were no effects on *ABCG2* mRNA levels. IFN- γ increased P-gp function (10^3 and 10^4 pg/mL; p < .01), but down-regulated *ABCB1* mRNA levels (p < .05) 24 h after treatment [Figure 5j](#) and [Table 2](#).

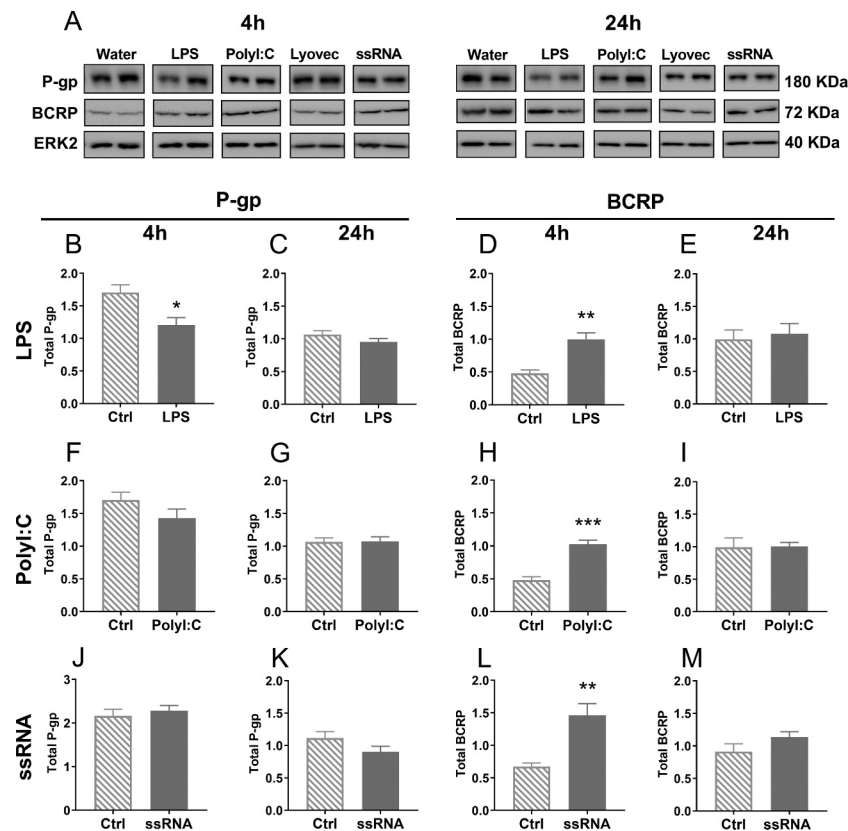


Figure 3. Total P-gp and BCRP protein expression in BECs, 4 h or 24 h after treatment with LPS (b–e), PolyI:C (f–i) or ssRNA (j–m) and their respective controls, measured using Western blot. Representative images of P-gp, BCRP and ERK2 (loading control) bands were cropped and are shown in A. N = 4–5/group. Dose: 10^6 pg/mL. Statistical analysis: unpaired Student’s t-test. Data are presented as mean \pm SEM. * $p < .05$, ** $p < .01$ and *** $p < .001$.

In contrast, IFN- γ decreased BCRP function both 4 h (10^3 pg/mL; $p < .05$) and 24 h (10^3 and 10^4 pg/mL; $p < .01$) after treatment Figure 5(k, L), though there were no effects on ABCG2 mRNA levels.

Discussion

The current study has demonstrated, for the first time, that bacterial and viral infection-mimics can impact drug transporter activity in human brain endothelial cells *in vitro*. hCMEC/D3 cells exposed to TLR ligands and pro-inflammatory cytokines exhibit altered P-gp and BCRP activity and expression. Interestingly, we also showed that these transporters were regulated in opposite directions by all infection mimics, which may indicate a compensatory regulation between P-gp and BCRP function in human BECs. These findings may have considerable implications for biodistribution of drugs in the CNS during bacterial and viral infection or inflammation.

LPS, a TLR-4 agonist, caused transporter-specific time-dependant changes in function. LPS induced a long-term (24 h) increase of P-gp activity, which, *in vivo*, would likely be associated with reduced concentrations of P-gp substrates in the brain, as shown previously using brain capillaries.^{20,21} Conversely, LPS exposure (24 h) decreased BCRP activity. Considering that altered drug transporter function can profoundly impact the accumulation and efficacy of CNS-targeting drugs, findings in the present study may be extremely important in the context of CNS infection-related therapeutics.

Chikungunya (CHIKV), Dengue virus (DENV), Human Immunodeficiency virus (HIV), Japanese Encephalitis (JEV), SARS-CoV-2, Zika virus (ZIKV) and West Nile virus (WNV) all belong to the class of ssRNA viruses.^{22,23} In the present study, we have shown that the viral-mimic ssRNA inhibits P-gp function both in the short (4 h) and longer-term (24 h). This may indicate potential

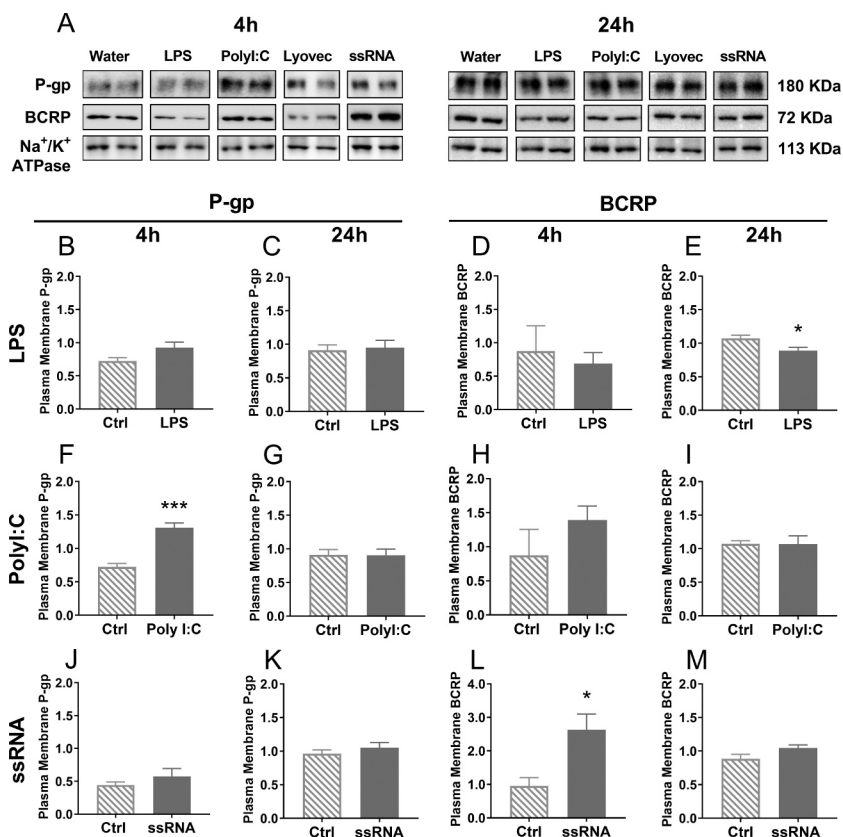


Figure 4. P-gp and BCRP protein expression in the plasma membrane 4 h or 24 h after treatment with LPS (b-e), PolyI:C (f-i) or ssRNA (j-m) and their respective controls, measured using Western blot. Representative images of P-gp, BCRP and Na⁺/K⁺-ATPase (loading control) bands were cropped and are shown in A. N = 4–5/group. Dose: 10⁶ pg/mL. Statistical analysis: unpaired Student's t-test. Data are presented as mean ± SEM. *p < .05 and ***p < .001.

implications in drug transfer across the BBB *in vivo*, in infections induced by ssRNA viruses. To our surprise, BCRP function was increased by ssRNA exposure at 24 h. Further studies are clearly required to determine the impact of ssRNA exposure on drug and xenobiotics accumulation in animal and human brain.

Cytomegalovirus (CMV), DENV and ZIKV activate the host' immune system through TLR-3 stimulation,^{24–26} therefore we used the TLR-3 agonist PolyI:C as an alternative viral mimic. We demonstrated that drug transporter function is regulated in a TLR type-specific manner, since ssRNA and PolyI:C induced opposite effects on P-gp and BCRP function. Human endometrial endothelial cells also present distinct responses to TLR-3 and TLR-7 agonists, based on the distinct inflammatory and antiviral immune pathways that they activate.²⁷ Nevertheless, infections mimicked by PolyI:C may profoundly influence the accumulation of potentially toxic compounds by reducing BCRP activity.

Taken together, our results provide evidence that there may be viral-specific effects on human brain protection through modification of P-gp and BCRP function in the brain vascular endothelium.

All infection-mimics had opposite effects on P-gp and BCRP function. LPS and PolyI:C stimulated P-gp function, but inhibited BCRP activity. In contrast, ssRNA decreased P-gp function whilst increasing BCRP activity. P-gp and BCRP can modulate the transfer of cytokines and other inflammatory mediators from the brain parenchyma to the blood, suggesting that infection itself might alter the accumulation of these compounds in the brain. While P-gp and BCRP share common inhibitors and substrates, their functional relationship is not well understood. In this regard, *Abcb1a*^{-/-} mutant mice exhibit threefold higher levels of *Abcg2* mRNA in brain capillaries compared to wild type animals,²⁸ suggesting a BCRP compensation for the loss of P-gp. Results from the present study, provide the first evidence of a potential compensatory relationship between P-gp

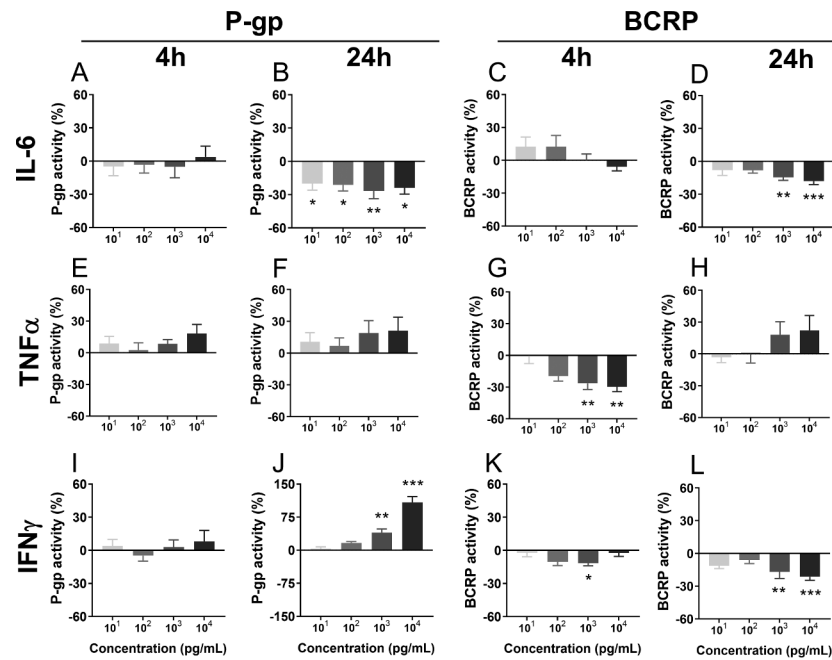


Figure 5. P-gp and BCRP function 4 h or 24 h after treatment with the pro-inflammatory factors IL-6 (a-d), TNF- α (e-h) or IFN- γ (i-l), measured using the Ca-AM and Ce6 assays. Dose: 10^1 – 10^4 pg/mL. N = 6–8/group. Values are expressed as a percentage of controls (vehicle). Data are presented as mean \pm SEM. Statistical analysis: Kruskal-Wallis test, followed by Dunn's multiple comparison test. * $p < .05$, ** $p < .01$ and *** $p < .001$.

Table 3. ABCB1 and ABCG2 mRNA levels following IL-6, TNF- α or IFN- γ (10^3 pg/mL) treatment (4- or 24 h), measured using qPCR.

	ABCB1		ABCG2	
	4 h	24 h	4 h	24 h
Ctrl	1.00 \pm 0.13	1.00 \pm 0.06	1.00 \pm 0.44	1.00 \pm 0.04
IL-6	1.17 \pm 0.06	0.63 \pm 0.05***	1.40 \pm 0.55	0.77 \pm 0.04*
Ctrl	1.00 \pm 0.13	1.00 \pm 0.06	1.00 \pm 0.22	1.00 \pm 0.04
TNF- α	1.29 \pm 0.12	0.85 \pm 0.06	1.26 \pm 0.09	1.09 \pm 0.05
Ctrl	1.00 \pm 0.13	1.00 \pm 0.06	1.00 \pm 0.22	1.00 \pm 0.04
IFN- γ	0.95 \pm 0.09	0.54 \pm 0.08*	1.00 \pm 0.07	1.05 \pm 0.15

Statistical analysis: unpaired Student's *t*-test. N = 4–5/group. Mean \pm SEM. * $p < 0.05$ and *** $p < 0.001$ versus respective controls.

and BCRP at the level of transporter function in human BECs. Further studies are required to better understand this important functional relationship at the level of brain protection.

Another interesting finding of this study is that modification of transporter function in BECs by bacterial and viral mimics seems to occur independent of total protein or mRNA abundance. Studies have reported several post-translational mechanisms regulating P-gp and BCRP activity in BECs, with no changes in gene or total protein expression. They include the insertion/removal of P-gp into/from the cell surface by Caveolin-1, ^{29,30} ERM (ezrin, radixin, and moesin) proteins, ³¹ or small GTPases. ³² In addition, phosphorylation ^{33,34} or glycosylation ³⁵

can also promote changes in drug transporter function, independent of changes in absolute transporter levels.

There were no observable changes in plasma membrane P-gp associated with altered P-gp function in human BECs after infection. In this case, other regulatory mechanisms such as changes in the lipid microenvironment or protein-membrane composition ^{36,37} may be contributing to the observed changes in P-gp function. Furthermore, indirect regulation of drug transporter activity mediated by the actin filament-associated protein (AFAP)-1 has been recently shown to inhibit P-gp function without changing total cellular or plasma membrane levels of P-gp protein in BECs. ³⁸ Further studies are required to elucidate which signaling

pathways are involved in the regulation of *P*-gp activity following infection.

In the case of BCRP, LPS (bacterial infection mimic) decreased transporter activity and plasma membrane BCRP protein at 24 h while total cellular BCRP levels remained unchanged, suggesting a possible change in BCRP intracellular translocation induced by LPS. Similar regulation of BCRP has been previously reported in human embryonic stem cells³⁹ and vascular endothelial cells.⁴⁰

Pro-inflammatory cytokines, including IL-6, TNF- α , and IFN- γ , represents a critical component of the response to bacterial and viral infection.⁴¹ We found in the present study that exposure to IL-6 decreased *P*-gp function and *ABCB1* mRNA levels in BECs, consistent with previous findings in BECs derived from guinea pigs at various stages of development.⁶ IL-6 (24 h) also down-regulated BCRP function and *ABCG2* mRNA levels in human BECs *in vitro*, which was consistent with a previous study that utilized 72 h exposure in the same cell line (72 h).⁴² TNF- α is a critical mediator of viral neurotrophic infections⁴³ and in the current study, there were no effects of TNF- α on *P*-gp function or *ABCB1* mRNA levels. However, BCRP activity was acutely (4 h) reduced after TNF- α treatment, consistent with previous reports in human, porcine, and guinea pig BECs.^{42,44}

IFN- γ plays a central role in viral replication and pathogenesis such as the ZIKV,⁴⁵ and disrupts the BBB, inducing the migration of inflammatory cells into the brain.⁴⁶ Therefore, we hypothesized IFN- γ would impact BBB function by downregulating drug transporter activity in human BECs. Indeed, IFN- γ exposure resulted in a significant reduction in BCRP function at 4 h and 24 h. In contrast, we found increased *P*-gp activity 24 h after exposure to IFN- γ . IFN- γ mediates BBB disruption following JEV infection in mice.⁴⁷ Thus, we can speculate that increased *P*-gp activity after exposure to this cytokine represents an adaptive mechanism in response to increased BBB permeability. Further studies are needed to address this possibility in human BECs. Notwithstanding, this study has identified the potential of IFN- γ to modulate both *P*-gp and BCRP at the human BBB, *in vitro*.

Although the current study focused on drug transporter activity and expression, infection/inflammation also changes other aspects of the

BBB structure. Indeed, tight junctions (TJ), represent important modulators of BBB integrity following exposure to infective/inflammatory stimuli. Studies have demonstrated that the presence of Claudin-5, and its ratio to other claudins, are crucial to normal BBB function.^{48–50} BBB breakdown induced by downregulation/degradation of TJ is a common feature of virus-induced neuropathogenesis. *In vivo* and *in vitro* studies have reported increased BBB permeability with concomitant downregulation of TJ proteins following different models of viral encephalitis, as well as exposure to infective agents, infection mimics, and pro-inflammatory mediators.^{51–61} Clearly, further studies are required to investigate the relationship of TJ and drug transporter function during infection.

Although the calcein-AM activity assay is a commonly used method for *in vitro* assessment of *P*-gp function and that calcein has been used as *P*-gp substrate for several years by different research groups,^{7,62–69} some studies report that calcein could also be substrate for other ABC transporters, including MRP1 and MRP2.^{70–72} There is no evidence in the literature of *ABCC1*/MRP1 or *ABCC2*/MRP2 responsiveness (altered expression or function) following infection in human BBB. However, considering hCMEC/D3 plasma membrane expression levels of MRP1 (1.65 ± 0.23 fmol/ μ g protein) compared to *P*-gp (3.87 ± 0.39 fmol/ μ g protein),⁶⁹ it is possible that the response we found may be driven by a combination of *P*-gp and MRP1 efflux. We recognize this is a potential limitation and highlight the importance of future studies to dissect the specific roles of *P*-gp and MRP1 in brain protection during infection.

In conclusion, we have shown, for the first time, that bacterial and viral mimics influence the expression and function of major drug transporter systems in hCMEC/D3 cells. The current findings are important in the context of bacterial/viral CNS insult, that include *Streptococcus pneumoniae*, *Haemophilus influenzae*, *Neisseria meningitidis*, arboviruses, coronavirus, enterovirus, herpesvirus, poliovirus and rotavirus.^{26,73–75} These pathogens can induce the onset of meningitis/encephalitis, leading to severe cerebral injury and long-lasting neurological dysfunction. We provide *in vitro* evidence that stimulation of TLRs is associated with disrupted *P*-gp and BCRP function in human

BECs. Should a similar effect occur *in vivo* brain protection against xenobiotics would be compromised and there altered biodistribution of drugs in the CNS could result. Further studies with co-culture models would be extremely relevant to highlight the possible influence of astrocytes and microglia on brain endothelial drug transporter response to TLRs ligands. Also, given the transporter-specific sensitivity of P-gp and BCRP in BECs to different infection-mimics, polymicrobial infections may result in diverse patterns of P-gp and BCRP dysfunction in the brain endothelium, and this concept should be further investigated. Our results highlight the need to assess the effects of infection-mimics, and of important human pathogens on the exposure of xenobiotics and accumulation of therapeutic drugs at multiple times across the life course, *in vivo*.

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Disclosure of interest

The authors report no conflict of interest.

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