Stimulatory G-Protein α Subunit Modulates Endothelial Cell Permeability Through Regulation of Plasmalemma Vesicle-Associated Protein

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Endothelial cell leakage occurs in several diseases. Intracellular junctions and transcellular fashion are involved. The definite regulatory mechanism is complicated and not fully elucidated. The alpha subunit of the heterotrimeric G-stimulatory protein (Gsα) mediates receptor-stimulated production of cyclic adenosine monophosphate (cAMP). However, the role of Gsα in the endothelial barrier remains unclear. In this study, mice with knockout of endothelial-specific Gsa (GsαECKO) were generated by crossbreeding Gsafl/o mice with Cdh5-CreERT2 transgenic mice, induced in adult mice by tamoxifen treatment. GsαECKO mice displayed phenotypes of edema, anemia, hypoproteinemia and hyperlipoproteinemia, which indicates impaired microvascular permeability. Mechanistically, Gsα deficiency reduces the level of endothelial plasmalemma vesicle-associated protein (PLVAP). In addition, overexpression of Gsα increased phosphorylation of cAMP response element-binding protein (CREB) as well as the mRNA and protein levels of PLVAP. CREB could bind to the CRE site of PLVAP promoter and regulate its expression. Thus, Gsα might regulate endothelial permeability via cAMP/CREB-mediated PLVAP expression.

Keywords: endothelial permeability, Gsα, CREB, PLVAP, edema

INTRODUCTION

The endothelial barrier plays a crucial role in organ function. Dysregulated endothelial permeability contributes to many pathological progresses and affects the treatment of diseases. Transendothelial fluid sieving is controlled by the vascular barrier and hydrostatic, oncotic forces that drive movement across the endothelium, as described by the Starling equation (Claesson-Welsh et al., 2021; Michel, 1997). Increased endothelial permeability is a prominent feature of many diseases such as asthma,
arthritic, chronic bowel disease, cancer, infections, trauma, ischemic stroke, and other conditions where leakage due to increased endothelial permeability could result in edema, impaired function and morbidity (Middleton et al., 2004; Sandoval and Witt, 2008; Tomita et al., 2021). Intracellular junctions connect adjacent endothelial cells and govern the extravasation of plasma and the exchange of its macromolecular constituent. Some extravasation processes occur in a transcellular fashion. Solutes or cells are taken up by endothelial cells and are transported within a vesicle-like structure from the luminal to the abluminal side under inflammatory conditions (Wettchureck et al., 2019). These vesicle-like structures are usually caveolae and vesicular-vacuolar organelle (VVO) (Dvorak and Feng, 2001). Plasmalemma vesicle-associated protein (PLVAP) is an endothelial cell-specific single-span, type II membrane glycoprotein and forms homodimers in situ (Stan et al., 1999a; Stan, 2004). This membrane protein is involved in the formation of the diaphragms that bridge endothelial fenestrae (Stan et al., 1999b). In some organs, such as the lung, PLVAP is required for the formation of caveolae stomata and transendothelial channels (Simionescu et al., 1983). Homozygous stop mutation of Plvap gene in a newborn cause fetal protein-losing enteropathy (PLE) (Broekaert et al., 2018). Homozygous disruption of the Plvap gene in a mixed background leads to growth retardation and decreased survival. The organs with fenestrated capillaries in PLVAP knockout mice showed diaphragm disappearance of caveolae, transendothelial channels (TEC) and fenestrate causing hypoproteinemia and compromising endothelial barrier function. Endothelial PLVAP specifically germine deletion mice had similar features to global PLVAP knockout mice (Stan et al., 2012). Besides, endothelial cell-specific deletion of PLVAP in adult mice induced by tamoxifen were also observed with the loss of plasma protein from circulation and edema formation in multiple organs. Simultaneously, LPS/endotoxin-induced lung injury increased lung endothelial permeability through PLVAP (Jones et al., 2020). Thus, PLVAP plays an indispensable role in endothelial barrier function. However, the regulation of PLVAP expression remains unclear.

The alpha subunit of the stimulatory G protein (Gsα) is expressed in many cell types and is responsible for receptor-stimulated cyclic adenosine monophosphate (cAMP) generation (Weinstein et al., 2007). Gsα signaling has been revealed to play significant roles in skeletal development, neurite formation, neurobiology of learning and memory, inflammatory reactions, tumorigenesis, and others (Cong et al., 2019; Dimitrov et al., 2019; Kelly et al., 2008; Rao et al., 2016; Sarma et al., 2015). We previously reported that smooth muscle-specific Gsa deletion mice displayed severe intestinal obstruction due to decreased contractility of the intestinal smooth muscle (Qin et al., 2017). Moreover, Gsa deficiency in smooth muscle cells promoted smooth muscle phenotype switching and exaggerated angiotensin II-induced abdominal aortic aneurysm formation in mice (Qin et al., 2019). However, the role of Gsa in regulating endothelial cell homeostasis and permeability remains unknown.

In this study, we explored the function of endothelial Gsa in vivo using endothelial-specific Gsa knockout mice and found that Gsa deficiency caused many phenotypes such as edema and hypoproteinemia.

**MATERIALS AND METHODS**

**Generation of Endothelial-Specific Gsα Knockout Mice**

All mice comprised of a C57BL/6J genetic background. The endothelial-specific Gsa knockout (Gsa<sup>ECKO</sup>) mice were generated as follows. Gsa<sup>flox/flox</sup> mice (Chen et al., 2005) were crossbred with transgenic mice expressing CreERT<sup>2</sup> under the control of the Cdh5 promoter (Wang et al., 2010). Recombination was induced in six-week-old mice by daily intraperitoneal administration of tamoxifen (Sigma-Aldrich, St. Louis, MO, United States) dissolved in ethanol: corn oil (1:10) solution with the dose of 1 mg/d for 5 consecutive days. Gsa<sup>flox/flox</sup> Cre-mice were used as control (CTR) for all experiments. Mice were housed in individually ventilated cages under standard housing conditions (22°C, 12 h light/dark cycle), with ad libitum access to chow and water. All animal procedures were approved by and conducted in accordance with the National Institutes of Health Guidelines and with the approval of the Animal Care and Use Committee of Shandong University (Approval No. DWLL-2018-018).

**Blood Pressure Measurement**

The heart rate (HR), systolic, diastolic and mean blood pressures of conscious adult mice were recorded indirectly and noninvasively using a tail-cuff system (BP-2010E; Softron, Tokyo, Japan). After the mice were placed in a hop pocket, the sensor was positioned at the base of the tail. The pocket was kept in a prewarmed box at 37°C, and blood pressure was measured for 20 min at the same time every day. Animals were acclimated to the system for 7 consecutive days before blood pressure measurement.

**Evans Blue Dye Extravasation Assay.**

CTR and Gsa<sup>ECKO</sup> mice underwent Evans blue dye extravasation assay as described (Sangwung et al., 2017). EB dye was obtained from Solarbio (Beijing, China). Male CTR and Gsa<sup>ECKO</sup> mice were injected intravenously with 20 mg/kg body weight of sterile EB dye. Mustard oil diluted to 5% in mineral oil was applied to the dorsal and ventral surfaces of the ear using a cotton swab, which was repeated after 15 min. Mice were anesthetized with ketamine and xylazine (100 mg/kg+5 mg/kg respectively, i.p.) and photographs were taken 30 min after the injection of EB dye. The whole body was perfused with phosphate buffer saline (PBS), and the lung, heart, kidney, liver, and skin were harvested. The EB dye was extracted from the organs with 1 ml of formamide overnight at 55°C and measured spectrophotometrically at 600 nm. The amount of EB extracted in formamide was calculated against a standard curve of known EB concentrations.

**Transmission Electron Microscopy**

Mice were anesthetized with ketamine and xylazine (100 mg/kg+5 mg/kg respectively, i.p.). Organs were harvested,
fixed in 2.5% glutaraldehyde/4% paraformaldehyde fixative, and then cut in 1 mm² blocks. The blocks were then rinsed with 0.1 M sodium cacodylate and postfixed in 1% OsO₄/0.1 M sodium cacodylate. Then the blocks were briefly rinsed, stained with Kellenberger’s uranyl acetate, dehydrated through graded ethanol and embedded in Epon812 resin (Sigma-Aldrich). Ultrathin sections (20–40 nm) were cut and stained with uranyl acetate and lead citrate. The sections were examined with JEOL-1200EX using a bottom-mount MORADA-G2 camera.

**Blood Sampling and Biochemical Analysis**

The blood was isolated and immediately collected in microcontainer tubes coated with clotting activators containing a gel separator (BD Biosciences, Plymouth, United Kingdom). Blood was allowed to clot for at least 30 min before serum separation by centrifugation at 3,000 rpm for 15 min. The biochemical analysis was performed using the automatic biochemical analyzer (Chemry 240) to measure albumin, alanine aminotransferase (ALT), aspartate aminotransferase (AST), blood urea nitrogen (BUN), creatinine, urine protein, high-density lipoprotein (HDL), low-density lipoprotein (LDL), triglycerides (TC), and total cholesterol (TG) levels. For routine blood tests, the blood was isolated and immediately collected in EDTA-coated microcontainer tubes (BD Biosciences). Blood cell compositions were analyzed using a ProCyte Dx Hematology Analyzer (IDEXX, Westbrook, ME, United States) following the manufacturer’s instructions.

**Histology**

The mouse organs were harvested after euthanasia, fixed in 4% formalin and embedded in paraffin. 5 μm thick sections were cut and stained with hematoxylin/eosin (H&E). Images were acquired using PRANORAMIC SCANII (3D Histech, Budapest, Hungary).

**Western Blot Analysis**

Human umbilical vein endothelial cells (HUVECs) and organ tissues were homogenized in RIPA lysis buffer containing protease and phosphatase inhibitors. Lysates were separated by SDS-PAGE and transferred to PVDF membranes. Target proteins were probed with specific antibodies overnight and then incubated with secondary antibodies conjugated with chemiluminescent molecules. This was followed by detection of chemiluminescent reagents (Millipore, Burlington, MA, United States) using the Bio-Rad ChemDoc MP following the manufacturer’s instructions. The sheared chromatin was incubated with 1 μg rabbit IgG or anti-CREB antibody (Millipore). ChIP-DNA was purified and eluted in 50 μl of elution buffer, and 2 μl immunoprecipitated DNA was subjected to PCR using the following primers: 5'- TTCCTCGCGTCTCAAG TTCAT -3' and 5'- ATTGCATAGGAGGCCGTAC -3'.

**Luciferase Reporter Assay**

The DNA fragment from the human PLVAP promoter extending from −1727bp to −1bp, was cloned into the pG3 Basic vector to generate the wild-type Luc construct. The mutant construct with deletion of the CRE site in the PLVAP promoter was generated. For luciferase assay, the luciferase reporter plasmid was transfected into HEK-293T cells in 24-well plates by using lipofectamine 2000 (Thermo Fisher Scientific, Waltham, MA, United States). The p-RL-TK plasmid carrying the Renilla luciferase under control of the thymidine kinase promoter was co-transfected as an internal control. After 24 h, cells were treated with forskolin and luciferase activity was analyzed using the Dual Luciferase Assay Kit (Beyotime, Nantong, China).

**Cell Culture**

HUVECs were purchased from ScienCell (Santiago, MN, United States; Lot Number 28433), and cultured in endothelial cell medium (ECM, ScienCell) with 1% endothelial cell growth supplement (ECGS, ScienCell) and 5% FBS. The culture surfaces were pre-coated by fibronectin (ScienCell) which was diluted in PBS (1:100). Cells from the fourth to sixth generations were used for the experiments. At 80%–90% confluence, HUVECs were transfected with control, Gsa, or CREB siRNA (GenePharma, Shanghai, China) using Lipofectamine RNAiMAX Transfection Reagent (Thermo Fisher Scientific). According to the manufacturer’s instructions, the culture medium was replaced with Opti-MEM (Gibco) and the RNA-lipid complexes were added to cells followed by incubating for 6 h at 37°C and replacing with normal medium. After 48 h, cells were collected for analysis. For the virus-mediated gene transduction, HUVECs were infected with adenovirus-expressing Gsa or lentivirus-expressing CREB (Vigenbio, Jinan, China) for 48 h followed by analysis. H89 and forskolin were purchased from Abcam.

**RNA Extraction and Quantitative RT-PCR**

Total RNA was extracted from HUVECs using ReliaPrep RNA Cell Miniprep (Promega, Madison, Wisconsin, United States). cDNA was reversed-transcribed using the PrimeScript RT
FIGURE 1 | Edema and impaired postnatal survival in GsαECKO mice. (A) Representative immunofluorescent staining of aortas from CTR and GsαECKO mice to determine Gsα expression. Scale bar, 100 μm. (B) The growth curve for male CTR and GsαECKO mice after tamoxifen injection, n = 5 per group. (C) GsαECKO mice were larger than CTR, transparent gelatin substance in subcutaneous tissue (c) and edema facial ministry (d). (D) Kaplan Meier survival plot of CTR and GsαECKO mice after tamoxifen injection, n = 15 per group. ****p < 0.0001 between 2 indicated groups by Log-rank test. (E) Histology of skin harvested from CTR and GsαECKO mice. Scale bar, 500 μm. (F) GsαECKO mice showed ascites, evidenced by fluid accumulation in the peritoneum of the mice. (G) HR (heart rate) of GsαECKO mice and CTR, n = 5 per group. (H) Systolic pressure (SBP) of CTR and GsαECKO mice n = 10 per group. (I) Mean blood pressure (MBP) of GsαECKO mice and CTR, n = 10 per group. (J) Diastolic blood pressure (DBP) of GsαECKO mice and CTR, n = 10 per group. Data are shown as mean ± SEM. *p < 0.05; **p < 0.01; ***p < 0.001 between 2 indicated groups by 2-tailed Student t-test.
reagent kit with gDNA Eraser (Takara, Otsu, Shiga Prefecture, Japan). PCR was performed using TB Green Premix Ex Taq II (Takara) with the Roche Light Cycler 480II. The following primers were used: PLVAP: forward 5′-CCGGTCTCCTA CACGAACA -3′ and reverse 5′-TGAAGAGCAAAGCAT CGCA -3′. β-actin: forward 5′-CATTGATTTTGCTATCCA GGC-3′ and reverse 5′-CTCCTTAATGTCACGCAGT-3′.

Statistical Analysis
Data were expressed as mean ± SEM and analyzed using GraphPad Prism 9 (GraphPad Software Inc., San Diego, CA, United States). Statistical comparisons of two groups using an unpaired 2-sided Student t-test. For comparisons among more than 2 groups, the One-Way ANOVA and Bonferroni post-tests were used. *p < 0.05 was considered statistically significant.

RESULTS

Endothelial-Specific Deletion of Gsα in Mice Caused Edema and Impaired Postnatal Survival
To determine the biological significance of endothelial Gsα expression in adult mice, Gsα<sup>flox/flox</sup> mice were crossbred with Cdh5-CreERT<sup>2</sup> mice to generate Gsα<sup>flox/+</sup>/Cre + mice, which were further intercrossed to obtain Gsα<sup>flox/flox</sup>/Cre + mice. These mice were induced Cdh5-CreERT<sup>2</sup> activity when 6 weeks old by intraperitoneal injection of tamoxifen for 5 consecutive days to delete Gsα in endothelial cells (referred to as Gsα<sup>ECKO</sup> mice). The littermate Gsα<sup>flox/flox</sup>/Cre-mice with the same dose of tamoxifen were used as controls (CTR). Immunoﬂuorescent assay was performed to detect Gsα expression.

![Figure 2](https://example.com/figure2.png)
**FIGURE 2** | Loss of endothelial Gsα increased microvascular permeability. (A) Representative gross images of CTR and Gsα<sup>ECKO</sup> mice after EB dye injection. The black arrow indicated EB dye exudation in Gsα<sup>ECKO</sup> mice. (B–F) Quantification of EB dye extravasation of heart, liver, lung, skin, and kidney. n = 5 per group. (G,H) Transmission electron micrographs of capillaries from lung and kidney of CTR and Gsα<sup>ECKO</sup> mice. The arrowhead indicated the absence of diaphragms. Scale bar, 50 nm. Data are shown as mean ± SEM. *p < 0.05; **p < 0.01; ***p < 0.001; ****p < 0.0001 between 2 indicated groups by 2-tailed Student t-test.
in the aortic endothelium from CTR and Gsa\textsuperscript{ECKO} mice, which showed that Gsa decreased significantly in the endothelial cells of Gsa\textsuperscript{ECKO} mice compared with CTR (Figure 1A). The average body weight of Gsa\textsuperscript{ECKO} mice increased significantly at 3 weeks after tamoxifen treatment progressing in severity as the mice aged (Figure 1B). At the same time, Gsa\textsuperscript{ECKO} mice showed edema of lower body part primarily and spread to the upper trunk and facial ministry (Figure 1C). Transparent gelatin substance was observed in the subcutaneous tissue (Figure 1C). Staining the ventral skin with H&E, which exhibited distended a layer of subcutaneous tissue (Figure 1E). Gsa\textsuperscript{ECKO} mice also developed prominent pleural effusion (Figure 1F). Strikingly, endothelial Gsa deficiency mice succumbed starting 8 weeks after tamoxifen injection (Figure 1D). The HR and blood pressure of Gsa\textsuperscript{ECKO} mice were lower compared to those of CTR (Figures 1G–J).

Gsa\textsuperscript{ECKO} Mice Increased Microvascular Permeability

The movement of albumin from the vascular to the extravascular compartment was evaluated by using the Evans Blue (EB) dye extravasation assay. EB binds tightly to plasma proteins (especially albumin) and is normally retained in the vascular space, its extravasation indicating protein leakage into the interstitial space. Gsa\textsuperscript{ECKO} mice exhibited increased EB-albumin extravasation in organs with fenestrated endothelial cells especially in the skin. The appearance of Gsa\textsuperscript{ECKO} mice showed deeper blue in the ear, paw, subcutaneous gelatin, and perivascular adipose tissue (PVAT) (Figure 2A). Moreover, EB-albumin extravasation also increased in the lung, heart, kidney and liver (Figures 2B–F). These results suggested increased transendothelial permeability of albumin in Gsa\textsuperscript{ECKO} mice. Next, TEM measurement was performed to explore the ultrastructural differences. The caveola diaphragms of lung capillaries were absent in Gsa\textsuperscript{ECKO} mice, while the neck and bulb diameters of the caveola increased (Figure 2G). However, the inter-endothelial junctions appeared similar in CTR and Gsa\textsuperscript{ECKO} mice. In kidney peritubular capillaries, the electron opaque structure indicative of diaphragms was not found in Gsa\textsuperscript{ECKO} mice (Figure 2H). Thus, endothelial Gsa is necessary for the \textit{in vivo} maintenance of capillary diaphragms and its deficiency could cause gross changes in the caveolar structure.

Gsa\textsuperscript{ECKO} Mice Displayed Inflammation Infiltration in the Lung

Gsa\textsuperscript{ECKO} mice exhibited pleural effusion (Figure 3A), which coagulated into gelatin within a few minutes of being fetched. Pleural fluid contained high levels of protein and glucose, FIGURE 3 | The lung tissue of Gsa\textsuperscript{ECKO} mice was infiltrated with inflammation. (A) Pleural effusion was sucked from the thoracic cavity. (B) Presence of petechiae on the pleural surface of Gsa\textsuperscript{ECKO} mice lungs. (C) H&E staining of lung tissue sections from CTR and Gsa\textsuperscript{ECKO} mice. Scale bar, 20 μm. (D) Representative immunofluorescent staining of CD68 in lung tissues of CTR and Gsa\textsuperscript{ECKO} mice. Scale bar, 100 μm. (E) Quantification of CD68 positive area/DAPI area. n = 6 to 7 per group. Data are shown as mean ± SEM. *p < 0.05 between 2 indicated groups by 2-tailed Student t-test.
suggesting that loss of endothelial Gsα increased filtration of plasma into the extravascular spaces (Table 1). Gross examination revealed numerous petechiae on the pleural surface microvessels, indicative of red blood cells extravasation in the lung of GsαECKO mice (Figure 3B). Vascular abnormalities were not observed in other organs examined such as kidney, liver, spleen, and brain. H&E staining showed inflammatory cell infiltration and interstitial thickening in GsαECKO mice (Figure 3C). The immunofluorescent staining with CD68 in lung tissues indicated that macrophage infiltration increased in GsαECKO mice (Figures 3D, E). Increasing EB-albumin extravasation and fluid extravasation reflected by H&E staining indicated that endothelial Gsα deficiency impaired endothelial barrier function especially in the lung, which led to inflammatory infiltration.

**Loss of Endothelial Gsα Impaired Plasma Protein Homeostasis and Blood Composition**

Since endothelial Gsα deficiency in adult mice increases vascular permeability to albumin, we assessed the function of the liver and kidney. The plasma albumin level in GsαECKO mice was significantly lower than that in CTR (Figure 4A). The serum levels of ALT, AST and BUN were similar in CTR and GsαECKO mice (Figures 4B–D), although plasma creatinine was increased in GsαECKO mice (Figure 4E). GsαECKO mice exhibited normal kidney and liver histology (Supplementary Figure S1). Protein concentrations in the urine of CTR and GsαECKO mice were not significantly different (Figure 4F). These results illustrated that endothelial Gsα deficiency does not injure the kidney and liver, so hypoproteinemia of GsαECKO mice was not caused by abnormal liver or kidney. Routine blood test revealed that white blood cells (WBC) including lymphocytes and granulocytes were increased in GsαECKO mice (Figures

| TABLE 1 | Composition of ascites in GsαECKO mice. Pleural fluid was collected from the thorax of GsαECKO mice and analyzed for biochemical contents; n = 4. |
| Pleural Fluid | Mean | SEM | n |
| Glucose (mM) | 5.4488 | 0.28318 | 4 |
| Albumin (g/L) | 10.2708 | 0.26769 | 4 |
| Total Protein | 15.9173 | 0.62739 | 4 |

FIGURE 4 | Endothelial Gsα deficiency impaired plasma protein homeostasis and disrupted blood composition. (A–E) Quantification of Albumin, ALT, AST, BUN, and Creatinine in blood serum of CTR and GsαECKO mice. n = 5 to 6 per group. (F) Quantification of protein in the urine of CTR and GsαECKO mice. n = 5 to 6 per group. (G–K) Quantification of white blood cells (WBC), Lymphocyte, Granulocyte, Monocyte, and red blood cells (RBC) in the blood of CTR and GsαECKO mice. n = 5 to 6 per group. (L–M) Quantification of hemoglobin (HGB) and (red blood cell-specific volume) HCT in blood of CTR and GsαECKO mice. n = 5 per group. (N–O) Mean corpuscular hemoglobin (MCH) and mean corpuscular volume (MCV) of red blood cells of CTR and GsαECKO mice. n = 5 to 6 per group. Data are shown as mean ± SEM. *p < 0.05; **p < 0.01; ***p < 0.001 between 2 indicated groups by 2-tailed Student t-test.
Monocytes had no difference between CTR and Gsa\(^{ECKO}\) mice (Figure 4J). Compared with CTR, Gsa\(^{ECKO}\) mice also exhibited significantly decreased red blood cells (RBC) and hemoglobin (HGB) levels (Figures 4K,L). Furthermore, Gsa\(^{ECKO}\) mice displayed decreased red blood cell-specific volume (HCT) (Figure 4M), increased mean corpuscular hemoglobin (MCH) and mean corpuscular volume (MCV) (Figures 4N,O). These results indicated that macrocytic anemia may occur in Gsa\(^{ECKO}\) mice.

Knockout of Endothelial-Specific Gsa Mice Displayed Decreased Adipose Tissue

Based on the previous studies, the decreasing plasma protein followed by a certain degree of hypertriglyceridemia in PLVAP\(^{-/-}\) mice (Stan et al., 2012). Serum lipid levels in Gsa\(^{ECKO}\) mice showed increased plasma TG, T-CHO, LDL, and HDL (Figures 5A–D). Although Gsa\(^{ECKO}\) mice gained more average weight than CTR at 3 weeks after tamoxifen injection, adipose tissue in Gsa\(^{ECKO}\) mice decreased. White adipose tissue (WAT) deposits in the abdominal wall, retropitoneal, and gonadal fat pads were much less in Gsa\(^{ECKO}\) mice compared to that in CTR (Figure 5E). The inguinal white adipose tissue (ingWAT) of Gsa\(^{ECKO}\) mice showed decreased size and infiltration with gelatinous material. At the same time, mesenteric white adipose tissue (mWAT) nearly diminished in Gsa\(^{ECKO}\) mice (Figure 5F). The histological analysis of subcutaneous adipose tissue (SAT) and visceral adipose tissue (VAT) revealed no significant difference in adipocytes size between CTR and Gsa\(^{ECKO}\) mice (Figure 5G), suggesting that the production of adipocytes decreased in Gsa\(^{ECKO}\). Taken together, endothelial Gsa knockout mice not only displayed increased blood lipid but also decreased white adipose tissue deposit, which illustrated its role in the regulation of lipid metabolism.

Gs\(\alpha\) Deficiency Reduced Plasmalemma Vesicle-Associated Protein Expression in Endothelial Cells

Gs\(\alpha\)\(^{ECKO}\) mice displayed the phenotype of diaphragms deficiency in the lung and kidney, which was similar to that in endothelial PLVAP knockout mice. PLVAP was identified as the first known molecular component of diaphragms of fenestrated capillaries (Stan, 2004; Stan et al., 2004). Therefore, we investigated whether Gsa regulates PLVAP expression. The protein and mRNA levels of PLVAP were markedly decreased in lung tissue of Gsa\(^{ECKO}\) mice compared with CTR (Figures 6A,B). Then, knockdown of Gsa with siRNA decreased the phosphorylation of CREB and the protein and mRNA levels of PLVAP in HUVECs (Figures 6C,D).

![Figure 5](https://example.com/figure5.png)

**Figure 5** | Gsa\(^{ECKO}\) mice displayed decreased blood lipid and adipose tissue. (A–D) Quantification of TG, TC, LDL and HDL-C in blood serum of CTR and Gsa\(^{ECKO}\) mice. *n* = 5 to 6 per group. (E) Representative gross anatomy images of inguinal white adipose tissue (ingWAT), epididymal white adipose tissue (eWAT), perirenal white adipose tissue (pWAT) and interscapular brown adipose tissue (iBAT) from CTR and Gsa\(^{ECKO}\) mice. (F) Representative gross anatomy images of mesenteric white adipose tissue (mWAT) tissue from CTR and Gsa\(^{ECKO}\) mice. (G) Histology of subcutaneous adipose tissue (SAT) and visceral adipose tissue (VAT) of Gsa\(^{ECKO}\) and CTR mice. Scale bar, 100 \(\mu\)m. Data are shown as mean ± SEM. **\(p<0.01\); ***\(p<0.001\); ****\(p<0.0001\) between 2 indicated groups by 2-tailed Student t-test.
Also, CREB knockdown with siRNA suppressed PLVAP expression (Figures 6E,F). Moreover, H89, as a PKA inhibitor, inhibited CREB phosphorylation and PLVAP expression in HUVECs (Figure 6G). Thus, Gsα deficiency could decrease PLVAP expression in endothelial cells.

Gsα Regulates PLVAP Expression via CREB-Mediated Transcription

Gsα was overexpressed in HUVECs, and the protein and mRNA levels of PLVAP were significantly elevated (Figures 7A,B), which indicated that endothelial Gsα may contribute to the transcription of PLVAP. CREB is the Gsα/cAMP/protein kinase A (PKA)-mediated transcript factor (Zhang et al., 2020) and may regulate PLVAP expression. HUVECs infected with CREB lentivirus displayed increased PLVAP expression (Figure 7C). We also used the cAMP activator forskolin to stimulate the endothelial cells and it enhanced the PLVAP levels (Figure 7D). To determine whether Gsα regulates PLVAP expression via CREB-mediated transcription, the PLVAP promoter was analyzed by the Transcription Factor Database (http://jaspar.genereg.net), an Internet-based transcription-factor binding-site program, one CRE site in the PLVAP promoter was identified. To test whether CREB binds to the predicted CRE site on the PLVAP promoter, chromatin immunoprecipitation assay was performed. The result demonstrated that CREB could bind to the CRE site in the PLVAP promoter (Figure 7E). To further analyze the role of the CRE site in PLVAP promoter activity, we mutated the core CRE site of PLVAP promoter which was inserted into a luciferase plasmid and tested in HEK-293T cells. Forskolin treatment significantly increased the luciferase activity of the wild-type but not CRE mutant PLVAP promoter (Figure 7F). The results indicated that the CRE site of the PLVAP promoter is required for Gsα-induced PLVAP gene expression. Thus, our data demonstrated that Gsα regulates PLVAP expression via CREB-mediated transcription in endothelial cells (Figure 7G).
DISCUSSION

The monolayer of endothelial cells forms a semi-permeable barrier that controls substance exchange. In this study, we demonstrated that endothelial Gsα knockout mice displayed endothelial barrier impairment with increased microvascular albumin permeability and hypoproteinemia, which may contribute to the phenotype of edema and impaired
postnatal survival of Gsa\textsuperscript{ECKO} mice. The lung tissue of Gsa\textsuperscript{ECKO} mice also showed numerous petechiae and inflammatory cell infiltration, accompanied by hydrothorax and ascites. In addition, Gsa\textsuperscript{ECKO} mice exhibited hyperlipidemia and diminished adipose tissue. Furthermore, caveolae diaphragms were absent following Gsa deficiency. Mechanistically, endothelial Gsa regulated PLVAP expression through cAMP/PKA/CREB signaling pathway.

Endothelial cells are highly metabolically active and play a critical role in regulating visuomotor tone, permeability, angiogenesis and both innate and adaptive immunity, which has been recognized in most diseases (Aird, 2007). Endothelial cell heterogeneity refers to phenotypic diversity across the vascular tree. The exchange of substances between blood and underlying tissue takes place primarily in capillaries. When endothelium is exposed to inflammatory cytokines and other factors, vascular permeability increases and plasmid fluid as well as proteins extravasate, which could sustain under chronic inflammation and cancer (McDonald, 2001). Usually, the transportation of fluid and small solutes occurs in paracellular manners, while macromolecules are transferred through caveole VVOs and transendothelial channels. Caveole and fenestrae possess a thin non-membranous stomatal diaphragm that contains the PLVAP protein, which is a key mediator in transcytosis of albumin across endothelial cells, inflammation-induced permeability, and leukocyte migration (Zhao and Zhao, 2020). As shown in this study, endothelial Gsa knockout mice revealed significant similarity to PLVAP knockout mice, which suffered from edema, anemia, and hyperlipoproteinemia although its severity varied with the background and strain of the mice (Jones et al., 2020; Stan et al., 2012).

PLVAP expression could be modulated by several compounds, signaling molecules, and biological processes. Vascular endothelial growth factor (VEGF) could regulate PLVAP expression in a phosphatidylinositol 3-kinase (PI3K)- or p38 mitogen-activated protein kinase (MAPK)-dependent manner (Strickland et al., 2005). PLVAP upregulation by phorbol 12-myristate 13-acetate (PMA) required the activation of Extracellular signal-regulated kinase 1/2 (ERK1/2)-MAPK pathway (Stan et al., 2004). A disintegrin and metalloproteinase domain 10 (ADAM10)/Notch signaling could downregulate PLVAP during the development of glomeruli (Farber et al., 2018). In addition, transforming growth factor-β (TGF-β), inflammatory mediators such as tumor necrosis factor-α (TNF-α), and shear stress could regulate PLVAP expression (Wasserman et al., 2002). Our study revealed that Gsa/cAMP/CREB signaling stimulates PLVAP gene expression, which enhance our understanding of PLVAP regulation.

The endothelial Gsa deficiency or overexpression could affect CREB protein level, which was consistent with previous reports from our and other groups that Gsa deficiency in smooth muscle or cardiomyocytes resulted in CREB protein reduction (Qin et al., 2017; Yin et al., 2021). Phosphorylation at Ser133 in CREB may contribute to its protein stability (Costes et al., 2006; Mouravlev et al., 2007). The molecular mechanism about the regulation of CREB protein level and stability by Gsa needs to be further investigated.

In addition to the function of vascular tone and permeability, endothelial cells also play an important role in the regulation of tissue lipid uptake and metabolism. For example, CD36 deletion in endothelial cells could increase plasma free fatty acids (FFA) and postprandial TG levels (Son et al., 2018) and its deletion in lymphatic endothelial cells caused visceral obesity and insulin resistance (Cifarelli et al., 2021). Endothelial peroxisome proliferator activated receptor-γ (PPARγ) knockout mice exhibited increased serum FFA and TG levels, decreased adiposity and increased insulin sensitivity in responses to high-fat diet (Kanda et al., 2009). Capillary endothelial cells uptake of lipids might occur via caveolae-mediated transcytosis. Cav-1\textsuperscript{−/−} mice had high blood fatty acid (FA) and TG levels, which may be attributed to impaired transcytosis of albumin that carries FAs and other lipids (Schubert et al., 2001). Our results showed that Gsa\textsuperscript{ECKO} mice displayed lower serum albumin levels, which reduced the binding between lipoprotein lipase (LpL) and vascular endothelial cells, resulting in the decreased ability of TG clearance and increased blood lipid (Shearer and Kaysen, 2006). Besides, endothelial Gsa deficiency reduced white adipose deposition. These results further confirm the important role of endothelial cells in the regulation of lipid metabolism.

In conclusion, adult endothelial Gsa deficiency increased microvascular permeability, which contributed to edema, anemia, hypoproteinemia, and hyperlipoproteinemia in Gsa\textsuperscript{ECKO} mice. Mechanically, Gsa regulated PLVAP expression through CREB-mediated transcription. Thus, Gsa plays a vital role in regulating endothelial cell permeability, which may provide a new strategy for the treatment of endothelial permeability-related diseases.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author.

ETHICS STATEMENT

The animal study was reviewed and approved by the Animal Care and Use Committee of Shandong University. Written informed consent was obtained from the owners for the participation of their animals in this study.

AUTHOR CONTRIBUTIONS

LH, WZ, and CZ designed the study. LH, HL, XJ, JC, and XQ performed experiments and analyzed results. LH and WZ drafted the manuscript. MC, LSW, JG, JY, QZ, and CZ revised the
manuscript for important intellectual content. All authors read and approved the final paper.

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**SUPPLEMENTARY MATERIAL**

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fphar.2022.941064/full#supplementary-material
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