Shear Stress Regulates ABCA1-dependent Membrane Cholesterol Content in Endothelial Cells Facilitating H$_2$S-dependent Vasodilation

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Abstract:

   Endothelial cells (ECs) express an array of integral membrane proteins, including ion channels and transporters that contribute to blood flow regulation and cell-cell communication. Many of these membrane proteins are regulated by plasma membrane cholesterol content. The ATP-binding cassette family a1 (ABCA1) transporter is a regulator of membrane cholesterol content. We have shown increased ABCA1 mRNA expression and reduced EC membrane cholesterol in resistance mesenteric arteries compared to conduit arteries. Previous studies suggest shear stress (SS) can increase or decrease ABCA1 expression in a cell-type-dependent manner. *Hypothesis*: SS sustains lower EC membrane cholesterol concentration through ABCA1-mediated cholesterol transport, facilitating H_2S-mediated vasodilation. *Method*s: The effect of SS on ABCA1 and membrane cholesterol content was assessed in pressurized mesenteric arteries from male Sprague-Dawley rats and cultured human aortic endothelial cells. Pressure myography was used to assess the effects of ABCA1 inhibition on H_2S-mediated vasodilation. Filipin was used to assess EC membrane cholesterol content. *Results*: SS increased ABCA1 expression in the endothelium of mesenteric arteries and cultured human aortic endothelial cells and markedly reduced EC membrane cholesterol. Inhibition of ABCA1 increased EC membrane cholesterol content and abolished H_2S-induced vasodilation. *Conclusion:* SS facilitation of EC-dependent vasodilation appears to be mediated by membrane cholesterol content.

Introduction:

The endothelium comprises a monolayer of highly specialized endothelial cells (ECs) lining the luminal surface of all blood vessels and occupies a fundamental position in the cardiovascular system. The endothelium regulates complex intercellular interactions, including vascular tone, vascular permeability, and innate immune response[1,2]. The endothelium senses and integrates various stimuli to regulate vascular tone through the synthesis and release of vasoactive gas molecules, such as nitric oxide, carbon monoxide, and hydrogen sulfide (H_2S) [3].

H_2S has been reported to be an endothelial-derived hyperpolarization factor contributing to vasodilation[4] and exerting antioxidant and anti-inflammatory protective effects on the cardiovascular system[5]. The enzymatic synthesis of H_2S involves cystathionine-γ-lyase, cystathionine-β-synthase, and 3-mercaptopyruvate sulfurtransferase; the specific enzymatic pathways vary across vascular beds, adding complexity to the regulation of H_2S-mediated vasodilation[6,7]. Our previous work has shown that H_2S sensitivity in conduit arteries (>300 µm) is dependent on the content of membrane cholesterol in ECs, whereas resistance arteries (<150 µm) are inherently sensitive to H_2S[8].

Membrane cholesterol is a fundamental constituent of the lipid bilayer, impacting membrane fluidity, stability, and organization[9]. Interestingly, our lab has shown that native EC membrane cholesterol content is greater in EC from conduit arteries than in those from resistance arteries,[8] suggesting innate regional differences in membrane cholesterol content along the vascular tree. In addition, our single-cell RNAseq analysis supported by immunohistochemistry indicates that ECs from resistance arteries express greater

levels of the reverse cholesterol transport genes ATP-binding cassette protein A1

(ABCA1) and phospholipid transfer protein (PLTP) compared to ECs from conduit

vessels[10]. The ATP-binding cassette (ABC) transport proteins, a family of eight genes,

are responsible for transporting various hydrophobic compounds across membranes.

ABCA1 specifically facilitates membrane cholesterol efflux through a process called

reverse cholesterol transport in multiple cell types[11,12]. Taken together, these findings

suggest that reduced membrane cholesterol within resistance arteries enhances

endothelial-mediated vasodilation, supporting the postulate that membrane cholesterol is

an important mediator of EC function. However, despite the observed regional

heterogeneity in ABCA1 expression within the vascular tree, it is unknown whether

ABCA1 mediates innate differences in EC cholesterol content. Additionally, the

mechanism mediating the differential ABCA1 expression between conduit and resistance

arteries also remains unknown. Previous studies suggest shear stress (SS) can both

increase and decrease ABCA1 expression, potentially in a cell-type-dependent manner[13–

15].

Vascular SS is a mechanical force resulting from friction against the endothelium

generated by blood flowing along its surface. Vascular ECs sense SS through the

activation of mechanosensors, translating it into biochemical signals influencing cell

morphology, vascular tone, and gene expression[16,17]. As the luminal diameter of an artery

decreases, SS exponentially increases[18], suggesting that differences in SS may mediate

cellular heterogeneity along the vascular tree. Therefore, we hypothesized that SS

sustains lower EC membrane cholesterol concentration through ABCA1-mediated

cholesterol transport, facilitating H_2S-mediated vasodilation.

Methods

Cell Culture

Primary HAoECs (Cell Applications S304-05a, P4-P6) were used for all cell culture

experiments. Cells were plated on cell culture dishes coated with 0.1% Fibronectin

(Sigma, FC010) and maintained at 37°C, 5% CO_2 in endothelial growth media V2 (Cell

Applications). To expose cells to SS, ECs were seeded on μ-Slide 0.4 Luer ibiTreat (ibidi

80176). Unidirectional, recirculating, laminar flow was applied to confluent monolayers

using a peristaltic pump [6 dyne/cm^2, 20 dyne/cm^2, or 0 dyne/cm^2 for static control] with

an inline wind Kessel (to dampen pulsatile oscillations) and bubble trap.

Animals

Sprague-Dawley rats (Envigo, 250–300 g, 9–10 wk. old) were used for all

experiments. Rats were euthanized with a lethal concentration of pentobarbital sodium

(200 mg/kg I.P.), and mesenteric arteries were collected for experiments. The Institutional

Animal Care and Use Committee of the University of New Mexico School of Medicine

reviewed and approved all animal protocols. All protocols conformed to National Institutes

of Health guidelines for animal use.

Preparation of Endothelial Tubes

The mesenteric arcade was removed and immediately placed in ice-cold HEPES buffered

physiological saline solution (HPSS; in mM: 130 NaCl, 4 KCl, 1.2 $MgSO_4$, 4 $NaHCO_3$, 1.8

$CaCl_2$, 10 HEPES, 1.18 KH_2PO_4, and 6 glucose; pH adjusted to 7.4 with NaOH),

containing 10 µM sodium nitroprusside and 0.1% bovine serum albumin (BSA). Veins

were removed, and arteries were cleared of connective and adipose tissue. Arteries (50-

150µm) were flushed of blood and pretreated with probucol (100 µM in 2% pluronic acid)

or vehicle at 37°C for 1 hour in HPSS containing 10% FBS and 0.1% bovine serum

albumin. Endothelial tubes were isolated as previously described[19]. Briefly, arteries were

then placed in an HPSS digestion solution (10% FBS, 0.1% bovine serum albumin, 0.62

mg/mL of papain, 1.5 mg/ mL of collagenase II, and 1 mg/mL dithiothreitol) with continued

treatment with probucol or vehicle. The arteries were incubated in the digestion solution

for 33 minutes at 37°C. The digestion solution was replaced with ice-cold HEPES buffered

physiological saline solution (HPSS; in mM: 130 NaCl, 4 KCl, 1.2 $MgSO_4$, 4 $NaHCO_3$, 1.8

$CaCl_2$, 10 HEPES, 1.18 KH_2PO_4, and 6 glucose; pH adjusted to 7.4 with NaOH), 10%

FBS and 0.1% BSA and artery segments were gently triturated to disrupt the adventitia

and smooth muscle layers. Isolated EC tubes are then transferred using a transfer pipette

and allowed to adhere to a glass slide. A pap-pen was used to create a hydrophobic

pocket on the slide with sufficient surface tension to hold the EC tubes, antibody, and

other solutions within a confined area. A Hamilton 250µL glass syringe was used to apply

staining solutions and perform washes. EC tubes were mounted with ProLong Gold

antifade mounting reagent.

Quantification of Membrane Cholesterol Content

Isolated endothelial tubes were plated on glass coverslips and fixed with 4%

paraformaldehyde (20 min at RT). Endothelial cell membrane cholesterol was detected

by incubating cells with the fluorescent cholesterol marker filipin III (100 µg/mL) for 45

minutes at 37 °C under light-protected conditions. Coverslips were then mounted on

slides using Prolong Gold Antifade (Thermo Fisher P10144). Samples were imaged using

a fluorescent confocal microscope (Zeiss LSM 800 AxioObserver.Z1/7; Gottingen,

Germany) with a 353 nm laser (excitation) and a 465 nm long-pass filter (emission),

combined with a Plan-Apochromat 63/1.4 numerical aperture (NA) oil immersion

objective. A negative control lacking filipin was used to control for background signal.

Filipin staining was quantified using ImageJ[20]. A total of 4 tubes/artery from five animals

were analyzed. The mean fluorescence intensity of each endothelial tube was measured

and averaged to determine a mean fluorescence value for each animal.

HAoECs were stained as previously described above with filipin, and the samples

were imaged using a Zeiss LSM800. Images contained 8-10 images with 7-10 cells per

image for three independent experiments. Regions of interest were created at

approximately 1 mm from the membrane, both intra- and extracellularly, to isolate the

membrane from the intracellular compartments. Images were collected within one optical

section showing membrane max diameter to ensure the focus was on the cell center, and

the mean fluorescence intensity per cell was measured and averaged to determine

cholesterol content.

Quantification of ABCA1 Protein Expression

Endothelial ABCA1 expression was quantified using immunofluorescence in *en-*

face small arteries and HAoECs. Small (<150 µm) were cannulated and pressurized[21].

Briefly, pressurized mesenteric arteries were fixed with 4% paraformaldehyde for 20

minutes at room temperature. Samples were blocked with 5% non-donkey serum (NDS)

and treated with ABCA1 primary antibody (1:100) (Thermo Fisher, PA1-16789) overnight

at 4ºC. Endothelial tubes were then blocked with 5% NDS for 1 h at 37ºC and incubated

with donkey-α-rabbit 488 (1:1000) (Jackson ImmunoResearch Labs, 711-546-152) in 1%

NDS for 4 h at 37ºC. Samples were mounted with Prolong Gold Antifade DAPI (Thermo

Fisher, P36935), and z-stack images were acquired using confocal microscopy (Zeiss

LSM 800 AxioObserver.Z1/7) using a 488 nm laser (excitation) and 517 nm long pass

filter (emission) and a Plan-Apochromat 63/1.4 NA oil immersion objective. Secondary

antibody controls were used to verify the lack of non-specific staining. A total of 4

tubes/animal from five animals were analyzed. Z-stacks containing 14-16 slices each at

150nm were obtained, maximum projections were generated, and the mean fluorescence

intensity (MFI) of each endothelial tube was measured using ImageJ and averaged to

determine a mean fluorescence value for each animal.

For the HAoECs, ECs were exposed to shear stress (6 dynes/cm^2 or 20 dyne/cm^2)

or static conditions for 24 hours using a parallel flow chamber prior to fixation with 4%

PFA. HAoECs were blocked with 5% NDS and treated with ABCA1 primary antibody

(1:100) (Thermo Fisher, PA1-16789) or CSE (1:100 Proteintech, 12217-1-AP) overnight

at 4 ºC. HAoECs were then blocked with 5% NDS for 1 hour at 37 ºC and incubated with

donkey-α-rabbit 488 (1:1000) in 1% NDS for 4 hours at 37 ºC. Samples were mounted

with Prolong Gold Antifade DAPI. Images contained 8-10 images with 7-10 cells per

image for three independent runs, and the MFI was determined using ImageJ. The LXR

agonist T0901317 (100 nM) is a well-established inducer of ABCA1 expression and was

used to validate the specificity of the ABCA1 antibody (Fig. 6).

Application of SS to Mesenteric Arteries

Mesenteric conduit arteries (>300 µm) were cannulated and pressurized to

65mmHg using a gravity-based column system. Luminal flow was generated by adjusting

the column heights [$P_{arterial}$ ~130mmHg and P_{venous} ~35mmHg (HSS)]. Arteries were

continuously superfused with HPSS (5 ml/min, 37°C). Mean pressure was maintained at

65 mmHg. Vessels were exposed to flow for 6 hours, followed by a viability response

(Ach-dilation), before fixation with 4% PFA for 20 minutes while being continually

pressurized. Vessels were then cut along their longitudinal axis, secured *en face,* and

blocked with 5% NDS for 1 hour at room temperature. Vessel segments were then

incubated overnight at 4 °C with ABCA1 primary antibody (1:100 – PA1-16789, Thermo)

in 1% NDS in phosphate-buffered saline. Secondary antibody AF-488 (1:1000, Jackson

Lab) was applied for 1 hour at room temperature in 1% NDS in phosphate-buffered saline

and mounted with Prolong Gold. Five images from various parts of the artery were

captured per condition. ABCA1 intensity was measured using ImageJ.

RT-qPCR Studies

HAoECs were exposed to either static, 6 dynes/cm², or 20 dynes/cm² conditions

for 24 hours. In a separate set of experiments, HAoECs were treated with methyl-β-

cyclodextrin (MβCD, 1mM) to deplete membrane cholesterol or MβCD + cholesterol

(MβCD-Chol, 20mM) to increase membrane cholesterol content for 24 hours. RNA was

isolated using an RNA extraction kit (Macherey-Nagel, 740955.50) according to the

manufacture's recommendations. cDNA was generated with 0.1 µg of RNA using the

High-Capacity cDNA Reverse Transcription Kit (Thermo, 4374967). qPCR was performed

using the TaqMan Fast Advanced Master Mix (Thermo 4444556). TaqMan Gene

Expression Assay Primers for ABCA1 (4331182) and GAPDH (4331182) were purchased

from Thermofisher. GAPDH was used as a housekeeping gene to normalize the

expression of ABCA1. Data was normalized using the 2-delta CT method.

Vasodilation Studies

The mesenteric arcade was removed and immediately placed in ice-cold HPSS.

After the removal of veins and adipose tissue, third- to fourth-order mesenteric arteries

(with an inner diameter of 100–200 μm) were isolated and cannulated with glass

micropipettes, and gently flushed to remove blood from the lumen, as previously

described[22,23]. The artery was stretched longitudinally to approximate its in situ length and

pressurized with a servo-controlled peristaltic pump (Living Systems) to 75 mmHg. The

vessel chamber was superfused with HPSS at 5 ml/min at 37°C and allowed to equilibrate

for 15 min. To determine the contribution of ABCA1 on the H_2S-mediated vasodilatory

response, arteries were pretreated luminally with the specific ABCA1 antagonist, probucol

(100 μM), for one hour before arteries were pre-constricted to ~30–50% with the

thromboxane A_2 analog, U-46619 (Cayman Chemical). To assess H_2S-mediated

vasodilation, arteries were treated with sodium hydrosulfide (NaHS) at concentrations of

1, 10, and 30 μM (Cayman Chemical). Images were obtained using an Eclipse TS100

microscope (Nikon) and CCD100M camera to measure ID using edge-detection software

(IonOptix). Passive diameter was determined at the end of each experiment by

superfusing the vessel in Ca^{2+}-free HPSS HPSS containing: 130 NaCl, 4 KCl, 1.2 $MgSO_4$,

3.6 EGTA, $NaHCO_3$, 10 HEPES, 1.18 KH_2PO_4, and 6 glucose; pH adjusted to 7.4 with

NaOH), containing 10 μM sodium nitroprusside and 0.1% bovine serum albumin (in mM)

for 30 min. For all experiments, changes in vessel diameter (plateau response) are

expressed as the percent reversal of U46619-induced vasoconstriction.

Cholesterol efflux

240   HAoECs were cultured in the presence of probucol (100 µM) or vehicle for

24 hours. The media was then used to assess cholesterol efflux using Amplex-Red,

according to the manufacturer's instructions (Thermo Fisher A12216). A subset of

experiments were also exposed to static or HSS conditions for 24 hrs.

Statistics and Calculations

All data are expressed as means ± SD. Statistical comparisons were made using

Prism 9 (GraphPad Software). The specific statistical tests used are reported in the figure

legends. A probability of <0.05 was accepted as statistically significant for all

comparisons. Percent dilation was calculated as (NaHS diameter – U46619 diameter) /

(Calcium Free diameter – U46619 diameter) x100%.

Results

Effect of SS on ABCA1 expression and membrane cholesterol in HAoECs

We exposed HAoECs to a SS of 6 dynes/cm^2, 20 dynes/cm^2, or static conditions

for 24 hrs. Application of SS increased protein ABCA1 expression in HAoEC (Fig. 1a,b).

SS reduced Abca1 mRNA levels in HAoEC (Fig. 1c). Depletion of membrane cholesterol

reduced ABCA1 mRNA levels. Conversely, supplementing membrane cholesterol

increased ABCA1 mRNA levels (Fig. 1d). To validate that changes in ABCA1 were

associated with reductions in membrane cholesterol, we employed filipin staining to

demonstrate that exposing cells to SS reduces the cholesterol content of the EC

membrane (Fig. 2).

Effect of SS on ABCA1 expression in mesenteric conduit arteries

To assess the effect of SS on ABCA1 expression, pressurized mesenteric arteries were

exposed to either 12 or 20 dynes/cm^2 for 6 hours before fixation. SS elicited an increase

in ABCA1 expression compared to arteries exposed to 12 dyn/cm^2 SS (Fig.3). We have

previously examined ABCA1 expression in freshly isolated paraformaldehyde-fixed

endothelial cell tubes from both large and small mesenteric arteries.[24] ABCA1 expression

and membrane cholesterol were significantly lower in tubes from large arteries compared

to small resistance arteries. ABCA1 expression appears greater in freshly isolated EC

compared to arteries exposed to SS ex vivo. The difference in expression between the

freshly isolated EC and arteries exposed to 6 hours of ex vivo flow may be due to

differences in the absolute magnitude of shear stress in the two settings.

Effect of ABCA1 inhibition on membrane cholesterol and H$_2$S-mediated

vasodilation in mesenteric resistance arteries

Inhibition of EC ABCA1 in small mesenteric arteries (<150 μm) increased EC

membrane cholesterol and reduced cholesterol efflux from HAoECs (Fig. 4a-c).   In

addition, inhibition of ABCA1 significantly attenuated the SS-induced increase in

cholesterol efflux (Fig 4d). Administration of increasing concentrations of H$_2$S elicited a

dose-dependent vasodilation in arteries treated with the vehicle. Inhibition of ABCA1

attenuated H_2S-mediated vasodilation compared to control (Fig. 5).

Discussion

Endothelial function is crucial for maintaining vascular homeostasis, and

disruptions in endothelial signaling pathways can lead to the development of

cardiovascular disease. This study demonstrates a novel role for ABCA1 in regulating

membrane cholesterol levels in endothelial cells. To our knowledge, this is the first study

to demonstrate a role for ABCA1 in SS-mediated reductions in membrane cholesterol.

The findings suggest that ABCA1 regulation of membrane cholesterol impacts hydrogen

sulfide (H_2S)-mediated vasodilation with potential implications for the regulation of

microvascular function and disease pathophysiology. The key findings of this study are

(1) SS increased ABCA1 expression in endothelial cells but reduced mRNA transcripts,

(2) inhibition of ABCA1 increased membrane cholesterol levels and reduced cholesterol

efflux, and (3) ABCA1 inhibition attenuated H_2S-mediated vasodilation.

The regulation of membrane cholesterol by ABCA1 in resistance arteries has

significant physiological implications. Resistance-sized arteries are critical regulators of

vascular resistance and blood pressure, and their endothelial function is highly influenced

by membrane cholesterol content[25,26]. Our previous work demonstrated that five up-

regulated, differentially expressed genes involved in cholesterol transport

(Nfkbia, Apoe, Lrp1, Abca1, Pltp) were present in ECs from small mesenteric arteries

compared to ECs from large arteries. NF-kappa-B inhibitor alpha (*NFKBIA*) has been

shown to enhance cholesterol transport in macrophages by inhibiting NFKB,[27,28], whereas

ApoE facilitates cholesterol transport through direct binding with ABCA1[29,30]. The low-

density lipoprotein receptor-related protein (*LRP1*) has been shown to regulate lipid and

glucose metabolism and is a known co-activator of a transcriptional regulator of ABCA1,

PPAR gamma[31,32,33]. Moreover, phospholipid transfer protein (*PLTP*) functions as an

auxiliary protein that facilitates ABCA1-mediated cholesterol transport[34]. This gene

ontology pathway analysis related to cholesterol transport supports the postulate that

ABCA1 is a key determinant of membrane cholesterol heterogeneity.

Our previous work has shown that H_2S-mediated vasodilation in rat mesenteric

arteries involves the activation of EC transient receptor potential cation subfamily V

member 4 channel (TRPV4) channels and EC large conductance (BK_{Ca}) Ca^{2+}-activated

potassium channels (Morin et al. 2022). TRPV4 and BK_{ca} contain cholesterol

recognition/interaction amino acid consensus sequences (CRAC/CARC)[35,36]. Indeed,

membrane cholesterol has been shown to negatively regulate TRPV4 and BK_{Ca} channel

activity through direct protein-sterol interactions[35,37]. In addition to channel regulation via

direct protein-sterol interactions, changes in membrane cholesterol may alter signaling

microdomains. We have previously demonstrated that the depletion of membrane

cholesterol is required to demonstrate a role for TPRV4 channels in muscarinic receptor-

mediated vasodilation in rat gracilis arteries (Morin et al., 2022). In addition, we have

demonstrated that depletion of membrane cholesterol with MbCD in arteries with higher

cholesterol content (large mesenteric arteries) enhances H_2S dilation. Conversely,

loading arteries with MBCD-chol in arteries with lower native cholesterol content (small

arteries) abolished H_2S-mediated vasodilation. Interestingly, loading these arteries with

the cholesterol enantiomer, epicholesterol, did not affect H_2S vasodilation in these

arteries. Suggesting that the effect of cholesterol on H_2S vasodilation is due to protein-

sterol interactions rather than changes in physical properties of the membrane (e.g.,

membrane fluidity).[38] Our current findings suggest that increases in membrane

cholesterol through ABCA1 inhibition attenuates H_2S-mediated vasodilation in rat

mesenteric resistance arteries, and cholesterol efflux was reduced in HAoECs. Our

findings suggest that ABCA1 regulates endothelial membrane cholesterol by reducing

cholesterol and potentiating the dilatory effect of H_2S.

SS has been recognized as a key determinant of endothelial phenotype influencing

processes such as nitric oxide production[39], cellular polarization[40], inflammation[41,42], and

permeability[43]. A previous report demonstrated a potential mechanism for enhanced

ABCA1 expression in response to SS. Zhu *et al.* observed increases in key transcription

factors involved in lipid homeostasis and ABCA1 regulation in human umbilical vein

endothelial cells exposed to 2 dynes/cm^2 or 12 dynes/cm^2 SS. In addition, their findings

suggest that SS increases mRNA levels of the CYP27A gene, which is responsible for

oxysterol production. Additionally, mRNA levels of LXRα, an oxysterol-activated

transcriptional regulator of *Abca1,* were increased in response to SS[13]. In contrast, Li *et*

al. demonstrated that SS (40 dynes/cm^2) reduced ABCA1 expression in human brain

endothelial cells[14]. These discrepancies may be due to cell type differences or the

magnitude of the applied shear stress. Our data suggest that increases in SS increase

the expression of ABCA1, indicating that cholesterol transport mechanisms are actively

modulated in response to shear stress. We demonstrated that HAoECs exposed to SS

exhibit reduced membrane cholesterol levels compared to cells under static conditions.

In addition, this decrease in plasma membrane cholesterol was associated with a shift in

the localization of free cholesterol to an endosomal-like state and ABCA1 has been shown

to reside on the cell surface as well as endosomal compartments[44]. Notably, the

application of shear stress reduced ABCA1 mRNA transcript levels despite a significant

increase in ABCA1 protein. The sterol regulatory element-binding protein2 (SREBP2) is

a critical regulator of genes involved in cellular cholesterol homeostasis. Classically, a

decrease in cellular cholesterol leads to the activation and translocation of SREBP2 to

the nucleus, thereby promoting the expression of genes involved in cholesterol uptake

and de novo synthesis and suppressing the expression of cholesterol efflux genes[45]. In

the present study, we observed that SS increased ABCA1 protein expression, which is

associated with decreased membrane cholesterol and Abca1 mRNA levels. In addition,

we demonstrate that depletion of membrane cholesterol with MbCD decreased Abca1

mRNA. In contrast, cholesterol supplementation increased Abca1 mRNA levels (Fig. 1).

Taken together, these results suggest that SS may stabilize ABCA1, promoting

cholesterol efflux, and that the loss of membrane cholesterol results in a downregulates

Abca1 mRNA. One consideration in our cell culture model could be a change in the

degradation rate, such that higher flow rates may protect ABCA1 from calpain-mediated

degradation. Several studies have demonstrated the transcriptional and translation

regulation of ABCA1 in the presence of cholesterol-depleting agents such as MBCD or

Apo-A1. Studies suggest activation of ABCA1 to flux cholesterol from the cell resulted in

stabilization of the ABCA1 complex and reduction in transcripts. Arakawa et al.

demonstrate that apoA-1 binding to ABCA1 during cholesterol efflux protects the protein

from degradation, increasing protein levels, which is associated with a decrease in

ABCA1 transcript levels[46]. Over time, in response to continuous SS, a new set point for

cholesterol homeostasis would be established, balancing cholesterol efflux, uptake, and

synthesis[47].

This study provides evidence that ABCA1 mediates cholesterol transport and

regulates membrane cholesterol levels in endothelial cells, and that reductions in

membrane cholesterol content facilitate ion channel activity, contributing to H_2S signaling

and promoting vasodilation. Taken together, these findings provide compelling evidence

for a novel role for ABCA1 in vascular endothelial function. Our data also show that

ABCA1 is a key mediator of endothelial membrane cholesterol in endothelial cells and

that shear stress influences the expression of ABCA1, thereby regulating membrane

cholesterol levels.

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<u>Disclosures</u>

No conflicts of interest, financial or otherwise, are declared by the authors.

<u>Author contributions</u>

J.R.A., L.V.G.B., N.L.K., and J.S.N. conceived and designed research; J.R.A. performed

experiments; J.R.A. analyzed data; J.R.A., L.V.G.B., N.L.K., and J.S.N. interpreted results

of experiments; J.R.A. and J.S.N. prepared figures; J.R.A. drafted manuscript; J.R.A.,

L.V.G.B., N.L.K., and J.S.N. edited and revised manuscript; J.R.A., L.V.G.B., N.L.K., and

J.S.N. approved final version of manuscript.

Figure 1. Shear stress increases ABCA1 expression. (A) Representative images of

ABCA1 expression (yellow) in response to static, 6 dynes/cm^2, or 20 dynes/cm^2 in human

aortic endothelial cells. (B) Quantification of mean fluorescence intensity for ABCA1

expression. 8-10 images were acquired per condition (n=5/group) with 6 – 7 cells per

image. (C) ABCA1 mRNA expression normalized to GAPDH in response to static

conditions, 6 dynes/cm^2, or 20 dynes/cm^2 in HAoECs. (D) ABCA mRNA in response to

cholesterol manipulation in HAoECs. One-way ANOVA with Tukey's Post-hoc test was

used for statistical analysis.

Figure 2. Shear stress decreases membrane cholesterol. (A) Representative images of

membrane cholesterol expression in response to static, 6 dynes/cm^2, or 20 dynes/cm^2 in

human aortic endothelial cells. (B) Quantification of mean fluorescence intensity for filipin

III staining (orange) for the plasma membrane and (C) internal cholesterol content. 8-10

images per condition (n=3/group) with 6 – 7 cells per image were acquired. One-way

ANOVA with Tukey's Post-hoc test was used for statistical analysis.

Figure 3. Increasing shear stress increases ABCA1 expression. (A) Representative

images of ABCA1 expression in response to increasing shear stress in intact large

mesenteric arteries. (B) Quantification of mean fluorescence intensity expressed as fold

change for ABCA1 expression. Five images per artery were acquired per condition

(n=3/group). Nuclear stain DAPI (magenta), ABCA1 (green). Student's t-test was used for
statistical analysis.

Figure 4. Inhibition of ABCA1 increases membrane cholesterol and reduces cholesterol
efflux. (A) Representative images of membrane cholesterol in response to vehicle or
probucol in endothelial tubes isolated from small mesenteric arteries (< 150 μm). (B)
Quantification of mean fluorescence intensity for filipin III staining (yellow). 4 tubes/artery
from 5 animals. (C) Quantification of cholesterol efflux from HAoECs. One-way Anova
was used for statistical analysis. (D) Quantification of cholesterol efflux from HAoECs in
response to shear stress. Student's t-test or One-Way ANOVA with Tukey's Post-hoc test
was used for statistical analysis. **** P<0.0001, *** P< 0.0007.

Figure 5. Inhibition of ABCA1 attenuates H_2S-mediated vasodilation. (A) Representative
trace of vehicle treated artery, The red arrow indicates pre-constrictor U46619, black
arrows indicate H_2S dosage, and the blue arrow indicates Ca^{2+} free PSS. (B)
representative trace of probucol treated artery. (C) Summary responses of NaHS-
dependent dilation in arteries treated with vehicle (n = 5) or the ABCa1 inhibitor probucol
(100 μM, n = 5). Unpaired students t test was used for statistical analysis.

Figure 6. The liver X receptor agonist (T0901317) increases ABCA1 expression in
HAoECs. (A) Representative images of ABCA1 expression in response to vehicle
(DMSO) or T09 (10 μM and 100 μM). (B) Quantification of mean fluorescence intensity
for ABCA1 expression. 8-10 images per condition (n=3) with 6 – 7 cells per image. One-
way ANOVA with Tukey's Post-hoc test was used for statistical analysis.

Fig 1.

447   Fig 2.

452   Fig 3

Fig 4.

Fig 5.

Fig 6.

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