

Effects of Ebola Virus Glycoproteins on Endothelial Cell Activation and Barrier Function

Victoria M. Wahl-Jensen,^{1,2†} Tatiana A. Afanasieva,^{3†} Jochen Seebach,³ Ute Ströher,^{1,2}
Heinz Feldmann,^{1,2} and Hans-Joachim Schnittler^{3*}

*Special Pathogens Program, National Microbiology Laboratory, Public Health Agency of Canada,¹ and
Department of Medical Microbiology, University of Manitoba,² Winnipeg, Manitoba, Canada, and
Institut für Physiologie, Technische Universität, Dresden 01307, Germany³*

Received 25 October 2004/Accepted 1 May 2005

Ebola virus causes severe hemorrhagic fever with high mortality rates in humans and nonhuman primates. Vascular instability and dysregulation are disease-decisive symptoms during severe infection. While the transmembrane glycoprotein GP_{1,2} has been shown to cause endothelial cell destruction, the role of the soluble glycoproteins in pathogenesis is largely unknown; however, they are hypothesized to be of biological relevance in terms of target cell activation and/or increase of endothelial permeability. Here we show that virus-like particles (VLPs) consisting of the Ebola virus matrix protein VP40 and GP_{1,2} were able to activate endothelial cells and induce a decrease in barrier function as determined by impedance spectroscopy and hydraulic conductivity measurements. In contrast, the soluble glycoproteins sGP and Δ -peptide did not activate endothelial cells or change the endothelial barrier function. The VLP-induced decrease in barrier function was further enhanced by the cytokine tumor necrosis factor alpha (TNF- α), which is known to induce a long-lasting decrease in endothelial cell barrier function and is hypothesized to play a key role in Ebola virus pathogenesis. Surprisingly, sGP, but not Δ -peptide, induced a recovery of endothelial barrier function following treatment with TNF- α . Our results demonstrate that Ebola virus GP_{1,2} in its particle-associated form mediates endothelial cell activation and a decrease in endothelial cell barrier function. Furthermore, sGP, the major soluble glycoprotein of Ebola virus, seems to possess an anti-inflammatory role by protecting the endothelial cell barrier function.

Ebola virus (EBOV) causes a severe hemorrhagic disease in humans and nonhuman primates, with mortality rates as high as 89% (14, 33). The hemorrhagic disease caused by EBOV is characterized by generalized fluid distribution problems, hypotension, coagulation disorders, and a tendency to bleed, finally resulting in fulminant shock. Vascular instability and dysregulation are hallmarks of the pathogenesis in EBOV hemorrhagic fever (HF). Endothelial disturbances can be caused indirectly, by proinflammatory cytokines such as tumor necrosis factor alpha (TNF- α) released from EBOV-infected monocytes/macrophages, and directly, following virus infection of endothelial cells. Additionally, the EBOV transmembrane and soluble glycoproteins are regarded as major viral pathogenic determinants and are also thought to contribute to vascular dysregulation (8, 14, 16, 38, 47, 52, 63).

All glycoprotein forms are encoded by gene 4 of the EBOV genome. The primary product of this gene (80% of the transcripts) is a precursor of the nonstructural soluble glycoprotein (pre-sGP) that is posttranslationally cleaved by furin or a furin-like endoprotease into mature sGP and Δ -peptide, both of which are secreted products (54, 55). In addition to being observed *in vitro*, sGP has been detected in the blood of infected patients (34). RNA editing is necessary to express the

precursor of the structural type I transmembrane glycoprotein (GP_{1,2}) (34, 50). The precursor is posttranslationally cleaved by furin or a furin-like endoprotease into the disulfide-linked fragments GP₁ and GP₂ (35, 51, 53). The homotrimeric GP_{1,2} forms the spikes on the virus particle and is indispensable for receptor binding and fusion with the host cell membrane (18, 48, 59). *In vitro* studies have demonstrated that the full-length GP_{1,2} is cytotoxic to endothelial cells and thus may contribute to endothelial damage during EBOV HF (8, 52, 63). More recently it was demonstrated that a mutant EBOV lacking the editing site showed increased cytotoxicity, suggesting that editing might be a mechanism to regulate EBOV GP_{1,2} cytotoxicity (52). The active role of sGP has not been sufficiently elucidated. Previously it was suggested that sGP interacts with and inactivates neutrophils through binding to CD16b (23, 62), a concept that has been challenged by others (27). Due to the relatively high expression level of sGP during infection, its role in effectively binding antibodies that might otherwise be protective and/or in functioning as a mediator in the activation of target cells has been discussed (16, 34, 47). However, the functional role of the soluble glycoproteins in vascular dysregulation such as increased endothelial permeability, hemorrhage, and shock remains largely unknown (14, 38).

The vascular endothelium has multiple functions in maintaining homeostasis. It forms a unique selective barrier between the blood and tissue, controlling exchange of solutes and water. Balance of bodily solutes between blood and tissue and fluid homeostasis are regulated in two ways. Transport of macromolecules, such as albumin, through the endothelium fol-

* Corresponding author. Mailing address: Institut für Physiologie, Technische Universität, Medizinische Fakultät, Fetscherstraße 74, Dresden D-01307, Germany. Phone: 49-351-458 6007. Fax: 49-351-458 6301. E-mail: hans.schnittler@mailbox.tu-dresden.de.

† V.M.W.-J. and T.A.A. contributed equally to this work.

lows a transcellular pathway (29). At the same time, water and small-solute exchange occurs by a paracellular route through endothelial junctions preferred in capillaries and postcapillary venules (2, 28). Under physiological conditions the intercellular junctions are impermeable to macromolecules. In inflamed tissue, the endothelial barrier function is decreased due to reduced intercellular adhesion, leading to enhanced extravasation of small solutes and water that causes edema formation (2, 28, 40). In severe cases intercellular gap formation, allowing extravasation of larger macromolecules and water into the interstitial space and further contributing to the severity of the pathological condition, can be observed (28).

The functional integrity of the endothelium is provided by endothelial adherens, tight, and gap junctions (5, 9, 38, 49). The composition of endothelial junctions is heterogeneous within different organs and within the microvascular bed (45). The common distribution of adherens junctions throughout the vascular bed makes them crucial structures for regulating the paracellular barrier function, particularly in small vessels. The postcapillary venules, displaying mainly adherens-type junctions, are the target site for increased paracellular permeability (2, 6, 28) and leukocyte extravasation during inflammation (10, 26, 49) and EBOV infection (13, 14, 36). The backbone of adherens junctions is formed by transmembrane vascular endothelial cadherin (VE-cadherin, also known as cadherin-5). VE-cadherin interacts with β - and γ -catenins, and this complex is connected via α -catenin to the actin filament cytoskeleton, which is crucially involved in the regulation of the endothelial barrier function (5, 9, 37, 57). An earlier study has shown that stimulation with tissue culture supernatants derived from Marburg virus-infected monocytes/macrophages caused a reorganization of the VE-cadherin-catenin complex and was associated with intercellular gap formation (13). This phenomenon is comparable to changes observed under stimulation of endothelial cells with inflammatory mediators such as TNF- α (58). While the molecular mechanisms for the breakdown of endothelial barrier function are not completely understood, it is clear that the VE-cadherin-catenin complex and actin filaments are crucially involved. Increases in endothelial permeability are associated with changes in both VE-cadherin organization and formation of actin filament stress fibers that are able to contract, thereby providing the driving force for intercellular gap formation (5, 37, 57). The vascular endothelium also directs immune responses through the induction of cytokines, chemokines, and cellular receptors that recruit or activate immune cells. Proinflammatory cytokines (e.g., TNF- α , IFN- γ) activate the endothelium, causing expression of cell adhesion molecules (CAMs) (e.g., E-selectin, ICAM-1, VCAM-1). These molecules mediate the initial steps in leukocyte transmigration and mediate rolling and tight adhesion of leukocytes to the endothelium (46, 49, 60).

In the present paper we address the question of whether the soluble EBOV glycoproteins, sGP and Δ -peptide, influence endothelial activation and endothelial barrier function. We found that neither sGP nor Δ -peptide was able to activate endothelial cells or cause significant changes in barrier function. In contrast, virus-like particles (VLPs) expressing the transmembrane glycoprotein GP_{1,2} at the surface and thus mimicking the morphology of virus particles were able to activate endothelial cells and induce a moderate but sustained

decrease in barrier function. Most interestingly, sGP displayed an unexpected protective role in endothelial layer integrity when endothelial cells were treated with TNF- α , suggesting a possible anti-inflammatory role of sGP during EBOV HF.

(This work was performed by V. M. Wahl-Jensen in partial fulfillment of the thesis requirement for a Ph.D. from the University of Manitoba, Winnipeg, Manitoba, Canada, 2004.)

MATERIALS AND METHODS

Cell culture and virus. Endothelial cells were isolated from human umbilical veins and cultured as previously described (40). Cells were passaged once and grown to confluence in endothelial growth medium (Promocell, Heidelberg, Germany). All culture dishes were coated with cross-linked gelatin as described elsewhere (39). The human embryonic kidney cell line 293T and Vero E6 cells (ATCC 1568) were maintained in Dulbecco's modified Eagle medium supplemented with 10% fetal bovine serum, L-glutamine (2 mM), penicillin (100 U/ml), and streptomycin (100 μ g/ml). All cells were incubated at 37°C in a humidified 5% CO₂ environment. *Zaire ebolavirus* (ZEBOV) strain Mayinga was kindly provided by the Special Pathogens Branch, Centers for Disease Control and Prevention, Atlanta, Ga. Virus stocks were freshly prepared in Vero E6 cells using Dulbecco's modified Eagle medium with 2% fetal calf serum. Harvesting was performed at a time when no obvious cytopathic effect was seen, in order to limit contamination with released cellular proteins. Mock-infected Vero E6 cells were treated the same way in order to prepare a control (mock stock). Virus titration was performed as previously described (47).

Infection of endothelial cells. Confluent monolayers of human endothelial cells on coverslips were either mock infected or infected with ZEBOV at a multiplicity of infection of 10 (diluted in endothelial basal medium). After an adsorption period of 1 h, the inoculum was replaced with new medium (endothelial basal medium containing 5% human serum) and the cultures were incubated for various times at 37°C in a CO₂ incubator.

Production and purification of recombinant proteins and VLPs. Influenza virus hemagglutinin epitope-tagged (HA-tagged) recombinant ZEBOV sGP and Δ -peptide were generated through transient transfection and purified as described previously (56). Briefly, soluble glycoproteins were expressed in 293T cells and immunoaffinity purified using the HA tag. The dimeric and monomeric conformations of sGP and Δ -peptide, respectively, were conserved and were similar to those of wild-type proteins produced during virus infection (56). Purified proteins were quantitated using the DC protein assay (Bio-Rad, Mississauga, Ontario, Canada). ZEBOV VLP_{VP40/GP} and VLP_{VP40} were generated as previously described by transient transfection of 293T cells with a plasmid(s) encoding ZEBOV VP40 and/or EBOV GP_{1,2} and quantitated by electron microscopy particle counts and a DC protein assay (56). Prior to use, all recombinant glycoproteins, VLPs, and media were analyzed for endotoxin presence using the *Limulus* amoebocyte lysate test (BioWhittaker, Walkersville, Md.) according to the manufacturer's instructions. The relative molar concentrations of glycoproteins were calculated as follows: sGP, 6×10^{12} molecules per μ g; Δ -peptide, 4.3×10^{13} molecules per μ g; GP_{1,2} in VLP_{VP40/GP}, 9×10^5 molecules per particle.

Treatment of endothelial cells with proteins and VLPs. Recombinant proteins or VLPs were added to the culture medium of confluent endothelial cells at 10- μ g/ml or 50- μ g/ml quantities for proteins or at a 1:1 ratio of VLPs to cells. Cells were incubated at 37°C for 6, 12, or 24 h posttreatment in a 95% humidified, 5% CO₂ environment. Recombinant human TNF- α (100 ng/ml) (R&D Systems, Minneapolis, Minn.) served as a positive control, while HA-peptide (Roche, Laval, Quebec, Canada), used to elute the HA-tagged soluble glycoproteins from the affinity column, and purified supernatants of control vector-transfected cells served as negative controls.

Immunofluorescence assays. For immunofluorescence analysis, polyclonal goat antibodies to ICAM-1 and VCAM-1, mouse monoclonal antibodies to E-selectin and PECAM-1 (all from R&D Systems, Minneapolis, Minn.), and a monoclonal antibody to VE-cadherin (BD Biosciences, Heidelberg, Germany) were used. After the appropriate treatment, endothelial cells were fixed with 2% formaldehyde followed by permeabilization with 0.1% Triton X-100 for 10 min at room temperature. Cells were incubated with primary antibodies at a dilution of 1:100 overnight at 4°C. A corresponding secondary antibody conjugated to Alexa 488 (Molecular Probes, Eugene, Oreg.) or Cy3 (Jackson ImmunoResearch Laboratories, Bar Harbor, Maine) was applied for 1 h at a 1:400 dilution. Actin was visualized by incubating the cells with phalloidin conjugated with tetramethyl rhodamine isocyanate (TRITC; Sigma, Deisenhofen, Germany) for 30 min at

room temperature. Stained cells were postfixed with 2% paraformaldehyde and analyzed using a Zeiss microscope.

RT-PCR. Endothelial cells were treated with VLPs or mock supernatants as described above. At 0, 6, 12, and 24 h, cells were disrupted using the guanidinium isothiocyanate-based RLT buffer of an RNeasy Mini kit (QIAGEN, Hilden, Germany). RNA was isolated according to the protocol for animal cells provided by the manufacturer. cDNA was synthesized using a First Strand cDNA synthesis kit for reverse transcription-PCR (RT-PCR) (Roche, Mannheim, Germany) according to the manufacturer's instructions followed by conventional PCR with the following primers: ICAM-1 fwd, 5'-CAGTGACTGCTACTCGAGATCT-3'; ICAM-1 rev, 5'-CCTCTTGGCTTAGTCATGTGAC-3'; VCAM-1 fwd, 5'-CTGGAG GATGCAGACAGGAAG-3'; VCAM-1 rev, 5'-CCAATCTGAGCAGC AATCCGG-3'; E-sel fwd, 5'-AGT GGCCACGGTGAATGTGTA-3'; E-sel rev, 5'-CCCAGATGAGG TACTGTAAG-3'; GAPDH fwd, 5'-GGTTTCTAG ACGGCAGTCA-3'; GAPDH rev, 5'-TGGCAA TTCCATGGCACCGTC A-3'. The PCR included 94°C for 30 s, 65°C for 30 s, and 72°C for 1 min for 19 to 22 cycles, with a final elongation of 7 min at 72°C.

Impedance spectroscopy. Chambered slides for impedance spectroscopy were prepared by applying photosensitive lacquer (CRC Industries, Iffenzheim, Germany) on indium tin oxide (ITO)-coated polyester film (75 mm by 25 mm by 2 mm) (Delta Technologies Ltd., Stillwater, Minn.) Electrode areas were generated by exposing masked ITO slides to UV light for 3 min. Electrode areas were cleared of lacquer by treatment with a sodium hydroxide solution (0.7%, wt/vol). Chambers removed from LabTek slides (Nunc International, Naperville, Ill.) were fixed to the ITO with silicon, coated with cross-linked gelatin as described above, and seeded with endothelial cells. The transendothelial electrical resistance (TER) was determined as previously described (42). Briefly, an alternating voltage was applied, and the impedance magnitude was measured at frequencies between 10 Hz and 1 MHz between the electrode area of the ITO slide and a counterelectrode. The TER was calculated from the resultant spectra (42). Prior to addition of secreted glycoproteins or VLPs, the cells were allowed to equilibrate for approximately 1 h or until a constant TER spectrum was observed in order to establish a baseline resistance. Only the endothelial layers displaying more than 10 Ω cm² were used for experiments. At the termination of all experiments, a 3 mM solution of EGTA, which induces a decrease in TER, was added to all wells as a control. All electrical resistance data are presented as normalized to baseline resistance values (TER/TER₀). TER data are shown as means \pm standard errors. Data were compared by analysis of variance in conjunction with Bonferroni's adjustment (SigmaStat; SPSS, Erfurt, Germany). Values were considered to be statistically significant when P was <0.05 .

Transwell filter tracer assay and hydraulic conductivity. The transwell filter tracer assays were performed on Costar filters (diameter, 6.5 mm; pore size, 0.4 μ m) (Costar, Corning, N.Y.) by using fluorescein isothiocyanate (FITC)-labeled dextran (Sigma, Deisenhofen, Germany) as the tracer substance as described elsewhere (13). Briefly, endothelial cells were treated with VLPs, and 0.5 μ g/ml of FITC-dextran (4 kDa, neutral charge) was added to the upper compartment. Following an incubation time of 0, 3, 10, or 20 h, the FITC-dextran was measured in the lower compartment of the transwell filter system by fluorescence photometry. All steps were performed at 37°C.

Hydraulic conductivities of VLP-treated endothelial cell monolayers were measured and calculated as described elsewhere (40). Briefly, cells were cultured on commercially available Costar (Corning, N.Y.) polycarbonate filters (pore size, 0.4 μ m; diameter, 6.5 mm). Endothelial cells were treated with VLP_{VP40/GP} or VLP_{VP40}, and after 20 h a hydrostatic pressure of 10 cm of water column was applied to the upper side of the transwell filter system using a glass tube inserted into the filter. Water flux (J_v) was measured for the first 30 min, and hydraulic conductivity (L_p) was calculated according to Starling's equation, $L_p = J_v \cdot A^{-1} \cdot \Delta P^{-1}$, where A indicates the filter area (in cm²) and ΔP the applied hydrostatic pressure (cm of water column). Constant hydrostatic pressure was maintained by medium adjustment to the upper compartment of the filter. Hydraulic conductivity was expressed as cm \cdot s⁻¹ \cdot cm H₂O⁻¹. Measurements were performed in four independent experiments.

RESULTS

Activation of endothelial cells by Ebola virus glycoproteins.

It was recently reported that supernatants from ZEBOV-infected Vero E6 cells, but not mock-infected supernatants, were capable of activating primary human endothelial monolayers (20). This activation was demonstrated by transcriptional upregulation of the CAMs ICAM-1 and VCAM-1, although ex-

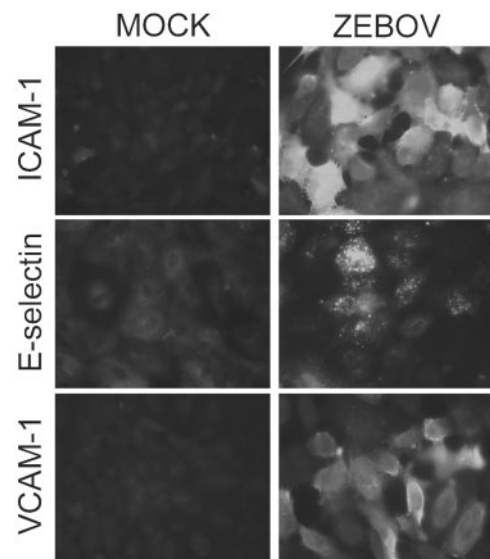


FIG. 1. EBOV-induced expression of CAMs in primary endothelial cells. Mock-infected (controls) and ZEBOV-infected endothelial cells were fixed at 12 h and stained with monoclonal antibodies to E-selectin, ICAM-1, and VCAM-1. Original magnification, $\times 400$.

pression of these and other CAMs on the surfaces of cells was not addressed. Whether this activation could be the result of an interaction between cellular receptors and ZEBOV soluble glycoproteins or particle-associated GP_{1,2}, independent of soluble host factors, remained uninvestigated. Therefore, we first investigated the upregulation of specific cell markers following EBOV infection of endothelial cells. Endothelial cells were infected with ZEBOV at a multiplicity of infection of 10. Expression of ICAM-1, VCAM-1, and E-selectin was demonstrated on the transcriptional level by RT-PCR (data not shown) and on the protein level by indirect immunofluorescence staining 12 h (Fig. 1) and 24 h postinfection. These results confirmed previously published data (20) but additionally demonstrated the expression of CAMs on the surfaces of infected endothelial cells.

To specifically determine the ability of ZEBOV secreted glycoproteins to activate primary endothelial cells, we expressed, characterized, and purified the major secreted glycoproteins, sGP and Δ -peptide. Both proteins were determined to be authentic proteins produced during natural virus infection (56). To assess the ability of particle-associated GP_{1,2} to induce activation, we simultaneously transfected VP40 and/or GP_{1,2} plasmids into 293T cells using a previously established protocol to generate ZEBOV VLP_{VP40/GP} and VLP_{VP40}, the latter of which served as a control (Fig. 2C) (56). Additionally, VLPs were characterized by electron microscopy (data not shown). All preparations were tested for the presence of endotoxin, and levels were found to universally occur below 0.21 endotoxin unit/ml, values equivalent to or less than that of media used in all experiments.

In order to further dissect whether soluble glycoproteins play a role in the upregulation of CAMs and whether replication is needed for the activation, we treated endothelial cells with either VLPs (1:1 ratio with cells) or secreted glycoproteins (10 μ g/ml or 50 μ g/ml) for 6, 12, or 24 h; 100 ng/ml of TNF- α

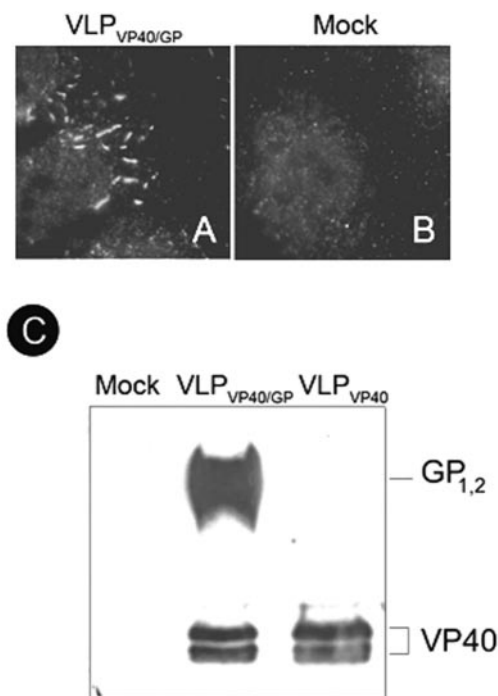


FIG. 2. Adhesion of virus-like particles to endothelial cells. Endothelial cells were treated with purified VLP_{VP40/GP} for 1.5 h, washed extensively, and stained with a monoclonal antibody directed against the transmembrane glycoprotein GP_{1,2} (ZGP12/1.1). Binding of VLP_{VP40/GP} to endothelial cells (A) is shown in comparison to a control (Mock) (B). Original magnification, ×1,000. The different VLP preparations were characterized by Western blotting using anti-GP_{1,2} and anti-VP40 antibodies (C), indicating the absence of GP_{1,2} in the VLP_{VP40}.

was used as a positive control to demonstrate endothelial activation. Like live virus, VLP_{VP40/GP} bound to endothelial cells, as demonstrated by immunofluorescence analysis (Fig. 2A and B), and strongly activated endothelial cells, as measured by upregulation of ICAM-1, VCAM-1, and E-selectin (Fig. 3A and B). Neither sGP nor Δ-peptide caused upregulation of ICAM-1, VCAM-1, or E-selectin over a time period of 24 h at any concentration tested (50 μg/ml and 10 μg/ml) (Fig. 3A, shown for 12 h after treatment with 50 μg/ml of glycoproteins). The time kinetics of endothelial activation were studied on the transcriptional level by RT-PCR using RNA isolated from cells either treated with the VLPs, mock treated, or treated with the soluble glycoproteins. In VLP_{VP40/GP}-treated cells, transcripts for ICAM-1, VCAM-1, and E-selectin appeared early, with maximal levels of expression at 6 h, and declined thereafter (Fig. 3B). In contrast, sGP and Δ-peptide (data not shown) as well as VLP_{VP40} (Fig. 3C, shown for E-selectin expression) did not upregulate the expression of CAMs, confirming the protein expression data shown in Fig. 3A. These results demonstrated that replication is not necessary to induce activation and that activation is induced by VLPs containing GP_{1,2}.

Change of barrier function by Ebola virus glycoproteins. In order to directly assess the abilities of ZEBOV sGP and Δ-peptide as well as VLPs to affect the endothelial cell barrier function, we used impedance spectroscopy to detect changes in TER. Endothelial cells were cultured on electrodes, secreted

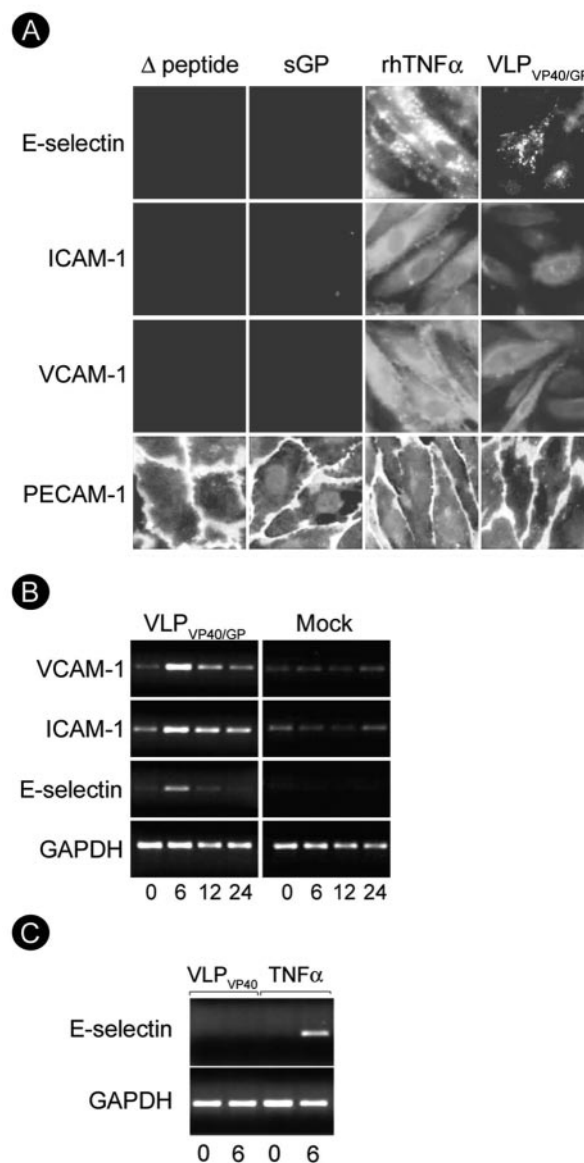


FIG. 3. Endothelial cell activation through soluble glycoproteins and virus-like particles. (A) Endothelial cells were treated with soluble glycoproteins (sGP or Δ-peptide) at a concentration of 50 μg/ml or with VLP_{VP40/GP} (1 particle/cell). Twelve hours posttreatment, cells were fixed and permeabilized, and activation was measured by immunofluorescence analysis using monoclonal antibodies directed against E-selectin, VCAM-1, and ICAM-1. TNF-α was used as a positive control (100 ng/ml). PECAM-1 staining was used to confirm the integrity of the endothelial monolayer. Only TNF-α and VLP_{VP40/GP} induced expression of CAMs. Magnification, ×400. (B and C) The kinetics of VLP-induced endothelial cell activation was further assessed on the transcriptional level by RT-PCR using primers described in Materials and Methods. Cells were either treated with VLP_{VP40/GP} or VLP_{VP40} (1 particle/cell) or mock treated and harvested for RNA isolation at the indicated time points (hours). Transcription of CAMs was clearly upregulated, with maximal levels at 6 h, after VLP_{VP40/GP} (B) but not mock (B) or VLP_{VP40} (C) treatment.

glycoproteins and VLPs were added to the confluent monolayers, and the TER was recorded over a period of 24 h. Even at high concentrations (50 μg/ml), sGP and Δ-peptide were unable to cause long-lasting changes in endothelial barrier func-

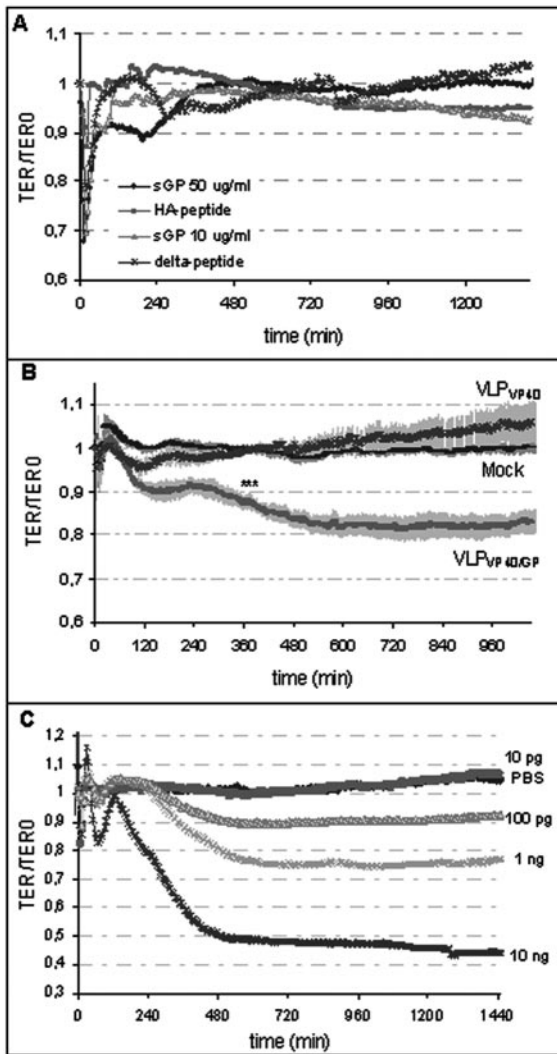


FIG. 4. Effects of soluble glycoproteins, virus-like particles, and TNF- α on endothelial barrier function. Confluent endothelial cell monolayers were treated with soluble glycoproteins (A), VLPs (1 particle/cell) (B), or different doses of TNF- α (C), and the TER was measured using impedance spectroscopy. Treatment of endothelial cells with sGP administered at 10 or 50 $\mu\text{g/ml}$ or with Δ -peptide administered at 50 $\mu\text{g/ml}$ showed no significant long-lasting changes in TER (A). Purified HA-peptide served as a negative control and was added at an equivalent concentration. In contrast to VLP_{VP40}- and mock-treated endothelial cells, treatment with VLP_{VP40/GP} resulted in a decrease in the endothelial barrier function (B). Change in barrier function was significant ($P < 0.05$) from the time points marked by asterisks. In addition, confluent endothelial monolayers were treated with increasing concentrations of TNF- α , and a dose-dependent decrease in TER was observed (C).

tion (Fig. 4A). The rapid transient drop in TER immediately after administration of sGP was concentration independent and therefore seems to be nonspecific. This was followed by a slightly prolonged TER decrease that was not statistically significant. The application of VLP_{VP40/GP} decreased the barrier function within the first 1 to 2 h (Fig. 4B) and was maintained at a 17% decrease for the duration of the experiment. This decrease in TER was found to be statistically significant ($P < 0.05$), and the extent of the decrease was comparable to that

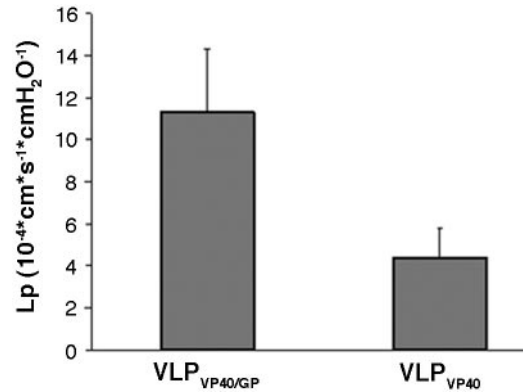


FIG. 5. Effect of virus-like particles on hydraulic conductivity (L_p) of endothelial cells. Confluent endothelial cells cultured on polycarbonate filter membranes were treated with VLP_{VP40/GP} or VLP_{VP40}. In contrast to VLP_{VP40}, VLP_{VP40/GP} increased the hydraulic conductivity approximately threefold.

for endothelial cell treatment with TNF- α at concentrations ranging from 0.1 to 1 ng/ml (Fig. 4C).

To further characterize the permeability-decreasing effect of VLP_{VP40/GP}, we measured endothelial permeability by two additional methods using a transwell filter system. Firstly, we used a 4-kDa FITC-dextran tracer to investigate macromolecular permeability through an endothelial monolayer grown on a filter. We did not observe a significant increase in FITC-dextran permeability following treatment with VLPs (data not shown). However, determination of the hydraulic conductivity clearly indicated a nearly threefold increase in water permeability following treatment of endothelial cells with VLP_{VP40/GP} but not VLP_{VP40} (Fig. 5). The results indicate that VLP_{VP40/GP} is able to increase endothelial permeability for water but not macromolecules and are in accordance with the moderate TER increase demonstrated in Fig. 4.

In order to investigate if the change in barrier function was associated with a redistribution of the adherens-type junction proteins, VE-cadherin, PECAM-1, and the actin filament system were visualized after treatment of endothelial monolayers with VLP_{VP40/GP} (1:1 ratio to cells) or different concentrations (10 $\mu\text{g/ml}$ [data not shown] or 50 $\mu\text{g/ml}$) of the soluble glycoprotein sGP or Δ -peptide for 24 h. Subsequently, cells were fixed and stained for VE-cadherin and actin filaments by using a VE-cadherin monoclonal antibody and TRITC-labeled phalloidin specific for actin filaments, respectively. Neither of the two secreted glycoproteins caused changes in VE-cadherin arrangement or actin reorganization (Fig. 6), effects that are typically seen after treatment of endothelial cells with harsh proinflammatory mediators such as thrombin or high concentrations of TNF- α . VLP_{VP40/GP} treatment did not result in a marked VE-cadherin rearrangement but caused a moderate increase in actin stress fiber formation (Fig. 6). This probably contributes to the moderate changes in barrier function observed by impedance spectroscopy and hydraulic conductivity (Fig. 4 and 5).

Combined effects of viral glycoproteins and TNF- α on endothelial barrier function. During an EBOV infection, the endothelium of the host is targeted by cytokines produced by activated monocytes/macrophages (primary target cells) (14)

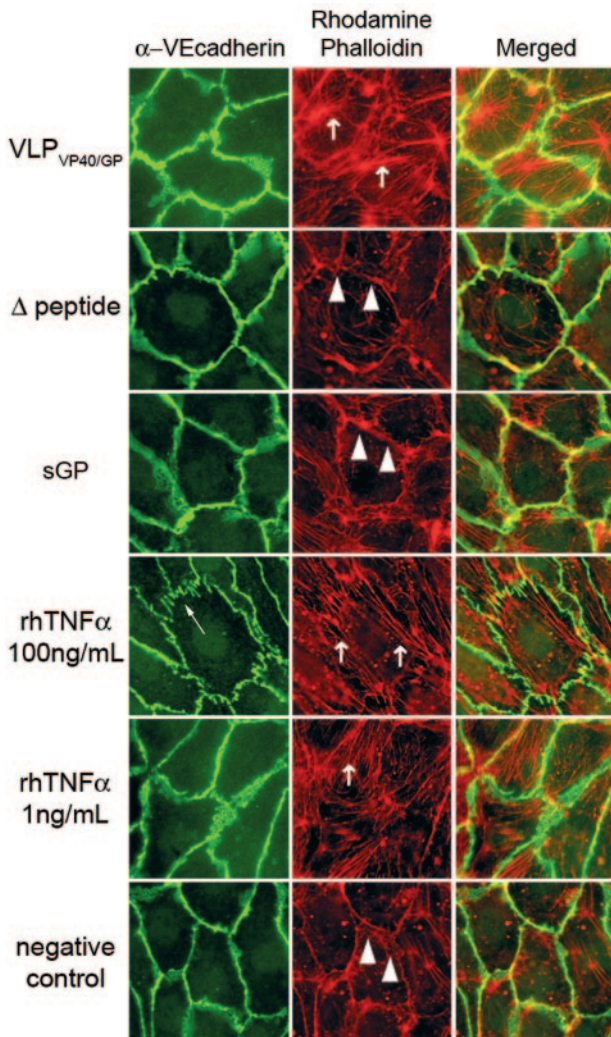


FIG. 6. Effects of soluble glycoproteins and virus-like particles on VE-cadherin and actin distribution in endothelial cells. Confluent endothelial monolayers were treated with either 50 μg/ml sGP or Δ-peptide, VLP_{VP40/GP} (1 particle/cell), or TNF-α (1 ng/ml or 100 ng/ml), which served as a positive control. Only TNF-α at 100 ng/ml was able to cause morphological rearrangement of VE-cadherin and actin stress fibers (arrows). VLP_{VP40/GP} and TNF-α at lower concentrations (1 ng/ml) caused a moderate increase in actin filament stress fiber formation. In control cells the junction-associated actin filaments were seen (arrowheads). Magnification, ×400.

and soluble viral glycoproteins that are released into the bloodstream (34). Thus, it seems likely that the endothelial barrier function is affected as a result of a combined function of all these different cellular and viral mediators. Therefore, we tested the effects of sGP and Δ-peptide on endothelial cell integrity in the presence of TNF-α using impedance spectroscopy. We first determined the amount of TNF-α that produces a moderate but reliable decrease in endothelial cell barrier function (Fig. 4C). For the studies with the soluble glycoproteins, we used 1 ng/ml TNF-α, a dose that reproducibly decreased the barrier function by approximately 30%.

When administered simultaneously, VLP_{VP40/GP} and TNF-α had an additive effect on the decrease in endothelial barrier function that was not observed with VLP_{VP40} (Fig. 7A). This

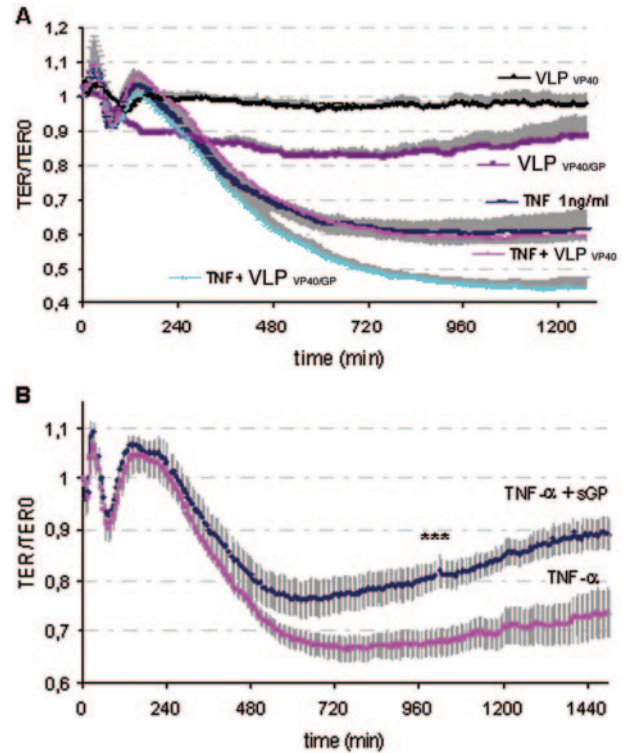


FIG. 7. Effect of simultaneous application of TNF-α and VLPs or sGP on endothelial cells. (A) Endothelial cells were treated with TNF-α (1 ng/ml) and VLPs (1 particle/cell) or sGP (10 μg/ml). Simultaneous application of TNF-α and VLP_{VP40/GP} led to further decreases in TER (approximately 15%), indicating their additive effect on the barrier function. (B) Simultaneous treatment with TNF-α and sGP led to a 15% to 20% recovery of the TNF-α-induced changes in barrier function. Changes in TER were significant ($P < 0.05$) from the time point marked by asterisks.

indicates the importance of particle-associated GP_{1,2} for this process. As demonstrated above (Fig. 4A), sGP did not affect the integrity of the endothelial cell monolayer within 24 h of incubation. However, when administered simultaneously with TNF-α, sGP reversed the barrier-decreasing effect of TNF-α by about 20% and thus allowed for recovery of the endothelial monolayer (Fig. 7B). This effect was statistically significant after 17 h ($P < 0.05$). Furthermore, this effect was sGP specific, since neither VLP_{VP40/GP} (Fig. 7A) nor Δ-peptide (data not shown) had a protective effect on endothelial cell barrier function if applied simultaneously with TNF-α.

DISCUSSION

Of all the viral hemorrhagic fevers, those caused by EBOV are the most severe (14, 33). The dramatic clinical presentation in humans and nonhuman primates as well as extensive laboratory data generated in the past has led to the idea that filovirus-induced disease is as much an immune syndrome as a vascular disease (14, 38). While the clinical picture has become clearer through in vivo experimental studies, the molecular mechanisms, particularly with respect to vascular dysregulation, remain elusive. Although vascular instability and dysregulation are thought to be disease-decisive symptoms, endothelial

cells are largely considered secondary target cells during EBOV infection. Primary replication occurs within monocytes/macrophages and dendritic cells (20, 22, 47), which become activated upon infection and produce active mediator molecules such as proinflammatory cytokines and chemokines but also release soluble viral glycoproteins, particularly sGP, which has been detected in the plasma of human patients (34). The pathogenic role of secreted cytokines in EBOV pathogenesis has been investigated in the past (3, 20, 22, 24, 47), but little is known regarding the role of the secreted glycoproteins in endothelial dysfunction. Nevertheless, it has long been hypothesized that the secreted glycoproteins, particularly sGP and Δ -peptide, may play an important role during EBOV pathogenesis and can function as mediators of endothelial and immune dysregulation (13, 15, 19).

In the past several studies have morphologically described the involvement of the endothelium in experimentally infected nonhuman primates and rodent models as well as in postmortem human material (4, 19, 31, 32, 64). Destruction of endothelial cells could be found only in postmortem human material (64) and some nonhuman primate models (30–32), but all investigators concluded that endothelial cell function was impaired during infection. To investigate the influence of the soluble and transmembrane glycoproteins on endothelial cell activation and barrier function, we used immunofluorescence analyses and RT-PCR as well as impedance spectroscopy and transwell filter systems, respectively. Impedance spectroscopy is a highly sensitive biophysical assay that provides a unique possibility to study the endothelial barrier function under resting (25) and shear stress conditions with high time resolution (11, 42). This technique determines the TER of a cultured endothelial cell monolayer and predominantly reflects the changes in paracellular permeability (42). Endothelial cell barrier function is frequently studied in transwell filter systems by analyzing the passage of tracer substances (10, 12, 13). In contrast to impedance spectroscopy, which reliably detects changes in barrier function of about 2% (42), tracer systems are limited in time resolution and sensitivity and thus allow for the detection of strong effects only.

In the present work we clarified the long-standing question as to whether the EBOV secreted glycoproteins sGP and Δ -peptide are able to influence endothelial cell functions. We determined that, in addition to its ability to induce transcriptional activation of ICAM-1 and VCAM-1 (20), ZEBOV infection of endothelial cells resulted in surface expression of those molecules as well as of E-selectin (Fig. 1). We found that, in contrast to sGP and Δ -peptide, VLP_{VP40/GP}, produced by transfection of VP40 and GP_{1,2}, and nearly identical in morphology to infectious virus particles, activated endothelial cells in culture and decreased the endothelial cell barrier function (Fig. 3, 4, and 5). These effects were not observed using VLPs lacking GP_{1,2} (VLP_{VP40}), indicating the importance of the transmembrane glycoprotein GP_{1,2} in the context of a virion particle for endothelial cell activation (Fig. 3, 4, and 5). The permeability-increasing effect of VLP_{VP40/GP} might allow for extravasation of small solutes and water and thus might contribute to the development of edema and shock observed during EBOV HF. The moderate permeability increase was confirmed by a moderate change in endothelial cell morphology following VLP_{VP40/GP} treatment, as indicated by largely intact

VE-cadherin staining along the cell junctions but formation of actin stress fibers (Fig. 6). A similar morphological change associated with permeability increase was also observed after treatment of endothelial cells with low concentrations of TNF- α (1 ng/ml) (Fig. 4 and 6). The data are in agreement with our earlier studies showing that Marburg virus infection did not enhance macromolecular permeability in a transwell filter system (13).

Geisbert et al. (20) recently demonstrated upregulation of mRNA transcripts of several genes associated with endothelial cell activation, including cyclooxygenase-2, inducible nitric oxide synthase, ICAM-1, and VCAM-1, following ZEBOV infection of endothelial cells. At the same time, gamma-irradiated EBOV almost failed to activate endothelial cells (only cyclooxygenase-2 mRNA transcripts were upregulated), and it was therefore proposed that endothelial cell activation was dependent on virus replication. This raises the question as to whether or not VLPs and virus subjected to gamma irradiation, a method commonly used to inactivate infectious high-containment agents, can be used interchangeably to represent replication-deficient virus particles. It is known that gamma irradiation changes structural features of proteins dramatically, particularly at high doses (17, 21). For example, alpha-crystallin activity was reduced by 40% after a dose of 4,000 Gy, and a dose of 60,000 Gy was used to inactivate EBOV (20). Thus, it seems reasonable to assume that the structure of glycoproteins on gamma-irradiated EBOV may be altered, resulting in insufficient binding of virus particles to endothelial cells and, in turn, insufficient activation, an explanation that still needs to be addressed experimentally.

During an EBOV infection, large amounts of proinflammatory cytokines are secreted from infected primary target cells (20, 22, 47), with the level of TNF- α in the blood of infected patients reaching 5 to 7 ng/ml. Therefore, it is reasonable to assume that the endothelial barrier function during EBOV infection may be affected as a result of a combined action of different cellular and viral mediators. To verify this hypothesis, we studied the effect of the primary soluble glycoprotein sGP on endothelial barrier function in the presence of TNF- α (Fig. 7B). In order to detect potential cumulative effects on endothelial cell barrier function, we used a TNF- α concentration of 1 ng/ml, which induced an approximately 30% decrease in barrier function (Fig. 4C). The moderate effect of VLP_{VP40/GP} on endothelial barrier function was further enhanced in the presence of TNF- α , thus contributing to the severity of endothelial damage even in the absence of direct endothelial cell infection. Surprisingly, when sGP was administered simultaneously with TNF- α , it caused a recovery of endothelial cell barrier function starting after approximately 10 h (Fig. 7B). This finding is intriguing and suggests that sGP may have an anti-inflammatory role in the course of EBOV pathogenesis. During EBOV infection, areas of focal tissue destruction can be seen in multiple organs. Interestingly, these areas largely lack infiltration of leukocytes, although neutrophil aggregation within the vascular system is observed in infected nonhuman primates (31, 32). This might indicate that the activation of the endothelium with the recruitment of neutrophils occurs but that the transmigration process is blocked or impaired, which could be related to the observed anti-inflammatory effect of sGP. TNF- α plays a pivotal role in establishing and orchestrat-

ing inflammatory and immune responses to infection (1, 43). As a consequence, many viruses have evolved countermeasures, such as some of the large DNA viruses (i.e., poxviruses) that encode immunomodulatory proteins which directly inhibit or modify antiviral activities of proinflammatory cytokines (7, 41). Future studies will have to address the mechanism behind the potential anti-inflammatory action of sGP. In particular, it must be clarified whether the effect is specific for TNF- α and, if so, whether sGP directly interferes with TNF- α or components of the TNF signaling pathway. In this study we used a TNF- α concentration (1 ng/ml) lower than that observed in vivo during infection, but the effect of TNF- α on endothelial cells in vivo can be less pronounced. It was shown that under shear stress, which resembles in vivo physiological conditions, endothelial cells are less responsive to TNF- α (44, 61). Thus, it would also be of interest to determine the effect of sGP in vivo.

In conclusion, we demonstrated that neither of the two major EBOV secreted glycoproteins, sGP and Δ -peptide, was responsible for endothelial cell activation and/or reduced endothelial cell barrier function. In contrast, VLPs containing the EBOV transmembrane glycoprotein GP_{1,2} were potent activators of endothelial cells and also induced changes in endothelial cell barrier function, indicating that the transmembrane glycoprotein GP_{1,2} in the context of a virus particle is an important pathogenic determinant. Interestingly, sGP was found to modulate the endothelial cell inflammatory response, a phenomenon that might play a critical role in EBOV pathogenesis.

ACKNOWLEDGMENTS

The excellent technical help of Sylvia Grossklaus and Christine Mund is gratefully acknowledged. We thank Ayato Takada for kindly providing the monoclonal antibody ZGP12/1.1 and Rainer Koch for assistance in statistical evaluations.

This study was supported by grants from the Canadian Institutes of Health Research (CIHR) (MOP-43921, awarded to H.F.), the Priority program of the SPP 1130 "Infection of the endothelium" (DFG-grant SCHN/430 3-1, awarded to H.-J.S.), and the "Med-drive-program" of the Medical Faculty Carl-Gustav Carus, University of Dresden (awarded to J.S.). V.M.W.-J. was supported by fellowships from the Manitoba Health Research Council (MHRC) and the Department of Medical Microbiology, University of Manitoba.

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