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Von Willebrand Factor multimerization and the polarity of secretory pathways in endothelial cells.

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-The three endothelial secretory pathways, constitutive, basal and regulated, release VWF in different multimeric states.
-Apical and basolaterally released VWF follow different secretory pathways, thus releasing differentially multimerized protein.

Abstract

The Von Willebrand factor (VWF) synthesised and secreted by endothelial cells is central to haemostasis and thrombosis, providing a multifunctional adhesive platform that brings together components needed for these processes. VWF secretion can occur from both apical and basolateral sides of endothelial cells, and from constitutive, basal and regulated secretory pathways, the latter two via Weibel-Palade bodies (WPB). Although the amount and structure of VWF is crucial to its function, the extent of VWF release, multimerization and polarity of the three secretory pathways have only been addressed separately and with conflicting results. We set out to clarify these relationships using polarized Human Umbilical Vein Endothelial Cells (HUVECs) grown on Transwell membranes. We found the regulated secretion of Ultra large (UL) molecular weight VWF predominantly occurred apically, consistent with a role in localised platelet capture in the vessel lumen. We found that constitutive secretion of low molecular weight (LMW) VWF is targeted basolaterally, toward the subendothelial matrix, using the adaptor protein complex 1 (AP-1), where it may provide the bulk of collagen bound subendothelial VWF. We also found that basally-secreted VWF is composed of UL-VWF, released continuously from WPBs in the absence of stimuli, and occurs predominantly apically, suggesting this could be the main source of circulating plasma VWF. Together, we provide a unified dataset reporting the amount and multimeric state of VWF secreted from the constitutive, basal and regulated pathways in polarized HUVECs, and have established a new role for AP-1 in the basolateral constitutive secretion of VWF.
Introduction

Von Willebrand factor (VWF) is a multimeric adhesive glycoprotein, synthesised and secreted by endothelial cells\(^1\) that is involved in many vascular processes and is central to haemostasis and thrombosis\(^2\). Circulating plasma and subendothelial VWF, both synthetized by vascular endothelial cells, have been ascribed different roles in haemostasis\(^3,4\). How endothelial cells control these two very different functional pools of VWF has not been previously explored.

The multimeric form of secreted VWF largely dictates its haemostatic efficacy, since higher molecular weight VWF multimers have a much higher binding affinity for platelets\(^5\). Endothelial cells have two differentiated surfaces, apical, facing the vessel lumen, and basolateral facing the subendothelial extracellular matrix. Therefore, to fully understand how and where different multimeric forms of VWF make their contribution, we need to know not only how much is secreted apically versus basolaterally, but also by which one of three pathways it is delivered and the degree of multimerization found in each pool of this protein.

All three secretory pathways begin at the trans-Golgi network (TGN) where VWF is either packaged into nascent Weibel-Palade bodies (WPBs), distinctive rod-shaped regulated secretory organelles, or secreted directly through the ‘constitutive’ secretory pathway by non-descript small anterograde carriers\(^6\). WPBs store VWF for regulated secretion upon endothelial activation but they can also fuse and release VWF in the absence of activation, a process termed ‘basal’ secretion to distinguish it from ‘constitutive’ secretion\(^7,8\), although both collect together in the supernatant of unstimulated cells. Low molecular weight –VWF (LMW-VWF) is secreted by the constitutive pathway, in contrast to the HMW-VWF released via stimulated WPBs\(^6,9\), consistent with the need to control release of highly pro-thrombotic material.

Previous studies have addressed the polarity of VWF secretion, with conflicting results\(^10-14\), partly because they were carried out before it was established that there were three pathways for secretion of VWF\(^8\). van Buul-Wortelboer and colleagues\(^10\) used Human Umbilical Vein Endothelial Cells (HUVECs) grown on collagen lattices to show that unstimulated (i.e. basal plus constitutive) VWF release occurs at a 1:3 ratio (apical:basolateral) although the VWF quantified as basolateral
release also took into account all the VWF cumulatively bound to the collagen fibres (80%), and so is an overestimation of the true basolateral rate of unstimulated release. When cells were stimulated with phorbol 12-myristate 13-acetate (PMA), most of the VWF was released to the apical side, with no detectable change in the amount released basolaterally. In the same year Sporn and colleagues\textsuperscript{11} showed, using HUVECs grown on gelatin-coated polycarbonate membranes (3.0 µm pore size), that the amount of unstimulated VWF release (measured by 2h metabolic labelling and analysed by sodium dodecylsulfate (SDS) –polyacrylamide gel electrophoresis (PAGE)) into apical or basolateral chambers was approximately equal, and contained similar multimeric forms, including multimers of all sizes. When cells were stimulated with the calcium ionophore A23187, 90% of the additional VWF came out basolaterally, with only high multimeric VWF being released. The same pattern was reportedly seen with PMA stimulation.

A later study, again using HUVECs grown on a collagen matrix\textsuperscript{12}, showed that PMA-stimulated release was largely apical, with no change to the levels of basolateral release compared to unstimulated cells. Another study measuring the polar release of a variety of different molecules secreted by endothelial cells, including VWF\textsuperscript{13}, showed that HUVECs grown for 10 days on uncoated polycarbonate membranes (pore size 0.4 µm) secreted unstimulated VWF at a 2:1 ratio (apical:basolateral). No stimulant was used in this study to measure regulated VWF release. One final study\textsuperscript{14} employed the polar Madin-Darby Canine Kidney II (MDCK-II) epithelial cell line stably transfected with VWF cDNA to study VWF polar secretion. Here again, unstimulated VWF release accumulated at a 2:1 ratio (apical:basolateral) compared to a ratio for PMA stimulated release of 4:1. Finally, live cell imaging of individual regulated exocytic events has demonstrated fusion of WPB to both basolateral\textsuperscript{15} and apical surfaces\textsuperscript{16}. While no comparative analysis was done, this provides confirmation that evoked secretion can occur from both domains.

Altogether, the direction of VWF secretion, by which pathway(s) it is released, and the multimeric state of the releasate at both surfaces are currently unclear. In particular, the relative multimeric state and the polarity of VWF secreted by the basal pathway have never been established. We therefore addressed all these issues together and in parallel.
Finally, the origin of plasma VWF, which is the best characterised pool of this molecule and is the pool assayed for diagnostic purposes, is unclear. While it is expected to be generated from highly multimerized material that is then cleaved by ADAMTS-13 down to the multimeric pattern observed in plasma, whether it arises from basal or regulated exocytosis is yet unclear.

Not only is the basic description of VWF secretion confused, but the machinery for polarized targeting of these pathways in endothelial cells is also poorly understood. Since WPB are the regulated secretory VWF carriers, and AP-1 is essential to their formation\textsuperscript{17}, this adaptor protein complex is a strong candidate to be in control of this pathway. Loss of AP-1 also increases unstimulated secretion but whether this is affecting constitutive, basal or both pathways is unclear. Furthermore, AP-1 is thought to mainly play a role in basolateral targeting in polarized epithelial cells\textsuperscript{18} and its role in polar trafficking in endothelial cells has never been explored.

Here, we report the polarity and multimeric state of VWF secretion from constitutive, basal and regulated secretory pathways in polarized HUVECs, establish a new role for AP-1 in the constitutive secretion of VWF, and suggest a likely origin for plasma VWF.
Materials & Methods

Cell Culture

HUVECs pooled from multiple donors were obtained from Lonza (Lonza Walkersville, Walkersville, MD). Cells were maintained in HUVEC Growth Medium (HGM), composed of M199 (Gibco, Life Technologies) supplemented with 20% Fetal Bovine Serum, (Biosera), 30 µg/ml endothelial cell growth supplement from bovine neural tissue and 10 U/ml Heparin (both from Sigma-Aldrich) and used within 4 passages.

Secretion assay and vWF quantification by ELISA

Secreted VWF was obtained by incubating confluent cells in serum-free (SF) medium (M199, 0.1 mg/ml BSA, 10mM HEPES-NaOH, pH 7.4) for 60 minutes. The medium was collected and the cells incubated further with SF medium containing secretagogues for 30 minutes. The remaining cells were then lysed to determine total VWF levels. Relative amounts of VWF were determined by sandwich ELISA as previously described19.

Additional Materials & Methods can be found in Supplementary Information.
Results

Multimeric states of endothelial cell VWF secreted from three different pathways

Since the multimeric state of secreted VWF largely dictates its function, we determined the multimeric pattern of VWF secreted by the three known pathways from endothelial cells on plastic dishes, comparing releasate collected from confluent HUVECs and running the same amount of VWF in each sample (as measured by ELISA) on a sodium dodecylsulfate (SDS) /agarose gel. Brefeldin A (BFA), known to block direct constitutive release from the Golgi was used to distinguish between constitutive and basal VWF release during unstimulated conditions and two different widely used secretagogues, Histamine and phorbol 12-myristate 13-acetate (PMA) were used to stimulate regulated release. As previously reported\textsuperscript{20,21} secretagogue-stimulated secretion, termed ‘regulated’ release, was mainly composed of Ultra-Large (UL) and very high molecular weight multimers (HMW-VWF), while unstimulated release contained a broad range of multimer sizes including a higher proportion of low molecular weight multimers (LMW-VWF) and prominent dimer bands (Figure 1A and B). When BFA was used to block constitutive secretion leaving only basal VWF releasate, a multimeric pattern very similar to that of stimulated samples was observed with the exception that a strong dimer band was always present.

These patterns suggest that although both are released during unstimulated conditions, constitutive and basally secreted VWF are differently multimerized and so could have very different functions. At the same time, while both believed to arise from WPBs, basal and regulated VWF also have different multimeric patterns, namely the presence of a dimer band in basal secretion, also suggesting a significant difference between these two pools of VWF (Figure 1C).

HUVECs form a tight monolayer on Transwell membranes

To study the polarity of VWF secretion, we first established the conditions for filter-grown HUVECs to form a confluent polar monolayer. HUVEC seeded at a
density of $10^5$ cells per cm$^2$ on polyester membranes (pore size 0.4 µm), formed a continuous monolayer within 24 h. These cells are tightly packed, and acquire tight junction and adherent junction markers, such as VE-Cadherin, β-Catenin and ZO-1 (Figure 2A), that are retained for several days post seeding. HUVECs are extremely thin cells (3-4µm at their thickest) (Figure 2Bi), contain overlapping edges (Figure 2Bii-iii) and form tight junctions between neighbouring cells (Figure 2Biv-v). Grown under these conditions, HUVEC monolayers form a barrier to diffusion (Figure 2C). As previously described, treatment of confluent HUVEC monolayers with Thrombin opened gaps between cells$^{22}$ while treatment with 100µM Histamine for 30 minutes did not$^{23}$ (Figure S1 and S3).

**Constitutive, Basal and Regulated VWF polar release**

To measure unstimulated secretion, confluent HUVECs were washed, and incubated in Serum Free medium (SF) for 1 h, and then supernatant collected from the apical and basolateral chambers were assayed for the amount of VWF by ELISA. We found at least twice as much VWF accumulated apically compared to that released basolaterally, and this ratio was maintained cells grown for several days (Figure 3A).

Since VWF secreted with no externally-added stimulant is a mix of constitutive and basal secretory pathways, BFA$^8$ was used to distinguish between these two pathways. Surprisingly, BFA only reduced basolaterally secreted VWF (Figure 3B) revealing that most constitutive VWF secretion is directed towards the basolateral side of endothelial cells. Conversely, VWF released during BFA treatment and thus representing basal secretion, is secreted predominantly apically, with only a small fraction secreted basolaterally.

Since we were not expecting a constitutive VWF carrier to be targeted with any particular polarity, we determined if a synthetic secretory cargo lacking any targeting information would also be targeted basolaterally in this cell-system. To address this, HUVECs were transfected with lumGFP$^{24}$, a synthetic cargo that has been shown to be secreted via the constitutive pathway in endothelial cells$^{25}$. lumGFP secretion from membrane-grown cells was collected from both apical and basolateral chambers and a GFP ELISA showed equal amount of lumGFP released
apically and basolaterally (Figure 3C). Critically, this release was effectively blocked in both directions by BFA. Since lumGFP is secreted in a non-polarized fashion in polarized endothelial cells, this suggests that normal VWF constitutive secretion is dependent on active sorting into basolaterally-targeted carriers.

In order to establish the polarity of VWF secreted via the regulated pathway, cells on membranes were stimulated with various secretagogues. In all cases, more than 70% of secreted VWF accumulated apically (Figure 3D) and this same pattern was unchanged irrespective of how long the cells were grown on filters before assay (Figure 3E). As predicted, BFA did not change the pattern of regulated (i.e. stimulated minus unstimulated) VWF secretion (Figure 3F).

The degree of multimerization seen in VWF being secreted from the different pathways in different directions was then analysed. Equivalent amounts (units) of VWF in each sample (as measured by ELISA) were loaded and analysed by agarose gel electrophoresis. Although our ELISA results predicted that unstimulated apical VWF should be mostly basal secretion, composed of mostly UL-VWF multimers together with a dimer band, we still do see a high proportion of LMW-VWF in this sample, which disappears with BFA (Figure 3G, lane 1 and 3). This would suggest that unstimulated apical VWF releasate is composed of mostly UL multimeric basal VWF but also contains a small amount of constitutive VWF (i.e. BFA sensitive), which, with its LMW-VWF multimers, is probably underestimated in an ELISA and may be overrepresented on a gel.

Our ELISA results also suggest that unstimulated basolateral VWF secretion is mainly composed of constitutive VWF, with a smaller proportion of basal secretion (Figure 3B). This is matched by the multimer gels, where LMW-VWF multimers, which disappear with BFA, are seen along with the basally secreted UL-VWF bands (Figure 3G, lane 2 and 4).

We also noticed that when BFA was used, leaving only basal VWF, a strong dimer band remained in the apical unstimulated releasate but surprisingly disappeared from the basolateral sample (Figure 3G, lane 3 and 4). Further, in the Histamine stimulated samples, along with the expected UL-VWF multimers, a dimer band unpredictably appeared in the apically secreted material (Figure 3G, lane 5).
and 7). We currently have no explanation for this minor but surprising and consistently present phenomena.

**AP-1 and VWF polar targeting**

Most studies investigating the role of AP-1 in polarized targeting have done so by knocking-out or downregulating the amount of the AP-1µ1 subunit, one of the core subunits of AP-1. Most of this work has investigated the role of the epithelial-specific\(^{26}\) AP-1µ1B subunit, and while HUVECs do not express this transcript (**Figure 4A**), they do express the more ubiquitous AP-1µ1A variant, also implicated in polarized trafficking in some studies\(^{18}\). Here we used specific siRNA to effectively down regulate AP-1µ1A in HUVECs (herein referred to as siAP-1M1) using two rounds of nucleofection (**Figure 4B**). Similar to previous results from our lab using a VWF-expressing HEK293 cell model\(^{17}\), downregulation of AP-1µ1A in HUVECs impairs WPB formation (**Figure S2**) with a resulting increase in unstimulated VWF secretion (**Figure 4C**, non-striped bars) plus a decrease in stimulated VWF release (**Figure 4D**), when cells were grown on plastic wells. Critically, this was not due to Golgi fragmentation in siAP-1M1 treated cells\(^ {27,28}\) (**Figure S2**) since we know this would affect WPB biogenesis\(^ {29}\). Using BFA, we can also now show that most of this increase in unstimulated VWF release in siAP-1M1 treated cells is due to an increase in secretion via the constitutive pathway, since a larger proportion was blocked with BFA (**Figure 4C**). These results confirm that AP-1 is necessary for the sorting of VWF into WPB, and suggest that if this is disrupted, then the displaced VWF is continuously secreted through the constitutive pathway.

In order to determine if AP-1 is also involved in the polarized targeting of VWF from the different secretory pathways, during the second round of siRNA nucleofection HUVECs were seeded onto filters, and allowed to grow for two days. Importantly, siAP-1M1 treatment did not alter monolayer permeability even after Histamine stimulation (**Figure S3**). Secreted VWF was collected and assayed by ELISA revealing an increase in the total amount of unstimulated VWF secreted as is seen on plastic-grown cells (**Figure 5A**, total bars). As with cells grown on plastic (**Figure 4C**), this increase reflected a rise in the amount of VWF secreted via the constitutive pathway (**Figure 5B**, larger proportion blocked by BFA treatment).
Critically, as opposed to control cells where constitutive secretion of VWF is mostly basolateral, in siAP-1M1 treated cells it is secreted in both directions, losing its exclusive basolateral targeting. Ablation of another AP-1 subunit (γ-adaptin) confirmed these results (Figure S4), strongly suggesting that AP-1 is involved in the basolateral targeting of constitutive VWF carriers in endothelial cells.

To confirm if this effect on constitutive secretion was specific to VWF, we again used the lumGFP construct to check if AP-1 had any effect on the polarity of its secretion. The amount, sensitivity to BFA and polarity of lumGFP secretion (Figure 5C) were not affected in siAP-1M1 treated cells, confirming a specific effect of AP-1 on the polarity of VWF constitutive carriers and not a general function of this adaptor on all constitutive carriers in endothelial cells. The polarity of basal (Figure 5B, hatched bars) and regulated VWF release (Figure 5D) was not affected in siAP-1M1 treated cells, with the majority of VWF still being released apically.

Multimer gels of VWF secreted from siAP-1M1 treated cells were analysed and, as predicted from the secretion assay results (Figure 5A, B, D), we observed little change in the multimeric state of VWF secreted from siAP-1M1 treated cells compared to control cells. The multimeric pattern from Histamine-stimulated cells was identical and that from unstimulated cells was very similar when compared to control cells (Figure 5E) except for an increase in the proportion of LMW-VWF bands in the siAP-1M1 samples, due to the increase in the proportion of constitutively secreted VWF (Figure 5E, most clearly seen by the increase in tetramers and hexamers in lane 3-4 vs 1-2). This was not unexpected since, in addition to the increased constitutive release, basal and even regulated exocytosis are both still occurring, albeit at reduced levels.
**Discussion**

In this study, we establish for the first time, the relative amounts and multimeric states of VWF released through all three known secretory pathways; constitutive, basal and regulated release. We show that constitutively secreted VWF is composed of mainly LMW-VWF multimers; that basally secreted VWF, originating from the continuous release of VWF from WPB in the absence of added stimuli, is composed of mostly UL-VWF multimers together with a prominent dimer band; and that regulated VWF release is exclusively composed of UL-VWF multimers (Figure 1). Since the discovery of the basal pathway for VWF secretion from endothelial cells\(^8\), the multimeric state of this continuous source of VWF had not been previously established and thus its functional role has never been addressed. Our current findings suggest that the multimeric state of VWF secreted from the three pathways is different and so could have different physiological functions.

We go further and establish the amount and multimeric states of VWF from all three pathways in a polar endothelial context. The need to revisit the polarity of VWF secretion, mainly investigated 25 years ago, reflects firstly the fact that this analysis dates from before the clarification by Giblin et al\(^8\) of the three possible secretory pathways taken by VWF (constitutive, basal and regulated). Secondly, the literature from that time is somewhat confusing, with different results being presented. Finally, the relationship between all three pathways and the multimeric state of VWF has not been previously addressed, let alone in a polarized cell system. We have attempted to put all these aspects together into an overall comprehensive picture of polarized VWF secretion and multimerization using primary human endothelial cells.

We find that the largest amount of stimulated (regulated) VWF secretion occurs apically, with varying total amounts depending on the secretagogues used (Figure 6). This is in line with the majority of previous *in vitro*\(^{10,12,14}\) and *in vivo* studies\(^{30}\). This form of secreted VWF is also composed of mainly UL-VWF multimers. This is in line with the currently accepted, yet experimentally unsupported dogma, that most highly multimerized VWF is secreted mainly towards the apical (lumen) surface of endothelial cells, which we now prove experimentally for the first time. The lumen of the blood vessel is where circulating platelets can be recruited by
locally released UL-VWF upon stimulation, and where endothelial-anchored VWF strings can initiate a haemostatic plug, in a tightly localized and controlled fashion\textsuperscript{31}. Thus, regulated VWF release can be thought of as the critical pool for emergency use only, where apically released VWF strings along with basolaterally released anchoring VWF is restricted to the site of injury, while being of the highest platelet-binding activity.

We also find that twice as much unstimulated VWF accumulated apically compared to that released basolaterally, in line with reports from Narahara et al\textsuperscript{13} and Hop et al\textsuperscript{14}. We go further and use BFA to distinguish between VWF released constitutively versus basally, observing for the first time that basal VWF secretion occurs predominately to the apical side of endothelial cells, with only a small fraction being secreted basolaterally (Figure 6). These findings suggest that this form of VWF, continuously secreted along the vascular tree could be the major source of plasma VWF. Once secreted from endothelial cells through the basal pathway, this UL-VWF form is cleaved by the endothelial secreted\textsuperscript{32} metalloprotease ADAMTS-13\textsuperscript{31}, generating the more intermediate VWF multimers found in plasma (Figure 1), allowing control of levels of highly pro-thrombotic VWF in circulation.

Most in vitro work on exocytosis has focussed on regulated release, and the short collection times (15-60min) used in analyses emphasise the relative amount of secretagogue-driven release, giving the impression of this being the main source of VWF in plasma. However, we suggest that this may be mistaken: if basal release is only 5-20\% of secretagogue-driven secretion (depending on the secretagogue used) per unit time, then for regulated and basal secretion to both generate an equal amount of circulating VWF, 5-20\% of the WPB-containing endothelium would need to be activated at any given time- this seems extremely unlikely.

Our data are also consistent with the need for targeting of VWF into WPB for appropriate multimerization to occur during biosynthesis: both regulated and basal pathways release highly multimerized VWF, in contrast to the constitutive pathway. Why we see a significant dimer band in basally secreted VWF is as yet unclear, although the internal milieu of these two carriers (basal WPBs vs regulated WPBs) are likely to be different, perhaps having different retentions times inside the cell that could potentially drive differential multimerization patterns. Whether the dimers only
reflect the differential maturation of the two populations of WPB, or actively contribute to some difference in function is as yet unknown.

Previous findings from our lab (see Supplementary Figure S4C in 29) and others31,33,34 have shown that unstimulated cells only rarely form VWF strings under flow. This is consistent with our proposed role for basal VWF, again suggesting that, although being of similar high multimeric states (except for the dimer band), basal and regulated VWF play different functional roles. Whether basally-secreted VWF is incapable of producing strings or this difference simply reflects the exocytic mode of this secretory pathway: low numbers of randomly-distributed exocytic events will preclude co-operative action in bringing together the material from more than one exocytic event that could well be needed to drive string formation.

Finally, the requirement to sustain a steady source of plasma VWF to maintain adequate haemostasis is crucial, as observed in Type 1 and Type 3 von Willebrand disease patients35. Basally released VWF provides such a source since it is (1) continuously secreted, (2) of a high multimeric state such that it could act as the precursor of plasma VWF, (3) does not produce VWF strings to initiate thrombus formation, and (4) is secreted mostly apically towards the vessel lumen, four critical requirements for an adequate and reliable source of circulating plasma VWF.

Finally, our results also show that constitutive VWF release is, surprisingly, exclusively basolaterally targeted and We also report a novel role for AP-1 in this delivery along with its known role in WPB biogenesis17. This is in agreement with its long established role in basolateral targeting in epithelial cells. How AP-1 acts in sorting VWF at the TGN is puzzling. The AP-complexes involved in sorting during the formation of clathrin coated vesicles are generally recruited to membranes by binding to the cytoplasmic tails of integral transmembrane proteins36. How then does VWF, a protein with no transmembrane domains, recruit AP-1? To date, there is no identified abundant integral membrane VWF-binding protein to recruit AP-1. We note that heterologous expression of VWF results in formation of WPB in some cells but not others37, thus some cell-specific machinery must be in operation, but this elusive component has not yet been identified.

Finally: why is constitutively released VWF actively targeted for basolateral release? Substrate-bound VWF is known to promote anchoring along with collagen,
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at least in vitro, of circulating plasma VWF$^{38,39}$ and platelets$^{40}$ under shear flow, which is critical in producing a haemostatic plug during vascular injury when the entire endothelial layer has been lost and the subendothelial matrix is exposed to shear stress. We note that endothelial cells specifically target this material for incorporation into the matrix via AP-1, implying that this is not non-specific leakage but a deliberate provision of this pool, and that the VWF provided in this way is of low multimeric state. Since the vast majority of in vitro studies investigating the role of immobilised VWF use human plasma as their source of VWF (which contains both LMW and HMW, but is devoid of UL multimers) it could be that constitutively secreted VWF from endothelial cells is itself the source of functional sub-endothelial VWF, and/or it acts to increase the efficiency of recruitment of plasma VWF. Further in vitro studies comparing the functional differences when either UL-VWF or LMW-VWF are used as source material for matrix-bound VWF would help to dissect this issue.
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Authorship Contributions

M.L.S. designed, performed the research, analysed the data and wrote the paper.
D.F.C. designed research and wrote the paper.

Disclosure of Conflicts of Interest

The authors declare no conflict of interest.
References


Figure Legends

Figure 1. Multimeric states of endothelial cell VWF secreted from three different pathways.

(A) Representative multimer pattern of secreted VWF collected for 30 minutes from unstimulated, treated with 5µM of Brefeldin A (‘+BFA’) for 1 h prior and during secretion collection, treated with 100µM Histamine or 100 ng/mL of PMA. VWF ELISA was used to measure the amount of VWF secreted in all conditions and the same units of VWF were loaded on an SDS/agarose gel (migrations shown from top to bottom). A parallel sample of human plasma is shown for comparison (‘Plasma VWF’). Relative VWF amount ratios in this experiment: 1:1:5:16; Constitutive: Basal: Histamine: PMA.

(B) Representative line plots from the multimer patterns shown in (A). A.U; Arbitrary Units.

(C) Diagram of multimer patterns of plasma VWF and VWF secreted via the Constitutive, Basal and Regulated secretory pathways from endothelial cells. UL, Ultra Large; HMW, High Molecular Weight; LMW, Low Molecular Weight.

Figure 2. Confluent HUVECs on polyester membranes form a polar monolayer.

(D) Immunofluorescence of confluent HUVECs grown on polyester membranes for the times indicated (days) and stained for the cell’s nucleus (DAPI)(blue), VWF (red) and either VE-Cadherin, β-catenin or ZO-1 (green). Scale bar 50µm.

(E) TEM of a cross-section of a HUVEC (i) with the TGN to the right of the nucleus (*). Two examples of TEM cross-sections of overlapping tips between neighbouring HUVECs (ii-iii) and the close contact formed between the cell membranes of two adjacent cells (iv-v) with ‘fuzz’ like structure typical of Tight Junctions in epithelial cells (arrows). Scale bar: i 2µm, ii-iii 1µm, iv-v 300nm.

(F) Dextran permeability assay, where 1 mg/ml of 40kD-FITC-Dextran was loaded on the apical chamber of empty membranes (‘No cells’), 2 day confluent unstimulated HUVECs (‘-‘), or treated with 100µM Histamine
('+Hist') or 2.5 U/ml Thrombin ('+Thro') to open extracellular pores. The amount of FITC-Dextran in the basolateral chamber was measured after incubation with FITC-Dextran in the apical chamber for 30 minutes. Mean of triplicate wells (with SD), as % of ‘No cells’ control sample. (*p<0.05, **p<0.005, n.s. not significant).

**Figure 3. Polarity of VWF secretion.**

(A) Unstimulated VWF secretion on Transwell membranes of HUVECs grown for the time indicated (in days), expressed as a % of secreted VWF from the total VWF in each sample (secreted plus lysate). Results standardized to ‘D1’ sample. Representative experiment, mean of three replicates with SD.

(B) Unstimulated VWF secretion from HUVECs grown on Transwell membranes of HUVECs grown for 2 days. Cells were either left untreated (‘-BFA’) or were treated with 5µM of Brefeldin A (‘+BFA’) for 1 h prior and during secretion collection from both chambers. Results standardized to control ‘-BFA’ sample. Mean of three independent experiments with SEM. (**p<0.005).

(C) Lum-GFP secretion on Transwell membranes. HUVECs were nucleofected with lum-GFP construct and seeded on Transwell membranes for 30 h prior to secretion assay. Cells were treated with either Control (‘-BFA’) or 5µM of Brefeldin A (‘+BFA’) for 1 h prior and during secretion collection from both chambers. Amount of lumGFP secreted expressed as a percent of total GFP in each sample (secreted plus lysate). Representative experiment, mean of three replicates with SD. (**p<0.0005).

(D) Stimulated VWF secretion on Transwell membranes of HUVECs grown for 2 days. Cells were stimulated with either 100µM Histamine (‘Hist’), 100µM Histamine plus 100µM Adrenaline plus 100mM IBMX (‘Hist +Adr +IBMX’), 40ng/mL VEGF, or 100 ng/mL PMA for 30 minutes while secreted VWF was collected from both chambers. Amount of VWF secreted expressed as a percent of total VWF in each sample (secreted plus lysate). Representative experiment, mean of three replicates with SD.
Histamine VWF secretion on Transwell membranes of HUVECs grown for the time indicated (in days), expressed as a percent of total VWF in each sample (secreted plus lysate), relative to control (‘D1’) sample. Results standardized to ‘D1’ sample. Representative experiment, mean of three replicates with SD.

Histamine VWF secretion on Transwell membranes of HUVECs grown for 2 days. Cells were treated with either Control (‘-BFA’) or 5µM of Brefeldin A (‘+BFA’) for 1 h prior and during 30 min 100µM Histamine stimulation and secretion collection from both chambers. Results standardized to control ‘-BFA’ samples. Mean of three independent experiments with SEM.

Representative multimer pattern of secreted VWF collected from unstimulated and Histamine stimulated HUVECs. Cells were treated with either Control (‘-BFA’) or 5µM of Brefeldin A (‘+BFA’) for 1 h prior to and during VWF secretion collection in both chambers. Unstimulated secretion was collected for 1 h while 100µM Histamine stimulated secretion was collected for 30 min. The same units of VWF (as measured by ELISA) was loaded for all samples. VWF ELISA quantification corresponding to these samples are shown in Figure 3B and F. A, apical chamber; B, basolateral chamber. Lanes 1-4, unstimulated releasate, lane 5-8 Histamine stimulated releasate.

Figure 4. AP-1 regulates VWF sorting into WPBs in HUVECs.

PCR bands to detect the presence of AP-1 subunits in HUVECs and HT-29 cells. Primers specific for Actin (‘Act’), AP-1µ1A (‘µ1A’), AP-1µ1B (‘µ1B’) and AP-1γ1 (‘γ1’) were used for amplification of cDNA made from extracted RNA from cell lysates.

qRT-PCR measuring the efficiency of mRNA down-regulation of the AP-1µ1A subunit in HUVECs after 2 rounds of nucleofection with siRNA specifically targeting AP-1µ1A (siAP-1M1), 2 days apart. Mean of 12 experiments with SEM (***p<0.0005).

Unstimulated VWF secretion on plastic of HUVECs either Mock nucleofected or nucleofected with siAP-1M1. Cells were treated with either Control (‘-BFA’) or 5µM of Brefeldin A (‘+BFA’) for 1 h prior and during VWF secretion
collection. Results standardized to ‘Mock –BFA’ samples. Mean of 5 independent experiments with SEM. (**p<0.005).

(D) Histamine (100µM) stimulated VWF secretion on plastic of HUVECs either Mock nucleofected or nucleofected with siAP-1M1. Results standardized to Mock samples. Mean of 5 independent experiments with SEM. (**p<0.005).

Figure 5. AP-1 regulates basolateral sorting of VWF constitutive carriers.

(A) Unstimulated VWF secretion on Transwell membranes of HUVECs either Mock nucleofected or nucleofected with siRNA specifically targeting AP-1µ1A (siAP-1M1). Results standardized to Mock samples. Mean of 5 independent experiments with SEM. (*p<0.05, **p<0.005).

(B) Unstimulated VWF secretion on Transwell membranes of HUVECs either Mock nucleofected or nucleofected with siAP-1M1. Cells were treated with either Control (‘-BFA’) or 5µM of Brefeldin A (‘+BFA’) for 1 h prior and during VWF secretion collection. Results standardized to Mock samples. Mean of 3 independent experiments with SEM. (**p<0.005).

(C) Lum-GFP secretion on Transwell membranes. HUVECs were either Mock nucleofected or nucleofected with siRNA specifically targeting AP-1µ1A (siAP1) and lumGFP construct was added during the second round of nucleofection and seeded on Transwell membranes for 30 h prior to secretion collection. Cells were treated with either Control (‘-BFA’) or 5µM of Brefeldin A (‘+BFA’) for 1 h prior and during secretion collection in both chambers. Amount of lumGFP secreted expressed as a percent of total GFP in each sample (secreted plus lysate). Mean of 3 replicates with SD. (**p<0.005).

(D) Histamine (100µM) stimulated VWF secretion on Transwell membranes of HUVECs either Mock nucleofected or nucleofected with siRNA specifically targeting AP-1µ1A (siAP1). Results standardized to Mock samples. Mean of 7 independent experiments with SEM.

(E) Representative multimer pattern of secreted VWF collected from unstimulated and 100µM Histamine stimulated HUVECs either Mock nucleofected or nucleofected with siAP-1M1. Unstimulated secretion was collected for 1 h while Histamine stimulated secretion was collected for 30 min. The same units
of VWF (as measured by ELISA) was loaded for all samples. VWF ELISA quantification corresponding to these samples are shown in Figure 5A and D. A, apical chamber; B, basolateral chamber. Lanes 1-4, unstimulated releasate, lane 5-8 Histamine stimulated releasate.

Figure 6. The pathways of VWF secretion and their polarity.

Diagram of a blood vessel showing the relative amount (length of arrows) and direction of secreted VWF from the three different secretory pathways from endothelial cells. The direction of VWF secreted either to the apical (vessel lumen) or the basolateral side (subendothelial matrix) of endothelial cells from each pathways is indicated by the direction of the arrows.

Release from the constitutive pathway, which is Brefeldin A -sensitive, is exclusively to the basolateral side of endothelial cells, and is mostly composed of LMW-VWF multimers. Basal VWF release, the continuous release from WPBs, occurs mostly towards the apical side of endothelial cells with a small amount secreted basolaterally, and is composed of UL-VWF multimer together with a dimer band. Regulated release, originating from WPB after a secretagogue stimulant, occurs predominantly to the apical side with some release to the basolateral side and is composed of exclusively UL-VWF multimers.
Figure 2

A

Day 1 | Day 2 | Day 3 | Day 4

VE-cadherin

β-Catenin

ZO-1

B

C

% of Dextran permeated in 30 mins (as % of empty filters)

No cells  Hist  Thro  HUVECs

**  *  *
Figure 3

A) % of Unstimulated secreted vWF

B) % of Histamine secreted vWF (relative to Mock)

C) % of secreted lum. GFP

D) % of secreted vWF

E) % of Histamine secreted vWF

F) % of Histamine secreted vWF (relative to Mock)

G) Unstimulated release

Histamine stimulated

Lane        1       2       3       4      5       6       7      8

Lane 1-2: Unstimulated release
Lane 3-4: Histamine stimulated
Lane 5-6: Unstimulated release
Lane 7-8: Histamine stimulated
Figure 4

A) HUVEC and HT-29 cells with Act μ1A, μ1B, γ1 Act μ1A, μ1B, γ1 stained.

B) Bar graph showing AP-1M1 RNA levels with Mock and siAP-1M1 treatments.

C) Bar graph showing % of unstimulated secreted VWF with Mock, BFA, and siAP-1M1 treatments.

D) Bar graph showing % of histamine secreted VWF with Mock and siAP-1M1 treatments.
Figure 5

A

% of Unstimulated secreted VWF

Mock  siAP-1M1

B

% of Unstimulated secreted VWF

Mock  siAP-1M1

C

% of secreted hGFP

Mock  siAP-1M1

D

% of Histamine secreted VWF

Mock  siAP-1M1

E

Unstimulated release

Histamine stimulated

Mock  siAP-1M1

Mock  siAP-1M1

Lane 1 2 3 4 5 6 7 8
Figure 6

Endothelial VWF release

BLOOD VESSEL LUMEN

SUBENDOTHELIAL MATRIX

<table>
<thead>
<tr>
<th>Constitutive release</th>
<th>Basal release</th>
<th>Regulated release</th>
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From Weibel-Palade Bodies