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Role of reactive oxygen and nitrogen species in the vascular responses to inflammation

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Abstract

Inflammation is a complex and potentially life-threatening condition that involves the participation of a variety of chemical mediators, signaling pathways, and cell types. The microcirculation, which is critical for the initiation and perpetuation of an inflammatory response, exhibits several characteristic functional and structural changes in response to inflammation. These include vasomotor dysfunction (impaired vessel dilation and constriction), the adhesion and transendothelial migration of leukocytes, endothelial barrier dysfunction (increased vascular permeability), blood vessel proliferation (angiogenesis), and enhanced thrombus formation. These diverse responses of the microvasculature largely reflect the endothelial cell dysfunction that accompanies inflammation and the central role of these cells in modulating processes as varied as blood flow regulation, angiogenesis, and thrombogenesis. The importance of endothelial cells in inflammation-induced vascular dysfunction is also predicated on the ability of these cells to produce and respond to reactive oxygen and nitrogen species. Inflammation seems to upset the balance between nitric oxide and superoxide within (and surrounding) endothelial cells, which is necessary for normal vessel function. This review is focused on defining the molecular targets in the vessel wall that interact with reactive oxygen species and nitric oxide to produce the characteristic functional and structural changes that occur in response to inflammation. This analysis of the literature is consistent with the view that reactive oxygen and nitrogen species contribute significantly to the diverse vascular responses in inflammation and supports efforts that are directed at targeting these highly reactive species to maintain normal vascular health in pathological conditions that are associated with acute or chronic inflammation.

Keywords

Vasomotor dysfunction; Leukocyte–endothelial cell adhesion; Angiogenesis; Vascular permeability; Coagulation; Thrombosis; Free radicals

Introduction

Inflammation is a manifestation of immune system function that is triggered by microbial invasion and/or tissue injury. Accordingly, an inflammatory response is directed toward isolating and destroying invading microorganisms and injured cells and preparing the tissue for eventual repair and regeneration. Both the induction and the resolution phases of the inflammatory response are critically dependent on functional and structural changes in the

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microcirculation. These changes include impaired vasomotor function, the recruitment of leukocytes, diminished endothelial barrier function, angiogenesis, and enhanced thrombosis [1–6]. A variety of seemingly unrelated diseases that have been linked to inflammation, including cancer, obesity, diabetes, hypertension, ulcerative colitis, and Alzheimer disease, appear to exhibit all or most of these abnormalities in microvascular function. Given the growing recognition that inflammation is an underlying mechanism in many diseases associated with significant morbidity and mortality, much attention has been devoted to defining the mechanisms that link inflammation to microvascular dysfunction.

Two major effector systems that are frequently implicated in the vascular alterations associated with inflammation involve the generation of reactive oxygen and nitrogen species. The literature in this area is generally consistent with the view that the enhanced production of reactive oxygen species (ROS) and diminished bioavailability of nitric oxide (NO) that accompany an inflammatory response play a pivotal role in mediating the microvascular dysfunction and that restoration of the normal balance between ROS and NO will return vascular function to a normal state [7–12]. Although it remains unclear how the imbalance between ROS and NO levels caused by inflammation can exert an influence on responses as diverse as impaired vasomotor function, angiogenesis, endothelial barrier dysfunction, and thrombogenesis, the dual role of endothelial cells as a source of ROS and NO and as a major target of the signaling mechanisms activated by these reactive species may be important in this regard.

The objective of this article is to describe the contributions of ROS and NO to the microvascular alterations that are characteristic of an inflammatory response. The focus here is less on whether ROS and NO are involved in the vascular dysfunction of inflammation but more directed toward how these reactive species exert the vascular changes by interacting with specific targets in cells that either comprise or surround the blood vessel wall. For each of the five characteristic vascular abnormalities associated with inflammation (impaired vasomotor function, leukocyte recruitment, endothelial barrier dysfunction, angiogenesis, and enhanced thrombus formation) we provide background information on the physiological process and follow this with an analysis of the molecular signals that are targeted by ROS or NO to yield the observed vascular abnormality.

Vasomotor dysfunction

Inflammation is generally associated with an altered capacity of resistance vessels to respond to endothelium-dependent vasodilators and vasoconstrictors [1]. It is generally accepted that vascular smooth muscle tone can be regulated by adjacent endothelial cells. Arteriolar endothelium is separated from the underlying vascular smooth muscle by the internal elastic lamina, the mean thickness of which is approximately 1.5 μ m [13]. Endothelial cell projections through the internal elastic lamina bring the membranes of endothelial cells and myocytes into even closer apposition (0–200 nm) [14]. At points of contact, myoendothelial gap junctions (MEGJs) are present, with a greater number of MEGJs in the distal than in the proximal arteriolar segments. The MEGJs allow for direct bidirectional communication between the two cell types as evidenced by the use of gap junction inhibitors in isolated arterial segments [15] or coculture of endothelial and vascular smooth muscle cells [16].

Endothelium-dependent vasodilation (Fig. 1)

Interest in endothelium-dependent vasodilation stemmed from the observation that removal of the endothelial cells from isolated arteries prevented the vasodilator response to acetylcholine [17]. Since then, numerous other vasoactive substances, such as G-protein-coupled receptor (GPCR) ligands, as well as shear stress, have been shown to exhibit a

similar dependence on an intact endothelium for induction of vasodilation [18]. Based on pharmacologic blockade and bioassay approaches, endothelium-dependent vasodilation has been attributed to the release of prostacylin (PGI₂) and NO, which are referred to as endothelium-derived relaxing factors (EDRFs) [17,19,20]. Based on the relative impact of pharmacologic inhibitors of nitric oxide synthases (NOSs) and cyclo-oxygenases (COXs) on endothelium-dependent vasodilation, NO seems to be the dominant EDRF and therefore it has received the most attention.

The residual vasodilation after pharmacologic blockade of cyclooxygenase and NOS is (1) associated with vascular smooth muscle hyperpolarization and (2) sensitive to blockade of Ca^{2+} -activated K⁺ channels. This endothelium-dependent vasodilation has been attributed to some endothelium-derived hyperpolarizing factor or factors (EDHFs) [21,22]. In general, EDHF-induced vasodilation is more prevalent in distal than in proximal arterial segments [22,23], correlating with the relative distribution of MEGJs [14]. A number of candidate EDHFs have been proposed, including cytochrome P450 epoxygenase-derived epoxyeicosatrienoic acids (EETs) [24], mitochondrial- or oxidase-derived H₂O₂ [25–27], and potassium ions/channels or direct electrical coupling via MEGJs [22,23,28].

Because the EDRFs NO and PGI_2 can also hyperpolarize vascular smooth muscle [22], it has been suggested that the term EDHF should refer only to myocyte hyperpolarization as a result of the spread of endothelial hyperpolarization to vascular smooth muscle by direct electrical coupling via MEGJs and/or accumulation of K⁺ ions in the intercellular space between them [22,23,29]. Two distinct types of EDHF pathways have been proposed to deal with this issue: one involving endothelial cell hyperpolarization with spread of the hyperpolarization to the smooth muscle and another involving endothelial release of synthesized chemical mediators that activate K⁺ channels on smooth muscle [30]. The contributions of EDRFs and EDHFs to endothelium-dependent vasodilation vary depending on species, vascular bed, and segment (proximal versus distal) of arteriole being assayed [22,23,31–34]. Further, there are interactions between EDRFs as well as between EDHFs and EDRFs [35–37].

An initial event in endothelial cell activation in response to ligation of GPCR or shear stress is an increase in intracellular Ca²⁺. Ca²⁺ influx occurs via receptor-operated channels (ROCs) and/or store-operated channels (SOCs) [38,39]. ROC activation occurs in response to external stimuli (e.g., GPCR ligation), which stimulate phospholipase C/inositol-3phosphate (PLC/IP₃) signaling. SOC activation is mediated by intracellular signals generated when Ca²⁺ stores are depleted (e.g., endoplasmic reticulum). The molecular identity of the various Ca²⁺ channels remains unclear. Of relevance to endothelial-dependent vasoreactivity, some members of the transient receptor proteins (TRPs) have been shown to be major components of Ca²⁺ channel complexes [38,40]. For example, TRPC1 is believed to be an important ROC, as well as a SOC component, depending on whether it is incorporated into lipid rafts [41]. Transient receptor potential vallinoid type 4 (TRPV4) also plays an important role in Ca²⁺ influx [40]. Both GPCR- and shear-stress-mediated vasodilation are significantly blunted after genetic blockade of TRPV4 [42-44]. Interestingly, TRPV4 and TRPC1 are associated with caveolae [40] and can form a complex to more efficiently promote Ca^{2+} influx [45]. Despite the ambiguity surrounding the molecular composition of the various endothelial Ca²⁺ channels, an increase in intracellular Ca^{2+} drives both EDRF and EDHF pathways of vascular smooth muscle relaxation.

In the EDRF pathway, an increased intracellular Ca^{2+} activates endothelial (e) NOS, which generates NO during the conversion of L-arginine to L-citrulline. NO diffuses to the myocyte where it binds to the heme moiety of soluble guanylate cyclase (sGC) and displaces iron from its usual position in the porphyrin ring allowing sGC to catalyze the formation of

cGMP [20]. Increased levels of myocyte cGMP initiate signaling cascades that result in a decrease in intracellular Ca^{2+} and subsequent smooth muscle relaxation. The decrease in myocyte intracellular Ca^{2+} is a result of efflux of K⁺ due to activation of large conductance channels (BK_{Ca}) resulting in hyperpolarization of the myocyte membrane and reduced influx of Ca^{2+} [28,46]. Alternatively, NO can induce relaxation of smooth muscle via sGC-independent mechanisms (e.g., *S*-nitrosothiols) [20,47]. By contrast, PGI₂ ligation of the prostacyclin receptor, IP, results in the generation of cAMP, which initiates a signaling cascade leading to a decrease in myocyte Ca^{2+} levels via activation of BK_{Ca} [28,46].

With respect to the EDHF pathway, the increase in intracellular Ca^{2+} results in activation of small and intermediate conductance Ca^{2+} -activated K⁺ channels (SK_{Ca} and IK_{Ca}) located in endothelial caveolae and cell projections [22,23,29]. Genetic blockade approaches have revealed that functional SK_{Ca} and IK_{Ca} channels are critical for both agonist and shear-stress-induced EDHF pathways of vasodilation [32]. Movement of K⁺ out of the cell (due to activation of SK_{Ca} and IK_{Ca}) results in hyperpolarization of the endothelial cells, which can spread to the myocyte via current conduction and/or movement of signaling molecules through the MEGJs. In addition, the extracellular accumulation of endothelial-derived K⁺ can hyperpolarize adjacent myocytes via stimulation of Na⁺/K⁺-ATPase and/or inward rectifier K⁺ channels [22]. Finally, vascular smooth muscle hyperpolarization/relaxation induced by synthesized and diffusible EDHFs (e.g., H₂O₂, EETs) is a result of activation of BK_{Ca} in vascular smooth muscle [30].

Role of ROS and NO (Fig. 1)

NO derived from eNOS is an EDRF whose mechanism of action has classically been attributed to activation of sGC in vascular smooth muscle [20]. NO can also undergo interconversion to nitroxyl anion (NO⁻), which exists as HNO in an aqueous milieu [31,48]. HNO seems to contribute equally with NO to the acetylcholine-induced, sGC-dependent vasodilation in rat and mouse mesenteric arteries [30], indicating that HNO can also be considered an EDRF. Further complicating matters, NO may also substantially contribute to the EDHF-mediated vasodilation. For example, acetylcholine-induced, EDHF-mediated vasodilation (i.e., eNOS and COX independent) can be prevented by scavenging NO [31]. This can be attributed to NO released from endothelial stores independent of eNOS activation, with nitrites and/or S-nitrosothiols as the most likely storage pool of NO [20]. Alternatively, activation of endothelial SK_{Ca} and IK_{Ca} channels, a hallmark feature of the EDHF pathway, also enhances NO synthesis via eNOS and contributes to acetylcholineinduced vasodilation [49,50]. Endothelium-derived H_2O_2 has been proposed to be an EDHF based on the following lines of evidence. Endothelial superoxide and H_2O_2 production is increased in mesenteric and coronary arteries in which endothelium-dependent vasodilation is resistant to pharmacologic blockade of NOS and cyclo-oxygenase [34,51]. Catalase can inhibit both agonist- and flow-induced endothelium-dependent vasodilation in isolated arteries [26,35,51]. Catalase can also prevent acetylcholine-induced smooth muscle hyperpolarization and relaxation, whereas exogenous H₂O₂ can induce hyperpolarization and relaxation of denuded arteries [25,52]. The endothelial component of the H₂O₂mediated response has been attributed to enhanced Ca^{2+} release from cellular stores (e.g., endothelium reticulum) and the secondary activation of SK_{Ca} and IK_{Ca} channels [35,53]. There is evidence that H2O2 can interact with smooth muscle after extracellular release from endothelial cells [51] or by intracellular communication via MEGJs [52,53].

Endothelial generation of H_2O_2 requires the presence of Cu,Zn-superoxide dismutase (SOD) [54,55], supporting the contention that H_2O_2 is derived from the dismutation of superoxide. The endothelial sources of superoxide generation in response to ligation of GPCR or shear stress are not entirely clear but may be species, vascular bed, and/or stimulus specific. In human coronary arteries, flow-induced vasodilation and H_2O_2 production have been

attributed to mitochondrial respiration; neither NOS nor NADPH oxidase seems to be involved [56]. However, in the same vascular bed, bradykinin-induced vasodilation and the associated endothelial production of H_2O_2 are dependent on NADPH oxidase [57]. Acetylcholine-induced H_2O_2 production and vasodilation in mesenteric arteries have been attributed to NOS; neither mitochondrial respiration nor other oxidases (e.g., xanthine oxidase, NADPH oxidase) have been implicated in this response [58].

There are significant antagonistic interactions between endothelial-derived ROS and NO, which can have an impact on the endothelium-dependent vasodilation induced by agonists or shear stress. Superoxide can interfere with NO-induced activation of sGC by interacting with NO, an event ameliorated by Cu,Zn-SOD [7]. Conversely, the direct interaction between superoxide and NO can diminish the ROS component of endothelium-dependent vasodilation. In addition, oxidant stress can lead to direct inhibition of eNOS by inducing phosphorylation of Tyr657 [59] or the uncoupling of eNOS via oxidation of the cofactor tetrahydrobiopterin (BH_4) [60,61]. Another interesting interaction between NO and ROS has been uncovered in endothelial cells exposed to shear stress for up to 24 h [62]. Short durations of shear stress (2 h) increase both ROS and NO production by endothelial cells and, as the duration of shear stress increases, endothelial ROS production declines, whereas NO production continues to increase. The decline in ROS production has been attributed to NO-induced downregulation of NADPH oxidase. Finally, if an agonist can interact with GPCRs on endothelial cells and smooth muscle cells, the end result may not be predictable. For example, there is evidence that smooth-muscle-derived superoxide can negatively modulate endothelial-dependent, NO-mediated vasodilation [63]. In this scenario, ligation of a GPCR (e.g., 5-HT) on smooth muscle cells generates superoxide, which can traverse MEGJs to interact with eNOS-derived NO, resulting in diminished NO bioavailability for activation of myocyte sGC.

ROS and NO may act cooperatively in the endothelium-dependent vasodilation induced by agonists or shear stress. NOS-derived superoxide may be required for the generation of H_2O_2 implicated in the EDHF-mediated, endothelium-dependent vasodilation. Genetic deletion of all three isoforms of NOS results in the abolition of acetylcholine-induced H_2O_2 production and vasodilation in mesenteric arteries [34]. Because BH₄ bioavailability was not affected, it was assumed that NOS uncoupling was not involved in the generation of superoxide and subsequent formation of H_2O_2 . However, oxidation of BH₄, rather than BH₄ depletion per se, seems to be a prerequisite for eNOS uncoupling [64]. Furthermore, limiting concentrations of other factors that regulate eNOS activity (e.g., substrate) can also lead to eNOS uncoupling [65]. Finally, the validity of these observations has been questioned based on the rather severe phenotype of the triple-NOS knockout mouse [22]. Irrespective, blockade of NOS via pharmacologic or knockdown approaches supports an NOS/H₂O₂ pathway in acetylcholine-induced vasodilation [66]. It has been proposed that the NOS/NO EDRF pathway is dominant in arteries and the NOS/H₂O₂ EDHF pathway is dominant in arterioles [27].

On the other hand, exogenous H_2O_2 can induce endothelium-dependent vasodilation that is blocked by inhibition of NOS, indicating that H_2O_2 can activate eNOS [60]. However, this phenomenon appears to be dose dependent. Low concentrations ($\approx 50 \ \mu$ M) of H_2O_2 induced eNOS phosphorylation on Tyr657 (inhibitory site) while not affecting eNOS phosphorylation on Ser1177 (stimulatory site) [59]. These phosphorylation events diminished eNOS activation by bradykinin and inhibited acetylcholine-induced vasodilation. On the other hand, high concentrations of H_2O_2 ($\approx 500 \ \mu$ M) can induce eNOS activation via phosphorylation of Ser1177 and dephosphorylation of Thr495 (inhibitory site) [67]. The ability of H_2O_2 and NO to regulate each other's bioavailability argues in favor of endothelial compartmentation of the systems involved in ROS and NO generation [68].

Shear-induced vasodilation: endothelial mechanosensing/transduction

The vasodilation induced by shear stress is dependent on endothelial sensing/transduction of the shear induced by flowing blood. The proposed mechanosensors on the luminal surface of the endothelium include components of the glycocalyx (glycoproteins and proteoglycans), caveolae, and ion channels [28,40,69–73]. The major intracellular mechanotransduction element seems to be the endothelial cytoskeleton [70,74].

Functional components of mechanosensing/transduction

The glycocalyx decorating the endothelial cell surface consists of proteoglycans and associated glycosaminoglycan (heparan sulfates, chondroitin sulfates, and hyaluronan) side chains and terminal sialic acid [75,76]. Two of the major proteoglycan core proteins of the glycocalyx are syndecan and glypican; syndecan is a transmembrane proteoglycan and glypican is anchored to membrane lipid rafts/caveolae via glycosylphosphatidylinositol. Enzymatic destruction of the glycosaminoglycans (GAGs) can severely blunt shear-induced vasodilation [77,78]. The sensitivity of the glycocalyx to proteolytic degradation may result in a less impressive structure when endothelial cells are enzymatically isolated and cultured [79]. However, this may be a fixation artifact, because rapid freezing of the cells reveals a glycocalyx comparable to that noted in vivo [80]. Whether or not the glycocalyx is compromised during the isolation and culture of endothelial cells, there is a sufficient structural integrity to allow the cells to respond to shear stress in a manner analogous to the in vivo situation [79]. Platelet-endothelial cell adhesion molecule-1 (PECAM-1), a transmembrane glycoprotein that serves as one of the core proteins of the glycocalyx [75,76], can be activated (phosphorylated) by shear stress [81] or mechanical "tugging" with magnetic beads [72]. Flow-induced vasodilation is blunted in arterioles from PECAM-1deficient mice [82].

Caveolae are flask-shaped invaginations in the endothelial membrane that are enriched in cholesterol and sphingolipids and provide a microdomain for a variety of signaling complexes (e.g., Ca²⁺-handling proteins) [40,83,84]. Caveolin-1, a protein constituent of caveolae, plays a role in caveola assembly [85] and, more importantly, caveolin-1 interacts with various signaling components to regulate flow-induced vasodilation [86]. Caveolin-1-deficient arteries exhibit a marked reduction in flow-induced dilation that is rescued by selectively reconstituting caveolin-1 expression in endothelium [85].

Calcium channel activation and endothelial Ca^{2+} influx are two of the earliest events in shear-induced endothelial activation [70]. For example, shear stress activates TRPV4 in endothelial cells, resulting in Ca^{2+} influx [43]. Furthermore, flow-mediated vasodilation is inhibited by interfering with TRPV4 activity (genetic, pharmacologic, and siRNA blockade).

The endothelial cytoskeleton, an intracellular network of actin microfilaments, microtubules, and intermediate filaments, plays an important role in shear-induced alterations in endothelial cell morphology and flow-induced vasodilation [74]. Pharmacologic depolymerization or stabilization of the microfilament/microtubule network impairs flow-induced vasodilation without affecting agonist-mediated vasodilation [74].

A unifying hypothesis

The various mechanosensor/transducers implicated in the vasodilation induced by shear stress most probably do not operate independently, but rather work in concert as interconnected networks [70]. Individual blockade of the various proposed mechanosensors/ transducers does not distinguish whether they are involved in the sensing versus the transduction of the shear stress. It has been predicted that fluid drag within the glycocalyx reduces the shear stress to negligible levels at the endothelial cell membrane proper [76].

Thus, it seems unlikely that membrane structures (e.g., caveolae and associated receptors, enzymes, or ion channels) play an important direct role in the detection of shear by endothelial cells with an intact glycocalyx. However, the GAGs (e.g., hyaluronan) and core proteins (e.g., PECAM-1) of the glycocalyx that readily respond to shear stress can interact with the cytoskeleton [70,74] and signaling components of lipid rafts/caveolae [87,88]. Many of the relevant ion (K^+ and Ca^{2+}) channels in the endothelial cell membrane proper are associated with lipid rafts/caveolae [28,39,40,68,89]. For example, the endothelial Ca^{2+} influx induced by shear stress can be prevented by enzymatic disruption of the glycocalyx [70]. Collectively, the literature is consistent with a unified hypothesis encompassing an initial sensing of shear stress by glycocalyx components that ultimately activate signaling complexes located in lipid rafts/caveolae. This hypothetical scenario has been alluded to in the past with respect to flow-induced arterial remodeling [74] and warrants further attention, specifically to address the sequence in which the various mechanosensors/transducers operate to elicit the vasodilation associated with shear stress.

Linkage to ROS and NO

Endothelial production of NO induced by shear stress is inhibited by enzymatic degradation of GAGs in vivo and in vitro [73,87]. Selective enzymatic degradation of various GAGs indicates that heparan sulfates (HS), hyaluronan (HA), and sialic acid (SA) constituents, but not chondroitin sulfate, are critical for the shear-induced increase in endothelial NO (nitrite) [87,90]. Interestingly, the decline in endothelial NO associated with degradation of HS and SA, but not HA, can be prevented by scavenging ROS [90]. These findings indicate that only HA is directly involved in endothelial NO production; the role of HS and SA is indirect, serving to limit ROS generation and increase NO bioavailability. It has been proposed that HA can activate eNOS either through direct binding to CD44 in caveolae, which contain eNOS, or indirectly via the HA-rich glypican binding to caveolae (glypican/caveolae/eNOS axis) [87,90].

As mentioned above, degradation of the HS and SA constituents of the glycocalyx results in increased ROS production; consequently the inability to detect endothelial NO production probably reflects reduced NO bioavailability [90]. This is in contrast to observations that, in endothelial cells exposed to shear stress, H_2O_2 activates eNOS (Ser1177 phosphorylation) and increases NO production, an event associated with a decrease in catalase activity initiated by a protein kinase C γ (PKC γ)-mediated phosphorylation event [91]. Interestingly, PKCs are enriched in caveolae [68], but whether and how the glycocalyx is linked to PKC γ in caveolae remain unclear.

The glycocalyx-associated glycoprotein PECAM-1 plays an important role in flow-induced vasodilation, but its direct relationship with eNOS activation is controversial [70]. For example, PECAM-1 association with eNOS has been shown to increase [81] or decrease [88] with application of shear to the endothelium. Flow-induced vasodilation is reduced in PECAM-1-deficient coronary arteries, but the portion of the dilation attributable to NO is not affected and, indeed, eNOS Ser1177 phosphorylation and NO production are still noted in these arteries [92]. One issue worth exploring is the role of the endothelial mechanosensory PECAM-1/vascular endothelial growth factor receptor 2 (VEGFR2) signalosome (or complex) [93] in flow-induced vasodilation. The components of the complex are associated with caveolae [68] and VEGFR2 can activate eNOS in a ligand-independent manner and plays a role in flow-induced vasodilation [94]. Thus, blockade (genetic or otherwise) of either one of the components of the PECAM-1/VEGFR2 signalosome would more than likely render it nonfunctional and inhibit flow-induced vasodilation.

Caveolin-1 binding to eNOS is important in maintaining eNOS within the caveolae while inhibiting its enzymatic activity [68,70]. An increase in shear stress enhances phosphorylation of both caveolin-1 and eNOS (Ser1177) [91,95]. Subsequently, caveolin-1 disassociates from eNOS leaving eNOS free to interact with calmodulin, resulting in increased eNOS activity [68,70,84]. As mentioned above, flow-induced vasodilation is impaired in caveolin-1-deficient mice and this impairment can be rescued by selectively overexpressing caveolin-1 in the endothelium [85,86].

Although shear stress has been shown to increase ROS production by endothelial cells [62,90], little is known about the potential links between mechanosensory/transduction components of endothelial cells and ROS-generating systems. Enzymatic degradation of HS or SA, but not HA, prevents shear-induced ROS production [90]. These findings indicate that shear-induced ROS production is dependent on HS and SA components of the glycocalyx. This differs from shear-induced NO production, which seems to be primarily dependent on HA [90].

Flow-induced ROS production is more likely to be demonstrated in arteries derived from animals or humans with pathologies associated with cardiovascular complications. The impaired flow-induced vasodilation in experimental hypertension (SHR) has been attributed to NADPH oxidase activation [96]. Angiotensin II, which is known to activate endothelial cell NADPH oxidase, has been implicated in the impaired vasodilatory response associated with hypertension [97]. In coronary arteries from patients with coronary artery disease, the impaired flow-induced vasodilation has been attributed to mitochondrial-derived superoxide and H_2O_2 [98].

Endothelium-dependent vasoconstriction

In addition to mediating vasodilation via EDRFs and EDHFs, the endothelium can mediate vasoconstriction via release of factors that elicit contraction of the surrounding smooth muscle. These endothelium-derived contracting factors (EDCFs) were identified using approaches similar to those utilized to characterize EDRFs and EDHFs, i.e., endothelial denudation of blood vessels, pharmacologic blockade, and bioassays [99,100]. In general, EDCFs are uncovered (1) after pharmacologic blockade of EDRFS and/or EDHFs (e.g., NO) [100] or (2) during pathogenesis of cardiovascular disorders (e.g., hypertension) in which NO bioavailability is compromised [101,102]. The decreased NO bioavailability in various pathologic conditions seems to result from enhanced ROS production via cyclo-oxygenase [103] or NADPH oxidase [104]. Endothelium-dependent contractions can be enhanced by compromising NO bioavailability or amplifying ROS bioavailability [105].

The major candidate EDCFs are COX-derived prostanoids such as thromboxane, prostaglandin $F_{2\alpha}$, and PGI_2 , with PGI_2 receiving the most experimental support [99– 101,106,107]. As mentioned above PGI_2 is considered an EDRF that induces smooth muscle relaxation by activating IP receptors on myocytes. The PGI_2 -mediated vasoconstriction is a result of PGI_2 ligation of TP receptors, which induce smooth muscle contraction via either Ca^{2+} release from intracellular stores [99,105] or inhibition of the cGMP/cAMP vasodilator pathway [108]. Although both IP and TP receptors are expressed on smooth muscle, in the SHR model of hypertension, the IP receptors are dysfunctional [109], whereas the TP receptors are hyperresponsive [105].

Role of ROS and NO

EDCFs were uncovered at around the same time as EDRFs [101], yet other than the interaction between NO and superoxide, little is known about the potential direct contribution of endothelial ROS or NO to EDCF-induced vasoconstriction. This presumably

results from the complexities inherent in pathologic conditions (e.g., hypertension) wherein the mechanisms underlying the oxidative stress and decreased NO availability remain undefined. In one model (rat renal artery), with acetylcholine inducing an endotheliumdependent vasoconstriction under conditions of normal NO bioavailability, endothelial NADPH oxidase-derived H₂O₂ was shown to behave as an EDCF [110]. Nonetheless, the current consensus holds that ROS contribute to EDCF-mediated vasoconstriction under certain pathologic conditions. For example, acetylcholine increases endothelial oxidative stress in aortic endothelial cells from hypertensive (SHR) rats, but not in endothelial cells from normotensive (WKY) rats [103].

The following has been proposed to explain the role of ROS in EDCF-mediated vasoconstriction in SHR: upon ligation of GPCR by acetylcholine, there is excessive accumulation of intracellular Ca^{2+} in endothelial cells, which results in enhanced COX activation and ROS production [103]. ROS can further activate COX in endothelial cells and/or diffuse to the neighboring vascular smooth muscle to activate COX and produce more prostanoids. This positive feedback mechanism leads to excessive prostanoid generation that activates hyperresponsive TP receptors and induces smooth muscle contraction [101,105]. Furthermore, ligation of TP receptors on vascular smooth muscle enhances their stability (less internalization) via a ROS-dependent mechanism [111].

Leukocyte recruitment

The major function of the inflammatory response is to clear invading pathogens or damaged tissue/cells and initiate repair [112,113]. Signals derived from injured tissue or invading microbes activate resident sentinel cells, such as macrophages and/or mast cells. If the episode is minor, the macrophages/mast cells clear the interstitial debris/microbes. If the stress is more severe, the sentinel cells recruit circulating neutrophils (polymorphonuclear cells, or PMNs), and later monocytes, to the affected site to aid in the clearing process [112,114]. This sequential recruitment of PMNs and monocytes is believed to result from progressive activation of the endothelium and/or the ability of emigrated PMNs to facilitate monocyte recruitment [115]. To reach the site of injury/infection, leukocytes must first be captured by the endothelium via a coordinated series of adhesive interactions referred to as rolling, adhesion, and emigration [116–119].

Detection of tissue injury/infection by resident sentinel cells (Fig. 2A)

The initiation of an inflammatory response is dependent on the recognition of invading microbes and/or damaged tissue by resident immune cells. The major sentinel cells involved in this innate immune response are the macrophages and mast cells [120–124].With microbial invasion, these cells recognize highly conserved components of microbes termed pathogen-associated molecular patterns (PAMPs), such as lipopolysaccharide (LPS) [125].With sterile tissue injury, the sentinel cells detect substances released by damaged cells and/or the extracellular matrix (ECM), referred to as damage-associated molecular patterns (DAMPs), such as heat shock proteins, high mobility group protein-B1 (HMGB1), and hyaluronan [126,127]. Both macrophages and mast cells can be activated by PAMPs and DAMPs. Although mast cells seem to be capable of phagocytosis [128], clearance of bacteria and/or cellular debris is generally the responsibility of macrophages [114]. Upon activation, both mast cells and macrophages release various inflammatory mediators (e.g., platelet activating factor (PAF)), chemokines (interleukin-8 (CXCL8)), and cytokines (e.g., tumor necrosis factor (TNF), interleukin-1 (IL-1)) [129,130].

A major class of membrane receptors for PAMPs and DAMPs are the Toll-like receptors (TLRs) [125,127]. During infection, various components of bacterial cell membranes interact with different homodimers or heterodimers of TLRs on the cell membranes of

sentinel cells [131–133]. A classic example is the interaction of LPS with macrophage TLR4. During sterile tissue injury, proteins (e.g., HMGB1) released from damaged cells or breakdown products of disrupted ECM (e.g., hyaluronan) can also interact with TLRs (e.g., TLR4, TLR2) [127,131]. Some diversity in the system exists, because HMGB1 can also interact with RAGE (receptor for advanced glycation end products) [134]. There is also evidence that IL-1 α released from necrotic cells utilizes the IL-1R on sentinel cells to initiate an inflammatory response [135]. Activation of TLRs, RAGE, or IL-1R results in the activation/nuclear translocation of transcription factors involved in the inflammatory response (e.g., NF- κ B) [131,136–138]. Collectively, most of the available information indicates that TLRs are the major receptors for PAMPs and DAMPs and that there is a convergence of molecular pathways at the level of NF- κ B. However, it has been proposed that the intensity of the response can be downregulated at the postreceptor level by concurrent activation of a CD24–Siglec pathway [139].

Although cells of the innate immune system are generally believed to be the major sentinel cells, there is evidence indicating that parenchymal cells can serve in this capacity [140,141]. Cardiac myocytes, renal mesangial cells, alveolar epithelial cells, and endothelial cells also possess TLRs and can respond to DAMPs and PAMPs [142–147]. These cells can be converted to a proinflammatory phenotype when activated via TLRs. For example, cardiac myocytes can generate cytokines/chemokines and endothelial cells can increase adhesion molecule expression in response to the TLR4 ligand, LPS [143,145,146].

Tissue injury is often associated with infection, which makes an accurate assessment of the relative roles of PAMPs and DAMPs to the overall inflammatory response rather difficult [112]. Specific experimental models have been used to isolate the inflammatory response to sterile injury (e.g., ischemia/reperfusion; I/R) from that due to infection (e.g., LPS) [112,113]. Based on these approaches, it is well accepted that TLR4 plays an important role in PAMP (LPS)-induced responses [143,145], whereas both TLR4 and RAGE play important roles in sterile (I/R) injury and inflammatory responses [127,134,141]. However, even under these well-defined experimental conditions the potential for interactions between PAMPs and DAMPS should be kept in mind. For example, PAMPs can activate adjacent cells to release DAMPs [127,144], as well as interacting with each other (e.g., HMGB1 binds LPS) [148–151].

Role of ROS and NO (Fig. 2A)

As discussed above, DAMPs and PAMPs generated during injury/infection can activate perivascular (e.g., macrophages, mast cells) and intravascular (e.g., leukocytes) immune cells, as well as endothelial cells; all of which subsequently generate ROS at an accelerated rate. The increased ROS production can also generate oxidation products from host-derived cells/debris, which also can serve as DAMPs and propagate the inflammatory response [152].

Activation of mast cells increases intracellular ROS, which may be involved in the intracellular signaling that leads to secretion of inflammatory mediators [153,154]. For example, NADPH oxidase-derived ROS have been implicated in the generation of IL-8 production by IL-1 β -activated mast cells [155]. Although mast cells contain NOS and can generate NO [156], the role of NO in mast cell activation by inflammatory stimuli is equivocal [157,158]. In short, the available information, albeit limited, indicates that mast-cell-derived ROS and NO are used for intracellular signaling purposes, rather than being exported. By contrast, activation of macrophages results in substantial ROS production via NADPH oxidase and NO production via inducible (i) NOS, both of which are involved in the killing of pathogens [159–162].

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Activation of macrophages by LPS, a TLR4 ligand, induces ROS generation both intracellularly and extracellularly via NADPH oxidase [163,164], with intracellular ROS serving a signaling function (e.g., potentiation of TNFa secretion) and extracellular ROS affecting adjacent cells (e.g., cytotoxicity) [163]. With respect to intracellular signaling, activation of the TLR/IRAK-1 pathway in macrophages can (1) rapidly activate NADPH oxidase via the small GTPase Rac1 and (2) induce transcriptional events (e.g., NF-rB) to increase the expression of a NADPH oxidase subunit, Nox1 [165]. In a cell-free system, IRAK-4 has been shown to interact with and phosphorylate p47phox, leading to rapid activation of NADPH oxidase; a result confirmed in LPS-activated PMNs [166]. Whether a similar role for IRAK-4 is operative in macrophages awaits confirmation. The NADPH oxidase pathway is also critical for the LPS-induced conversion of endothelial cells to a proinflammatory phenotype, e.g., generation of chemokines [146]. Of particular interest with respect to pathology is the proposed propagation of ROS production, i.e., H_2O_2 can activate NADPH oxidase to generate further ROS [167]. Once generated extracellularly, ROS can activate both mast cells and macrophages [153,168,169], thereby amplifying the inflammatory response. Coculture approaches indicate that transfer of oxidants (e.g., H₂O₂) generated by immune cells (high output) can have an impact on adjacent nonimmune cells (neurons, endothelial cells) [163,169,170]. For example, challenge of macrophages with H₂O₂ results in an intracellular oxidant stress and CXCL8 production, events mediated by TLR2 [169]. On the other hand, H_2O_2 can elicit oxidant stress in HEK293 cells, which are devoid of TLRs [169]. The relative importance of receptor-mediated transfer versus passive permeation of H_2O_2 in the transfer of oxidant stress between cells warrants further attention.

Activation of macrophages with LPS also increases iNOS expression, a response that seems to be mediated via the NADPH oxidase/NF- κ B pathway [171]. Intracellular NO may directly interact with ROS (superoxide) and thereby dampen the inflammatory response [160,161,172]. In addition, NO may interfere with the generation of ROS by NADPH oxidase. NO can inhibit Nox1 expression in IL-1-activated mesangial cells via a cGMP signaling mechanism [173]. Alternatively, NO can suppress endothelial cell NADPH oxidase activity by S-nitrosylation of p47phox, rather than via a cGMP pathway [174]. Regardless of the mechanisms involved, NO appears to dampen intracellular oxidative stress/signaling. Indeed, NO is generally considered anti-inflammatory particularly with respect to recruiting PMNs to affected sites [175,176]. Finally, macrophage-derived NO has been shown to dampen mast cell activity [153].

Role of interstitial sentinel cells in leukocyte recruitment

The role of interstitial sentinel cells in promoting leukocyte emigration to sites of tissue injury/infection is well documented in a variety of organ systems [145,177–181]. The primary interstitial immune cells involved in inflammation are the mast cells [182] and macrophages [122]. With respect to mast cells, induction of mast cell degranulation in vivo results in leukocyte emigration across postcapillary venules and their subsequent interstitial migration toward the activated mast cells [177]. Furthermore, the leukocyte emigration elicited by either sterile injury (I/R) or infectious challenge (*Helicobacter pylori* extract or *Clostridium difficile* toxin) is prevented by mast cell stabilizers [177]. In vitro, co-incubation of activated mast cells with endothelial cells increases PMN adhesion to the endothelial cells, whereas co-incubation with nonactivated mast cells does not [158]. With respect to macrophages, depletion of regional macrophages in vivo prevents leukocyte infiltration in response to sepsis induced by peritonitis [183]. In vitro, activated macrophages promote PMN migration across endothelial monolayers [170].

As mentioned above, in addition to interstitial immune cells, parenchymal cells can function as sentinel cells. For example, alveolar type II cells can respond to TLR ligands by

generating cytokines/chemokines and promoting leukocyte emigration, i.e., they can be converted to a proinflammatory phenotype [184]. Similar phenotypic changes in cardiac myocytes can be induced by challenge with septic plasma [185] or anoxia/reoxygenation (A/ R; in vitro counterpart to I/R) [186]. These activated cardiomyocytes can promote leukocyte transendothelial migration (TEM) by releasing inflammatory mediators. Interestingly, activation of endothelial cells with TLR ligands can convert them to a proinflammatory phenotype both in vivo [145] and in vitro [187]. However, if the only responding sentinel cell to PAMPs or DAMPs is the endothelial cell, it is difficult to envision how the leukocytes reach the affected site without additional cues from other interstitial cells. This issue is underscored by the observation that systemic administration of LPS in mice expressing TLR exclusively in endothelial cells induces an increase in leukocyte adhesion to brain endothelium, but the leukocytes fail to emigrate into the brain parenchyma [145].

Role of ROS and NO

It has been proposed that NO derived from eNOS keeps vascular endothelium in a quiescent state and that inhibition or removal of this source of NO upregulates the inflammatory pathway [188]. Studies using intravital microscopy to assess leukocyte interactions with venular endothelium indicate that pharmacologic blockade of NOS leads to oxidative stress within the venular wall and adjacent interstitium. This is accompanied by a rapid (within 30 min) increase in leukocyte adhesion to and migration across venules [175,189]. Moreover, basal leukocyte adhesion is increased in eNOS-deficient mice [176]. Based on these in vivo studies, it has been proposed that eNOS-derived NO is anti-inflammatory with respect to leukocyte adhesion to endothelium, presumably because of its ability to neutralize superoxide [188,190].

Qualitatively consistent with in vivo studies, pharmacologic inhibition of NOS in endothelial cell monolayers results in an intracellular oxidant stress and increased adhesivity for PMN and monocytes [191,192]. However, although the oxidant stress is noted within 30 min after inhibition of NOS, the increase in PMN or monocyte adhesion is not apparent until 2–4 h later and is not as robust as that noted in vivo. If endothelial cells are cocultured with mast cells, PMN adhesion to endothelial cells increases within 30 min after blockade of NO, a response more consistent with in vivo studies [158,175]. The increase in PMN–endothelial cell interactions was attributed to mast cell activation (e.g., oxidant stress, PAF generation). Because the PMN adhesion to endothelium in the coculture setting was prevented by extracellular SOD, it was proposed that limiting NO bioavailability resulted in an increase in endothelial production of ROS, which activated the adjacent mast cells [158].

Challenge of cardiac myocytes with A/R (sterile stress) results in an oxidant stress (linked to increased H_2O_2 production) and induction of a proinflammatory phenotype, i.e., it promotes PMN TEM [186,193]. Similarly, macrophage activation during sepsis (infectious stress) can induce oxidant stress in adjacent endothelial cells with a resultant increase in PMN TEM [170]. This response was dependent on the activation of endothelial cell NADPH oxidase because genetic blockade of NADPH oxidase prevented the PMN TEM [170]. An interesting study in zebrafish larvae indicates that epithelial cells at a sterile wound edge (tail-fin tip amputation) generates a H_2O_2 gradient via Duox (a member of the NADPH oxidase family) to recruit neutrophils to the wound [194]. It was proposed that H_2O_2 could be directly chemotactic for PMNs or that PMN migration toward the wound was due to chemotactic factors released from cells activated by H_2O_2 . Collectively, these observations indicate that interstitial sentinel cells (e.g., myocytes, macrophages) that incur an oxidant stress upon activation can "transfer" ROS (probably as H_2O_2) to other cells (e.g., endothelial cells) and thereby facilitate PMN recruitment.

Endothelial cell capture of leukocytes (Fig. 2B)

To reach the site of injury/infection, circulating leukocytes must first be captured by the endothelium, i.e., adhere to endothelium. The capture of leukocytes by vascular endothelium is a highly regulated process involving sequential engagement of various families of endothelial and leukocyte adhesion molecules [116,117,119]. In general, leukocyteendothelial cell adhesive interactions occur in postcapillary venules. As leukocytes leave the small-caliber capillaries and enter the larger postcapillary venules, they are forced toward the endothelial lining by hydrodynamic forces, thereby allowing them to interact with the endothelium [195]. If an inflammatory focus is present locally, the leukocytes form weak adhesive interactions with the endothelium, referred to as tethering or rolling. These initial adhesive interactions involve P-selectin and E-selectin on the endothelium and PSGL-1 and ESL-1 on neutrophils. Monocytes can roll on endothelial selectins and on the cell adhesion molecule VCAM-1. Leukocyte arrest during rolling is attributed to activation of leukocyte integrins by chemokines or other activators (e.g., PAF) present on the surface of the endothelium. Activated CD11/CD18 integrins on neutrophils interact with ICAM-1 on endothelial cells, whereas activated VLA4 on monocytes interacts with VCAM-1 on endothelial cells. After adhering to the endothelium, leukocytes change shape and crawl to a preferred site for emigration. Although the adhesion cascade described is generally accepted as representative, there are numerous overlaps with respect to the roles of various adhesion molecules in the successful capture of PMN and monocytes, as well as some notable variations related to specific organ systems [116,117,119,196-198].

Leukocyte-endothelial cell interactions occur despite the presence of an extensive filamentous network covering the surface of the endothelium, i.e., the glycocalyx. The glycocalyx decorating the endothelial cell surface consists of proteoglycans and associated glycosaminoglycan (heparan sulfates, chondroitin sulfates, and hyaluronan) side chains [75,76]. The glycocalyx extends for approximately 500 nm from the surface of the endothelial cell. This compares to the 10- to 30-nm length of the endothelial adhesion molecules that are responsible for capturing circulating leukocytes [116,199]. Although it would seem that interactions between leukocyte adhesion molecules and their counterparts on endothelial cells would be severely hindered by the glycocalyx, the steric hindrance is overcome somewhat by the localization of PSGL-1 (ligand for endothelial selectins) on PMN microvilli, which are approximately 300 nm in length [200]. Furthermore, the PMNderived cationic protein myeloperoxidase may interact with negatively charged heparin sulfates and allow PMNs to overcome the electrostatic hindrance of the glycocalyx and penetrate it sufficiently to interact with endothelial adhesion molecules [201]. Finally, leukocyte capture by endothelial cells during inflammation is facilitated by partial degradation of the glycocalyx and shedding of its constituents (e.g., heparan sulfate), thereby exposing relevant adhesion molecules [199,202–204]. The degradation and shedding of the glycocalyx is likely to play an important role in leukocyte recruitment because heparan sulfates localized within this structure appear to regulate the presentation of chemokines on the endothelial cell surface, which is critical for the activation and capture of leukocytes [79,205].

The endothelial lining of the microvasculature is not simply a passive barrier to leukocyte emigration, but an active participant in the inflammatory response [117,206–208]. Endothelial cell activation has been conveniently subdivided into two types of functional responses: a rapid response, which is transcription independent, and a more delayed response, which is transcription dependent [206]. According to this paradigm, the rapid response is generally initiated by ligands of GPCRs, such as histamine, PAF, LTB₄, and CXCL8. Activation of GPCRs results in three major functional events of relevance to endothelial cell–PMN interactions. First, activation of endothelial matrix metalloproteases (MMPs) results in local degradation of the glycocalyx, which facilitates leukocyte–

endothelial cell adhesive interactions [204]. Second, activation of myosin light-chain kinase induces mobilization of Weibel-Palade bodies, which transport P-selectin [206] to the endothelial cell membrane and CXCL8 to the endothelial glycocalyx [209]. Third, activation of phospholipase A₂ ultimately results in the generation of PAF from phosphatidylcholine, which is subsequently secreted and binds to the endothelial glycocalyx [206]. The rapid response is short-lived (<30 min) and serves to tether leukocytes (via P-selectin) and activate them (via PAF and CXCL8). The delayed response is generally elicited by LPS or cytokines, such as IL-1 and TNFα [206]. Activation of their respective receptors results in an intracellular signaling cascade culminating in NF-κB/AP-1-mediated transcription of genes encoding adhesion molecules (E-selectin, ICAM-1, VCAM-1) and chemokines (e.g., CXCL8). The adhesion molecules are expressed on the endothelial cell membrane, whereas the chemokines are bound to the glycocalyx. The delayed response is long-lived (> 12–24 h) and allows for further activation of neutrophils and monocytes (via CXCL8 for neutrophils and CCL2 for monocytes) and promotes firm adhesion to the endothelium (via ICAM-1 for neutrophils and VCAM-1 for monocytes).

During inflammation, the rapid and delayed functional responses of activated endothelial cells are not discrete. For example, inflammatory agents that elicit rapid responses can also elicit delayed responses, i.e., histamine can elicit an NF- κ B-dependent increase in VCAM-1 expression [210]. Thus, in vivo, where the interstitial milieu consists of a variety of inflammatory mediators (cytokines/chemokines), the rapid and delayed phases of functional activation of the endothelium most probably occur as a continuum. A good example to illustrate this point is I/R-induced inflammation. In an in vitro model of I/R (A/R) in which only endothelial cell monolayers are exposed to A/R, there are two phases of PMN adhesion to the monolayers, a rapid phase occurring within 30 min (NF- κ B independent) and a delayed phase occurring at 120 min (NF- κ B dependent) [211–213]. However, in vivo the two phases of I/R-induced leukocyte adhesion to postcapillary venules are not as distinct, but appear to be a continuum. Leukocyte adhesion gradually increases over the first 1–2 h and subsequently increases more rapidly over the next 2 h [214].

Role of ROS and NO (Fig. 2B)

Under basal conditions, endothelial cell ROS production is generally attributed to superoxide generation during mitochondrial respiration [215], although NADPH oxidase may also contribute [216]. No appreciable oxidant stress is incurred under basal conditions, because ROS accumulation is limited by the scavenging ability of endogenous antioxidant molecules (e.g., SOD, glutathione, catalase) [217]. However, the endothelial cell activation that accompanies inflammation generally results in a large increase in ROS production that has been largely attributed to cytosolic and/or membrane-bound enzymes such as NADPH oxidase and xanthine oxidase [218]. Of these, NADPH oxidase is considered to be the most critical [167,218–220], in part because ROS production by NADPH oxidase can influence other ROS-generating sources, such as mitochondria (oxidative damage) and xanthine oxidase (oxidative activation), to enhance their production of ROS [167,218,221]. Although enzymatic generation of ROS usually involves an initial production of superoxide, it is rapidly converted to H_2O_2 , which is the most important ROS involved in intracellular signaling [215].

Activation of endothelial cells with some GPCR ligands results in an intracellular oxidant stress leading to rapid conversion to a proinflammatory phenotype [220]. For example, challenge of endothelial cells with thrombin or CXCL8 results in a rapid (30–60 min), but transient (<2 h), increase in ROS production, which is mediated by NADPH oxidase [222,223]. The thrombin-induced ROS production is associated with P-selectin expression (rapid functional response) [223]. Activation of endothelial cells with either LPS or TNFa also results in a rapid (30 min) generation of ROS in endothelial cells, which is primarily

dependent on NADPH oxidase and to a lesser extent mitochondria; xanthine oxidase and NOS (uncoupled) do not seem to be involved [164,224]. The LPS-induced ROS production by endothelial cells is more robust and prolonged than noted with the GPCR ligand CXCL8 [222]. The increase in oxidant stress induced by LPS and TNFa has been implicated in NF- κ B-mediated transcription of adhesion molecules and chemokines that are involved in leukocyte capture by endothelial cells (delayed functional response) [146,206]. Collectively, these observations support a continuum of endothelial cell activation characterized by the induction of an initial oxidant-mediated (NF- κ B-independent) proadhesive phenotype, followed by a subsequent, more prolonged (NF- κ B-dependent) response.

As mentioned above, the source and roles of ROS in mediating the rapid and delayed activation of endothelial cells may not be readily differentiated during an inflammatory response. For example, histamine (GPCR ligand) induces a ROS-mediated (NADPH oxidase-derived), NF-xB-dependent increase in endothelial VCAM-1 mRNA within 60 min [210]. Conversely, LPS-induced ROS production is mediated by a CXCL8/NADPH oxidase pathway [222]. In both instances, endothelial cell ROS play an important role in the induction of the proadhesive phenotype. A more direct (receptor-independent) method of inducing an oxidant stress in endothelial cells is to challenge them with either A/R or H_2O_2 [212,225]. This approach results in an initial (30 min) phase of PMN adhesion to endothelium (oxidant-induced PAF generation) followed by a delayed (4 h) phase of PMN adhesion (e.g., NF-kB-mediated adhesion molecule expression). The sources of ROS after A/R include mitochondria, xanthine oxidase, and NADPH oxidase [212,226]. Once generated, ROS can be self-perpetuating. For example, the H_2O_2 generated via NADPH oxidase can further activate the enzyme resulting in propagation of the H₂O₂ signal [167,221]. Left unchecked, such a "feed-forward" mechanism would be detrimental to cell viability. However, increases in endothelial oxidant production result in increases in the production of antioxidant enzymes (e.g., SOD) via NF- κ B [225] and Nrf2 [216]. The antioxidants minimize the oxidant stress and blunt the conversion to a proadhesive phenotype.

GPCR activation can also generate NO within endothelial cells. An interesting example is the bradykinin receptors, B2R (constitutive) and B1R (inducible) [227]. B2R activation results in eNOS activation and the transient production of NO. The role of NO in the development of a proadhesive phenotype in endothelial cells is unclear. As mentioned above, inhibition of endogenous NO production results in an NF-kB-mediated increase in endothelial surface expression of adhesion molecules and capture of leukocytes [192]. However, eNOS deficiency, per se, does not affect cytokine-induced increases in adhesion molecule expression or leukocyte adhesion to the endothelium [228]. It has been proposed that eNOS-derived NO, per se, does not directly modulate endothelial cell conversion to a proadhesive phenotype, but rather serves to dampen the impact of ROS by interacting with superoxide and preventing further ROS generation. In an oxidative environment eNOS can be "uncoupled," i.e., converted from an NO-producing enzyme to a ROS-producing one [190,229]. The uncoupling of eNOS is generally attributed to a deficiency in BH₄ due to its oxidation by NADPH oxidase-derived ROS. However, the relative importance of BH₄ levels on eNOS uncoupling has been questioned [230]. Regardless, eNOS uncoupling during oxidative stress would be predicted to exacerbate endothelial cell activation (conversion to a proinflammatory phenotype).

Endothelial NADPH oxidase-derived ROS can induce the expression of iNOS [231] as can activation of the B2R on endothelial cells [227]. How this may influence the overall activation of the endothelium is not entirely clear. Although iNOS uncoupling can occur in an oxidative environment, iNOS activity is relatively resistant to H₂O₂ in this respect [232]. Alternatively, the high output of NO from iNOS may serve to inhibit NADPH oxidase

activity via S-nitrosylation of p47phox, thereby inhibiting ROS production by NADPH oxidase [174] and limiting endothelial cell activation. Clearly, the precise role of NO generated via either eNOS or iNOS in the capture of leukocytes by endothelium remains unclear and additional work is warranted.

Both ROS and NO can have an impact on the integrity of the glycocalyx. In an I/R model of inflammation, ROS generated by endothelial cell membrane-bound xanthine oxidase was implicated in the local degradation of the glycocalyx [233]. Furthermore, exogenous administration of aqueous NO has been shown to prevent the shedding of the endothelial glycocalyx induced by I/R [234]. A causal link between either ROS or NO and glycocalyx integrity and leukocyte adhesion to endothelium seems likely [235]; however, it has not been firmly established.

Endothelial regulation of leukocyte transendothelial migration

Once captured by ICAM-1 and VCAM-1, the leukocyte adhesive interactions are strengthened by enrichment (clustering) of these adhesion molecules at the base of the adherent leukocytes, forming "docking structures" [118,208]. In addition, adhesion molecules implicated in TEM, such as PECAM-1 and CD99, are recruited to the docking structures via membrane recycling that involves vesicle-like trafficking to the site of leukocyte adhesion [118,208,236]. Under some experimental conditions (e.g., excessive activation of leukocytes [118]), endothelial cell projections have been observed enveloping the adherent leukocyte [237,238]. Leukocyte TEM can occur via either a paracellular or a transcellular pathway [116,118,119,196,239]. Regardless of the pathway utilized, it is currently accepted that active participation of the endothelium is critical for leukocyte TEM [207,240–242]. Although the membrane dynamics and adhesion molecules involved may be very similar for the para- and transcellular pathways [118,236,243], the endothelial cell signaling mechanisms involved in leukocyte TEM have been more extensively characterized for the paracellular pathway.

An obvious assumption regarding paracellular TEM is that it requires separation of adjoining endothelial cells. There are two major interendothelial adhesion junctions (IEJs) that directly interact with the actin cytoskeleton to maintain endothelial barrier integrity, i.e., adherens and tight junctions. The adherens junction (AJs) complex consists of vascular endothelial cadherin (VE-cadherin), whose extracellular domain forms homotypic adhesion interactions with VE-cadherin on adjacent endothelial cells [244]. The intracellular domain forms complexes with α -, β -, and γ -catenin, which allows for an interaction with the actin cytoskeleton. Within tight junctions (TJs) occludins/claudins form homotypic bonds with their counterparts on adjacent endothelial cells [240,245]. Their cytoplasmic domains form complexes with ZO-1/ZO-2 to allow for interaction with actin. The stability of the AJ and TJ complexes is primarily regulated by the balance of activities of protein tyrosine kinases (PTKs) and protein tyrosine phosphatases (PTPs) [246]. In general, phosphorylation of AJ and TJ components loosens junctions, whereas dephosphorylation stabilizes the junctions. For example, the PTP, VE-PTP, is associated with VE-cadherin and keeps the cadherin in a dephosphorylated state, favoring AJ stability [244]. By contrast, increased activity of PTKs can phosphorylate components of both adherent and tight junctions, leading to their disassembly, e.g., Rho kinase can phosphorylate occludin [247] and pyk2 can phosphorylate β-catenins [248].

Endothelial cell-associated MMPs have been shown to play an important role in leukocyte TEM, i.e., inhibition of MMP activity blunts leukocyte TEM [241,249,250]. Because MMPs are capable of degrading both VE-cadherin [251] and occludin [252], as well as inducing gap formation between endothelial cells, it has been proposed that endothelial cell MMPs contribute to the disassembly of the adherens and tight junctions during leukocyte

paracellular migration [241]. However, the role of MMPs in loosening IEJs is controversial. VE-cadherin is generally believed to be internalized rather than degraded [118] and the role of MMPs in the degradation of occludin is equivocal [250,253]. Further, the direct addition of MMP-2 to naïve (nonactivated) endothelial cells does not induce occludin cleavage or gap formation [252]. Interestingly, inhibition of MMPs has been shown to prevent formyl methionine leucine phenylalanine (fMLP)-induced (1) degradation of the glycocalyx and (2) PMN adhesive interactions with endothelium [204]. The possibility that endothelial-derived MMPs may modulate leukocyte TEM via localized degradation of the glycocalyx warrants further attention. It is conceivable that MMPs facilitate leukocyte migration via degradation of key components of both IEJs and the glycocalyx.

Role of ROS and NO

There is evidence that ROS and/or NO may be involved in the signaling of ICAM-1- and VCAM-1-dependent leukocyte TEM across paracellular junctions. Ligation of ICAM-1 on cytokine-activated endothelial cells with PMNs or antibodies results in endothelial cytoskeleton reorganization, an effect attributed to ROS generation by xanthine oxidase [254]. Ligation of endothelial cell ICAM-1 on nonactivated endothelium with leukocytes or antibodies has also been shown to activate eNOS (phosphorylation) and generate NO [255]. Pharmacologic blockade of eNOS or sGC, but not xanthine oxidase nor NADPH oxidase, inhibits leukocyte TEM. Blockade of eNOS also prevents VE-cadherin phosphorylation. Based on these and previous studies [256], it is proposed that eNOS-derived NO is critical for VE-cadherin phosphorylation and leukocyte TEM. Because of the paucity of information on the role of ROS and NO in ICAM-1 signaling that leads to leukocyte TEM, it is difficult to reconcile with any certainty why some studies implicate ROS and others support a role for NO. It has been proposed that ligation of ICAM-1 on nonactivated endothelium may favor NO generation, whereas ligation of ICAM-1 on cytokine endothelium would favor ROS generation [255].

VCAM-1 engagement by leukocytes or antibodies results in activation of NADPH oxidase and the resultant ROS generation is believed to facilitate leukocyte paracellular TEM via two potential mechanisms: phosphorylation of VE-cadherin [240,245] and activation of MMPs [241,242]. The current consensus is that VCAM-1 clustering at the docking structure results in Rac1 activation, which leads to activation of NADPH oxidase and the generation of ROS [118,240,245]. Other sources of ROS do not seem to be involved because only blockade of NADPH oxidase interferes with leukocyte TEM; blockade of xanthine oxidase or NOS does not [242].

NADPH oxidase-derived ROS are believed to activate the PTKs and inhibit the PTPs, thereby altering the PTK/PTP balance in favor of PTKs and facilitating VE-cadherin/catenin phosphorylation [240]. For example, in the context of leukocyte TEM, NADPH oxidase-derived ROS has been shown to activate pyk2 (kinase), which subsequently phosphorylates β-catenin and results in the disassembly of the VE-cadherin/catenin complex [248]. Furthermore, ROS generation by VCAM-1 ligation can activate kinase cascades (e.g., PKC/Src kinase pathway) [257,258] leading to VE-cadherin phosphorylation [259]. NADPH oxidase-derived ROS can also activate a phosphatase, PTP1B, the activity of which seems to be critical for leukocyte TEM [260]. The activation of PTP1B is a result of ROS-induced activation of the kinase PKCa. This observation underscores the complexity of ROS-mediated modulation of the PTK/PTP balance with respect to leukocyte TEM.

The enhanced production of NADPH oxidase-derived ROS elicited by VCAM-1 ligation has also been implicated in the activation of endothelial-cell-associated MMP-2 and MMP-9 [249]. A comparable level of MMP activation results after endothelial cell exposure to H_2O_2 , at concentrations detected after VCAM-1 ligation. MMP activation occurs within

minutes and pharmacologic blockade of MMP activity substantially blunts leukocyte TEM. Components of adherens and tight junction complexes are considered the likely targets of the activated MMP [241,251]. As discussed above, NADPH oxidase-derived ROS have also been implicated in the disassembly of endothelial cell junctions via phosphorylation-mediated events. The question therefore arises as to whether phosphorylation of junction components and MMP-mediated degradation of these components occurs simultaneously to ensure gap formation and facilitate leukocyte TEM. One possibility is that phosphorylation events occur with a mild inflammatory stimulus and as stimulus intensity increases proteolytic events begin to contribute. This possibility is supported by the observation that low concentrations of H_2O_2 induce occludin phosphorylation, whereas higher concentrations result in occludin proteolysis [252].

Vascular permeability

Under resting conditions, the endothelial lining of the microvasculature is fairly permeable to small-molecule solutes (e.g., glucose, <3-nm radius) yet it significantly restricts the movement of plasma proteins (e.g., albumin; \approx 6-nm radius). Transendothelial movement of protein and fluid can be envisaged as involving either paracellular (between cells) or transcellular (through cells) pathways or some combination of both [261–263]. During inflammation, chemical cues released by sentinel cells (e.g., macrophages, mast cells) activate endothelial cell signaling pathways, which target structural elements (e.g., actin/ myosin) that regulate vascular permeability. Widening of the interendothelial junctions and frank separation of endothelial cells (gaps) results in microvascular protein (and fluid) leakage into the interstitium (edema). Alterations in other endothelial structural elements (e.g., intracellular vesicles, glycocalyx, basement membrane) can also contribute to the barrier dysfunction that accompanies inflammation [261–263].

Structural determinants: endothelium

Ultramicroscopic studies (e.g., electron microscopy) have identified several structural elements in the endothelial lining of the microcirculation that can potentially serve as paracellular pathways for solute and fluid exchange [264]. Interendothelial junctions consist of protein complexes that couple adjacent endothelial cells and maintain them in close apposition. Within the IEJs there are two major complexes relevant to paracellular permeability: TJs and AJs. The paracellular width imposed by the adhesive interactions of AJ complexes is \approx 15–20 nm, whereas that imparted by TJ complexes is only \approx 5 nm [265,266], TJs being further subdivided into closed (<3 nm) or open (>3 nm) [266]. The density (and/or ratio) of TJs and AJs in the IEJs seems to determine their restrictive properties [261]. In most vascular beds, water and small hydrophilic solutes (<3 nm in diameter) are believed to freely traverse the capillary endothelium via IEJs, the exception being the brain, where TJ density is relatively high and their width may be <1 nm (blood– brain barrier) [261,267,268]. Within a vascular bed, the venular end of the microcirculation has a lower TJ/AJ ratio and the TJ are less well developed than at the arterial end. Furthermore, in regions where three venular endothelial cells converge (tricellular region) the IEJ width can be as great as 30 nm [269]. Because the diameter of albumin is $\approx 6-7$ nm, the venular end of the microcirculation is the major site of albumin leakage[1,270,271].

At the molecular level, TJs contain occludin and claudin, whose extracellular domains allow for homotypic interactions between endothelial cells, whereas within the AJs the extracellular domains of VE-cadherin allow for such interactions [270,272,273]. Their respective intracellular domains interact with proteins (e.g., VE-cadherin with catenins, occludin with ZO-1) that create structural/functional linkages with the actin cytoskeleton (e.g., cortical actin). Loss of function studies indicate that (1) claudin-5 plays a role in permselectivity of brain capillaries to small molecules (<0.8 kDa) [274] and (2) VE-cadherin

is important in restricting albumin leakage from heart and lung microvessels [275]. Although TJs and AJs are generally considered independent entities there is significant "cross talk" between these IEJ complexes, which could have an impact on endothelial barrier function [276–279].

In addition to the paracellular exchange pathway, macromolecular transport can occur via a transcellular pathway. Transcellular exchange involves specialized endothelial organelles such as fenestrae or vesicle-like structures (e.g., caveolae) [271,280-283]. Fenestrae (20-30 nm) are generally covered with a semipermeable diaphragm, except in the liver, where they are not [261,264,284]. Fenestrae appear to allow the free and rapid passage of water and small solutes, yet their diaphragms offer restriction to the movement of proteins (behaving as 5-nm filters) [285]. Vesicles or caveolae (20-30 nm) associated with abluminal and luminal endothelial membranes have stomata that are generally covered by a diaphragm. It has been proposed that the caveolae can "shuttle" engulfed material (e.g., proteins) across the endothelium [261,264,286]. Alternatively, the vesicles may coalesce, forming transendothelial channels through which plasma constituents may traverse the endothelium. A unique potential transendothelial channel, the vesiculovacuolar organelle (VVO), has also been noted in close proximity to IEJs, predominantly in postcapillary venules [261,264,280]. The VVOs are grape-like clusters of vesicle-like structures (20–30 nm or multiples thereof) linked to one another via diaphragmed stomata and generally extending from the abluminal to the luminal aspect of the endothelium.

Although there is general agreement that both paracellular and transcellular pathways contribute to transendothelial protein and fluid exchange, the relative contributions of the two pathways are controversial. This issue may prove to be difficult to resolve, because there are indications that there may be structural and functional linkages between the two pathways. For example, the frequency of vesicle-like structures (e.g., VVOs) is much higher in the vicinity of IEJs and frequently the VVOs coalesce with the IEJs at various points [286]. Furthermore, caveolin-1 protein levels can influence the relative distribution and/or density of molecular constituents of IEJs [287]. Indeed, an attempt has been made to address a similar controversy regarding leukocyte movement across endothelium (paracellular vs transcellular) by proposing that there is an interaction between IEJ components with vesicle-like structures [236,288].

Structural determinants: glycocalyx and basement membrane

Endothelial cell-associated structures on both the luminal and the abluminal aspect of the cell have also been implicated in the regulation of vascular permeability [199]. The luminal surface of the endothelium is covered by a glycocalyx (200–500 nm in thickness) consisting of proteoglycans with GAG side chains (e.g., heparan sulfates) [76,289]. Enzymatic degradation of the glycocalyx components reduces its thickness and increases transendothelial albumin flux [290]. Conversely, stabilizing the glycocalyx (e.g., angiopoietin-1) decreases albumin permeability [291]. The overall negative charge of the GAGs is believed to impose a significant barrier to protein movement through the glycocalyx, but poses little hindrance to the movement of water [283,290].

The basement membrane (BM) adjacent to the abluminal aspect of the endothelium may also contribute to the microvascular permeability. Although the BM contains GAGs, in addition to collagen IV and laminin [283,292], the density of negative charges is too low to confer any significant charge selectivity with respect to albumin movement [293]. Overall, the limited amount of studies on the permeability properties of the BM suggest that it has minimal impact on protein flux across the microvasculature [283].

Functional assessments

Various in vivo and in vitro approaches have been used to assess the restrictive properties of the endothelial barrier [294–297]. In vivo approaches include whole-organ estimates of the osmotic reflection coefficient for plasma proteins (lymphatic flux analysis), microvascular bed protein leakage, in situ single-vessel estimates of hydraulic conductivity (Lp) or solute permeability (Ps), and discrete estimates (e.g., electron microscopy (EM)) of transcapillary protein movement. In vitro approaches have involved ex vivo single-vessel estimates of Ps or an assessment of endothelial cell monolayer Lp, electrical resistance (TEER), or transendothelial movement of protein. All of these approaches have their advantages and limitations. In vivo approaches allow for the assessment of barrier integrity in the context of a physiologic milieu, which also includes the contribution of various accessory structures (glycocalyx) and cells (e.g., leukocytes) [294,297]. However, the possibility that changes in solute flux may be a result of convective (filtration-dependent) and/or diffusive (surface area-dependent) movement, rather than alterations in the restrictive properties of the endothelium, must be taken into consideration. In vitro approaches allow for the assessment of various intracellular signaling pathways and their impact on the endothelial structures relevant to microvascular protein exchange under rigorously controlled conditions [294]. However, endothelial cell monolayers do not completely recapitulate the structural and functional properties of microvessels in vivo [294,296,298], i.e., endothelial monolayer permeability is generally higher than postcapillary permeability. Despite their limitations, both in vivo and in vitro approaches have yielded unique, yet complementary, information that has provided the basis for current concepts regarding the regulation of microvascular permeability.

Basal vascular permeability: role of ROS and NO

"Loss-of-function" approaches (pharmacologic and genetic blockade) have implicated NO in the regulation of basal vascular permeability. Pharmacologic blockade of NO production (exogenous and endogenous NOS inhibitors) increases basal microvascular permeability in a variety of vascular beds in vivo [299-303]. In addition to increasing vascular protein leakage, NO inhibitors also promote leukocyte- endothelial cell adhesion [175,300], and when the leukocyte adhesion is prevented (e.g., by immunoblockade of adhesion glycoproteins) the increased vascular protein leakage is also prevented [175,300]. However, efforts to "flush out" leukocytes from the microvasculature (e.g., blood-free perfusion of microvessels) have yielded equivocal results; blockade of NOS either did [302,304] or did not increase permeability [301,303]. This inconsistency may reflect the influence of perivascular interstitial cells on the vascular permeability response to alterations in NO bioavailability. For example, in blood-free preparations, the increase in albumin leakage and associated endothelial actin reorganization induced by NOS inhibitors are associated with mast cell degranulation [304]. Indeed, a role for both circulating leukocytes and interstitial mast cells has been proposed in modulating basal vascular permeability [1,305]. Consequently, although local inhibition of NOS in a vascular bed can increase microvascular permeability, it is difficult to differentiate the direct effects of NOS inhibitors on the endothelium per se from those resulting from inhibition of NO production by local circulating or interstitial accessory cells.

Time-course studies in blood-perfused mesenteric preparations have revealed an early (leukocyte-independent) and late (leukocyte- dependent) phase of albumin leakage in response to an NO inhibitor [300]. The late phase of protein leakage was directly related to the extent of leukocyte–endothelial adhesive interactions (adhesion and emigration). The early phase was attributed to a direct action on the endothelial cell, which is consistent with reports that NOS inhibitors can increase the flux of macromolecules across isolated glomeruli [306,307] or endothelial cell monolayers in vitro [308–310]. The endothelial

barrier dysfunction detected in monolayer preparations is associated with the formation of stress fibers and disruption of adherens junctions [310]. In both in vivo and in vitro models, a cyclic GMP analogue can reverse the functional/structural effects of NOS inhibition [300,310]. This observation implicates an NO/cGMP pathway in microvascular barrier stabilization under basal conditions.

Genetic blockade of eNOS has also yielded inconclusive results regarding the role of NO in the regulation of vascular permeability. In one study, basal albumin permeability was increased in multiple vascular beds of eNOS-deficient mice, a result confirmed in wild-type mice after long-term treatment with an NOS inhibitor [302]. The increase in transvascular albumin flux was attributed to open IEJs in all vascular beds examined, i.e., lung, heart, gut, and muscle. Transendothelial trafficking of caveolae was not affected. However, in another study it was noted that basal vascular albumin flux in the mesentery does not differ between eNOS and wild-type mice [311]. Mice deficient (heterozygous) in DDAH (dimethylarginine dimethylaminohydrolase) exhibit an increased basal albumin leak across pulmonary capillaries [310]. Because DDAHs metabolize the endogenous NOS inhibitor asymmetric methylarginine (ADMA) [312], this finding was attributed to inhibition of NOS. Furthermore, endothelial cell monolayers derived from DDAH1-deficient mice also exhibited an enhanced permeability, which could be restored by a cGMP analogue [310]. Collectively, the genetic blockade approaches targeting eNOS provide support for the contention that the NO/cGMP pathway plays a role in stabilizing microvascular permeability in the basal state.

By contrast, genetic approaches designed to enhance eNOS activity suggest that eNOSderived NO may destabilize the endothelial barrier. Under basal conditions, most of the intracellular pool of eNOS is associated with the scaffolding domain of caveolin-1, which prevents eNOS activation and function [313,314]. Therefore, it would be predicted that removal of this inhibitory function of caveolin-1 would result in increased NO production and a strengthening of the endothelial barrier against protein extravasation. However, caveolin-1- deficient mice are "hyperpermeable" as evidenced by an enhanced escape of plasma protein from the circulation and a corresponding accumulation of protein in a variety of organs [315]. It is corrected, however, by an inhibitor of NOS activity. The hyperpermeable state noted in caveolin-1-deficient animals was attributed to loosening of endothelial cells from the basement membrane and an increased permeability of IEJs [315]. Downregulation of caveolin-1 (siRNA for caveolin-1) in endothelial cell monolayers results in loss of both TJ (occludin and ZO-1) and AJ (VE-cadherin and catenin) proteins from the IEJ [316]. Another approach resulting in enhanced eNOS activity and NO production by endothelial cells is deletion of CEACAM-1 [317]. CEACAM-1-deficient mice are also hyperpermeable, as evidenced by increased basal protein accumulation in various tissues and ultrastructural evidence of disorganized IEJs.

The discordant results regarding the role of eNOS-derived NO on basal microvascular permeability are somewhat paradoxical. It has been suggested that in in vivo studies alterations in hemodynamic parameters (e.g., capillary surface area, hydrostatic pressure) can influence transvascular protein movement [295–297,318]. Of note is the argument that the enhanced vascular protein leakage noted in caveolin-1-deficient mice may not be due to alterations in microvascular permeability per se, but rather due to the NO-induced vasodilation. In this scenario, the vasodilation results in an increase in capillary hydrostatic pressure, which, in turn, simply increases the filtration of plasma proteins across an intact endothelial barrier [319,320]. Whereas hemodynamic alterations may be a confounding variable in states that increase endogenous NO production, they are unlikely to account for the increased vascular permeability noted during conditions that inhibit NO production [299].

Less attention has been devoted to the role of ROS in regulating basal endothelial barrier function. There is no published information on the effects of blockade of production or bioactivity of endogenous oxidants on basal microvascular permeability, as is the case for endogenous NO (e.g., NOS inhibitors). However, there is considerable evidence that exogenous oxidants increase permeability [321,322]. Based on the ability of superoxide to interact rapidly with NO, the overall consensus is that, in the basal state, the hyperpermeability associated with oxidant production is neutralized by the continuous endogenous NO production. This contention is supported by the following lines of evidence. In blood-perfused mesenteric venules, the early (leukocyte-independent) phase of albumin leakage induced by inhibition of endogenous NO production is associated with an increase in perivenular oxidant stress, which is blunted, along with albumin leakage, by treatment with antioxidants [189]. In in vitro studies, enzymatically generated superoxide can increase glomerular permeability to albumin, an effect prevented by an NO donor [306]. Finally, the increase in permeability of isolated glomeruli [306,307] or endothelial monolayers [308] induced by NO inhibitors can be rescued by inhibitors of NADPH oxidase or antioxidants.

Increased vascular permeability

Structural basis for hyperpermeability states-With respect to the paracellular pathway, disruption of IEJ complexes and widening of the interendothelial junctions may account for the enhanced vascular protein leakage in hyperpermeability states [261,262]. In general, alterations of the local PTK/PTP ratio in favor of PTKs promote phosphorylation of junctional proteins leading to their internalization and ultimately disassembly of the junctional complex [272,273]. In addition, the cortical actin adjacent to the IEJ is destabilized (F-actin converted to G-actin) [323], decreasing peripheral tension [324]. Finally, endothelial actomyosin-mediated contraction and central tension development pull the IEJ apart, creating "gaps" [6,324,325]. Actomyosin-mediated contraction is dependent on myosin light chain (MLC) phosphorylation, which, in turn, is regulated by the MLC kinase/MLC phosphatase balance [325]. In some hyperpermeability states, alterations in junctional complexes occur independent of alterations in cytoskeletal elements, i.e., blockade of actin/myosin interactions does not affect the disassembly of IEJ [296] or even gap formation [323]. It has been proposed that weak stimulation leads to junctional disruption of IEJ and stronger stimulation reinforces this response by initiating actinmyosin-based endothelial cell retraction [270,273].

With respect to the contribution of the transcellular pathway to hyperpermeability states, loss of diaphragms and widening of the VVO stomata [286] and an increase in the diameter of nondiaphragmed fenestrae have been implicated [326]. Little is known about the mechanisms involved, but it has been proposed that actin–myosin contractions contribute to these ultrastructural readjustments [280,286,327]. Increased vesicle trafficking has also been proposed to explain the increased albumin leakage that occurs during an inflammatory response [328].

Cellular and chemical mediators—Specific signals generated during injury/infection (PAMPs and DAMPs) can activate resident interstitial immune cells to initiate the inflammatory response. Perivascular mast cells have been implicated in the microvascular protein leakage associated with the inflammatory response induced by either infectious [329,330] or noninfectious insults [331,332]. Pharmacologic stabilization or genetic deletion of mast cells has been shown to blunt the leukocyte–endothelial cell adhesive interactions and vascular protein leakage in several experimental models of inflammation [1,177,333]. Furthermore pharmacologic activation of mast cells promotes leukocyte adhesion and protein leakage [177,334]. Analogous experimental approaches have also implicated resident macrophages in the microvascular protein leakage associated with leukocyte

invasion of inflamed tissue. For example, pharmacologic depletion of macrophages or blockade of macrophage-derived chemokines reduces PMN emigration and vascular protein leakage [335,336].

Impact of circulating immune cells—Pharmacologic or genetic blockade of neutrophil–endothelial adhesive interactions can blunt the leakage response in experimental models of inflammation [1]. Furthermore, the magnitude of the albumin leakage is directly correlated with the number of leukocytes adhering to or migrating across the endothelium in vivo (mesenteric venules) [300,332] and in vitro (endothelial monolayers) [337]. PMN-mediated increases in vascular permeability occur even though the underlying endothelium extends projections to envelop the adherent/migrating PMNs, forming endothelial domes; the leakage response is presumably due to the transfer of entrapped plasma proteins within the "dome" [237,325].

The PMN-induced leakage response occurs whether PMN are utilizing the paracellular or the transcellular pathway [237]. There is evidence that PMN-derived proteases (e.g., elastase) can disrupt IEJ complexes and induce endothelial cell retraction, thereby increasing paracellular permeability [294,338]. Alternatively, PMN ligation of endothelial adhesion molecules can elicit both actin–myosin-based endothelial cell tension development and IEJ disassembly, resulting in widening of the IEJ or gap formation [245,325]. Endothelial cell adhesion molecule-dependent signaling can also increase transendothelial albumin movement via a caveola transport pathway, without affecting the paracellular pathway [339,340].

While there is a large body of evidence supporting a role for neutrophil-mediated increases in vascular permeability during an inflammatory response [297], there is also evidence indicating endothelial barrier dysfunction can be uncoupled from neutrophil–endothelial cell adhesion [305]. In blood-perfused vascular beds, induction of infectious [329] or sterile [214] inflammation increases both vascular protein leakage and neutrophil–endothelial adhesive interactions [329,332]. However, the protein leakage response can be subdivided into an early and a late phase, the early phase being PMN independent and the later phase being PMN dependent. Furthermore, the increase in transendothelial albumin flux associated with macrophage/mast cell activation is noted in both blood-free preparations in vivo and endothelial monolayers sans neutrophils in vitro [304,341–344]. Finally, histamine, a mast-cell-derived mediator, can increase vascular albumin leak without affecting leukocyte–endothelial adhesive interactions [345]. Collectively, the data are consistent with a direct impact of inflammatory mediators on endothelial permeability, which is further reinforced by PMN–endothelial cell interactions.

Although T-lymphocytes (CD4⁺ and CD8⁺) are classically viewed as components of the adaptive immune system, they have been implicated in vascular responses to acute inflammation (innate immune system) [297]. Genetic and pharmacologic depletion of T-cells appears to blunt leukocyte–endothelial cell adhesion and the associated increase in vascular protein leakage in various models of inflammation [346–348]. Because T-cells can enhance neutrophil-mediated endothelial dysfunction [349], it is generally held that the role of T-cells in inflammation-induced protein leakage is a result of their ability to enhance neutrophil reactivity [297].

Circulating platelets have also been implicated as modulators of vascular permeability. Platelets can ameliorate the hyperpermeability response to inflammatory mediators in vascular beds supplied with a blood-free perfusate [297], but contribute to the hyperpermeability noted in experimental inflammation in vivo (blood perfused) [350]. The protective effect of platelets has been attributed to platelet-derived sphingosine (S1P)

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[297,351,352], whereas their hyperpermeability effects have been attributed to the formation of platelet–leukocyte aggregates. Interestingly, another source of circulating S1P, a substance noted for its stabilizing effect on the endothelial barrier [353], is red blood cells. Vascular albumin leak and the formation of interendothelial gaps are noted in the lungs of mice genetically deficient in plasma S1P, a situation rectified by transfusion with wild-type erythrocytes [354].

Role of ROS and NO—Both xanthine oxidase- and NADPH oxidase-derived ROS have been implicated in the vascular permeability associated with inflammation [355]. In I/R models, both the increase in venular/perivenular oxidant stress and the albumin leakage can be prevented by antioxidants, inhibitors of xanthine oxidase, or mast cell stabilization [356,357]. A proposed scenario is that oxidant production by endothelial xanthine oxidase results in activation of perivenular mast cells that, in turn, contribute to the inflammatory response, i.e., leukocyte recruitment and vascular protein leakage [297]. However, in other models of oxidant stress (systemic hypoxia), mast cells seem to be the primary producers of ROS that elicit the leukocyte recruitment and subsequent vascular protein leakage [358]. Alveolar macrophages (AM), which seem to be critical for I/R- or sepsis-induced pulmonary vascular leakage [183,335], can be activated by oxidant stress and also generate more ROS [359]. ROS generated by LPS-activated AMs have been implicated in endothelial barrier dysfunction. The AM-induced reduction in endothelial monolayer TEER is associated with a shift in the localization of the TJ components (occludin, ZO-1, claudin-5) from cell borders to cytoplasm(not loss thereof), an effect prevented by inhibition of NADPH oxidase [360]. Moreover, the increased transendothelial albumin movement induced by cytokine-activated AMs is abolished when NADPH oxidase-deficient AMs are used in the assay [342]. Circulating leukocytes can also contribute to perivascular oxidant stress and increase microvascular permeability [321,361]. PMN NADPH oxidase has been implicated in the pulmonary hyperpermeability induced by LPS [362].

The use of in vitro approaches utilizing endothelial cell monolayers indicates that endothelial cells can generate enough ROS to mediate increased monolayer permeability in the absence of any auxiliary vascular or perivascular immune cells [321]. Challenge of endothelial cells with anoxia/reoxygenation increases monolayer permeability, a response blocked by antioxidants or inhibitors of xanthine oxidase [294,363]. Direct exposure of endothelial cells to exogenous xanthine/xanthine oxidase or H2O2 can increase monolayer permeability, an effect blocked by intracellular delivery of catalase (conjugated to PECAM-1 antibody) [364]. Many agents (e.g., LPS, histamine, bradykinin, thrombin, VEGF) shown to increase endothelial monolaver permeability also increase endothelial ROS production via NADPH oxidase [57,365-368]. Although in some cases the mediatorinduced increase in monolayer permeability can be ameliorated by blockade of NADPH oxidase or ROS scavenging [364,367,368], in other cases, a cause-effect relationship may not be apparent. For example, thrombin-induced endothelial monolayer permeability is not prevented by intracellular delivery of antioxidant enzymes, catalase, or SOD [364]. Nonetheless, the literature is generally consistent with the view that accelerated intracellular ROS generation is an important underlying cause of the endothelial hyperpermeability induced by inflammatory mediators.

Delineation of the role of NO in the vascular permeability response associated with inflammation has been fraught with controversy [318,321,369]. As mentioned above, the general consensus is that NO serves to stabilize microvascular barrier function under basal conditions. However, during an inflammatory response, there is compelling evidence supporting a role for NO as both a mediator and an inhibitor of endothelial barrier dysfunction. In I/R-induced inflammation, the increase in vascular albumin leakage is associated with a reduction in local levels of NO (nitrate/nitrite levels), and the

hyperpermeability response is diminished by local administration of NO donors [332]. The protective effect of NO donors has been attributed to the ability of NO to interfere with auxiliary cell function, e.g., platelet–neutrophil interactions and/or mast cell degranulation [1,297,334]. Genetic blockade approaches seem to be divided on the role of auxiliary cell iNOS in the hyperpermeability response. LPS-induced pulmonary hyperpermeability is exacerbated in iNOS-deficient mice, a response attributed to macrophage iNOS [370]. In contrast, an in vitro coculture approach indicates that cytokine-activated AMs increase pulmonary endothelial monolayer permeability to albumin, a response not noted with iNOS-deficient AMs [371].

Genetic blockade approaches have also implicated eNOS in various in vivo models of acute inflammation. The hyperpermeability response observed in carrageenan- or zymosaninduced acute inflammation is largely blocked in eNOS-deficient mice [372]. Interestingly, the leukocyte extravasation noted in these models was not affected by genetic blockade of eNOS, suggesting an uncoupling between leukocyte adhesion and permeability responses. PAF-induced hyperpermeability in hamster cheek pouch and mesentery are (1) directly related to endothelial generation of NO [373] and (2) blunted in eNOS- but not iNOSdeficient mice [311]. Based on data generated from bone marrow chimeras, a vascular, rather than auxiliary, cell Akt/eNOS pathway has been implicated in the eNOS-induced hyperpermeability associated with acute inflammation in peripheral vascular beds [374]. In the pulmonary vascular bed, however, eNOS does not seem to contribute to the inflammation-induced hyperpermeability. For example, LPS-induced pulmonary hyperpermeability is not altered in eNOS-deficient mice [370]. Furthermore, the hyperpermeability associated with ventilator-induced lung injury is exaggerated in eNOSdeficient mice, but prevented in iNOS-deficient mice, with the latter response attributed to increased eNOS activity [375,376].

Some interesting explanations have been proposed to address the contradictory observations regarding the role of NO in modulating vascular permeability in vivo. The effects of NO may be tissue specific. For example, PAF-induced hyperpermeability in hamster cheek pouch is associated with enhanced NO generation, whereas the hyperpermeability response in lung is associated with a reduction in NO generation [377]. Furthermore, the PAF-induced hyperpermeability in systemic vascular beds seems to be driven by eNOS-derived NO [311,378], whereas the same response in the lungs seems to be dampened by eNOS-derived NO [376,379]. A second level of complication results from the relative roles of iNOS and eNOS in various experimental models. For example, LPS-induced vascular leakage in the mesentery seems to be mediated by induction of iNOS, and pharmacologic elevation of eNOS and/or prevention of iNOS induction alleviates the hyperpermeability response [380,381]. This phenomenon is consistent with the view that the effects of NO on barrier function may be concentration dependent; low levels serve to stabilize and high levels serve to destabilize [321,377].

In vitro approaches have specifically targeted the role of endothelial-derived NO on monolayer hyperpermeability, again with conflicting results. Exposure of endothelial monolayers to hypoxia induces an oxidant stress and increases monolayer permeability, events prevented by an NO donor and an analogue of cGMP [382,383]. Endothelial monolayer permeability (TEER) induced by LPS or cytokines can also be reversed by NO donors or cGMP analogues [384]. In contrast, challenge of endothelial cell monolayers with histamine or PAF results in increased NO production and monolayer permeability, and the hyperpermeability response is prevented by inhibition or depletion of eNOS [374,385]. Administration of NO (NO donors) has yielded equivocal results with respect to microvascular barrier function. Although NO donors elicit dose-dependent effects on the permeability of endothelial cell monolayers, some donors increase monolayer permeability

[310,386], whereas others decrease monolayer permeability [309,384]; in both instances, the effects of the NO donors involve the cGMP pathway.

The discordant results derived from in vitro models exposed to NO donors may reflect the somewhat artificial situation created by this mode of NO delivery to cells. It is likely that the quality (species) and quantity (rate, duration) of NO produced by intact cells differ from those derived from NO donors [387]. Further, exogenous application of NO donors subjects the entire intracellular environment to NO, whereas endogenous NO generation is likely to be compartmentalized [369]. For example, the PAF-induced increases in vascular permeability are dependent on eNOS internalization (via caveolae) and targeting to cytoplasmic locations [385]. On the other hand, acetylcholine, which does not alter barrier function, triggers internalization and compartmentalization of eNOS to the Golgi region [388]. Finally, the use of cyclic GMP analogues is also not without its own draw-backs. Moderate increases in intracellular cGMP serve to stabilize endothelial monolayer permeability, whereas higher levels diminish barrier function [389].

There is also evidence (albeit limited) that NO may play a role in the postinflammation recovery of endothelial barrier function. For example, the increase in paracellular gap formation and albumin leak induced by inflammatory mediators (e.g., PAF, bradykinin, histamine) is a transient phenomenon, lasting 30 min or less. The resealing of the endothelial barrier (gap closure and decreased protein leak) is inhibited by pharmacologic inhibition of eNOS [390]. S1P has also been implicated in barrier restoration after a hyperpermeability response (resealing of junctions and strengthening of cortical actin) via either endothelial cell receptor activation or an intracellular pathway [351,353,354]. Although S1P is a well-known activator of eNOS [391], the specific linkages between S1P, eNOS, and barrier resealing have not been firmly established.

Endothelial cell signaling (Fig. 3)

The increase in vascular permeability during inflammation requires activation of endothelial cells, such that they can make appropriate ultrastructural alterations favoring solute and fluid exchange. Endothelial activation is initiated either directly (via DAMPs and/or PAMPs) or indirectly (via release of mediators from auxiliary cells). In either case, endothelial activation results in cell signaling, which ultimately leads to an increase in the dimensions of the paracellular pathway, i.e., IEJ disassembly, loosening of cortical actin, and actomyosin contraction leading to interendothelial gap formation. Although endothelial activation can also lead to an increase in the dimensions of the transcellular pathway (VVOs) [280,286,327], less information is available regarding the cell signaling pathways involved in that process.

A multitude of signaling pathways have been implicated in the regulation of endothelial paracellular permeability. These pathways are to a large extent interrelated and a dominant feature is the role of the Rho GTPase family, specifically RhoA, Rac1, and Cdc42 [262,392,393]. In general, endothelial barrier stability is driven by Rac1/Cdc42, and endothelial barrier instability is driven by RhoA. Rac1/Cdc42 recruits IQGAP1 and inhibits its interaction with β -catenin, thereby strengthening AJ interaction with cortical actin; inhibition of Rac1/Cdc42 disassociates IQGAP1 from the complex and destabilizes AJs [393]. Furthermore, Rac1/Cdc42 recruits cortactin to strengthen cortical actin. Finally, Rac1/Cdc42 inhibits RhoA via several downstream targets. In response to ligation of GPCR (e.g., thrombin), PLC generates IP₃ and diacylglycerol (DAG). IP₃ releases Ca²⁺ from the endoplasmic reticulum to activate MLC kinase (MLCK) which, in turn, phosphorylates MLC to promote actomyosin-mediated endothelial contraction. DAG activates a transient receptor potential (TRP) channel to allow Ca²⁺ influx into the cell and activates PKC γ , which in turn activates RhoA. RhoA (via Rho kinase) inhibits MLC phosphatase (MLCP)

activity, thereby further enhancing actomyosin contraction [325]. Finally, Rho kinase inhibits Rac1 activation.

The increase in permeability may be self-limiting or can be resolved by activation of Rac1/Cdc42 and inhibition of RhoA. For example, GPCR ligation by thrombin causes a delayed activation of Rac1/Cdc42 via PKC γ activation of sphingosine kinase and generation of S1P [394]. Cdc42, at the level of the Golgi, enhances transport of junctional proteins toward the membrane [392]. Restabilization of the AJ complex results in further inactivation of RhoA via a VE-cadherin/p120 pathway. In addition, exogenous agents that increase intracellular cAMP (e.g., atrial natriuretic peptide) can inhibit RhoA and activate Rac1 [389,393].

The balance between Rac1/Cdc42 and RhoA dominance represents a generalized framework for understanding some of the fundamental signaling pathways involved in the regulation of endothelial barrier function. However, the in vivo situation is much more complex. For example, RhoA can also stabilize AJ complexes via Dia1, and Rac1 can lead to actomyosin contractions via PAK [393,395,396]. The specific effectors targeted by the Rho kinases can elicit different responses and it has been proposed that the effectors targeted by Rho kinases (e.g., Rac1) may be determined by different guanine nucleotide exchange factors [397].

Role of ROS and NO (Fig. 3)

Exposure of endothelial cells to oxidants (e.g., H_2O_2) or endogenous generation of oxidants in response to endothelial activation (e.g., VEGF) results in disruption of the IEJ, actomyosin contractions, gap formation, and an increase in endothelial permeability [252,321,368,398]. Extracellular H_2O_2 (from auxiliary cells) can convert endothelial xanthine dehydrogenase to xanthine oxidase [399] and can activate NADPH oxidase to generate oxidants intracellularly [167], thereby creating positive feedback loops perpetuating oxidant generation. Because the sole physiologic function of NADPH oxidase is the generation of superoxide (and H_2O_2) and the oxidase can be activated via a variety of inflammatory mediators [219], it has received the greatest attention.

Generation of oxidants by NADPH oxidase is facilitated by Rac1 binding to the enzyme [397]. For example, the H/R-induced increase in NADPH oxidase activity, ROS generation, and endothelial monolayer permeability is abolished in Rac1-deficient cells [400]. Activation of NADPH oxidase by Rac1 seems to be incongruous with the proposed role for Rac1 in endothelial barrier stabilization. Based on studies using thrombin as an agonist, it has been proposed that cytosolic Rac1 is responsible for the barrier-stabilizing effect, whereas membrane-associated Rac1 is involved in NADPH oxidase activation [367]. In VEGF-challenged endothelial cells, NADPH oxidase-derived ROS have been implicated in the phosphorylation of AJ proteins (VE-cadherin, β -catenin) and increased endothelial monolayer permeability [368]. The phosphorylation status of AJ proteins is dependent on the PTK/PTP balance [246]. ROS can either inactivate PTP or activate PTK via oxidation of critical cysteine residues, leading to phosphorylation of VE-cadherin and β-catenin and compromising IEJ integrity [246,398]. A similar mechanism may be operative at the level of actomyosin-mediated contraction, because NADPH oxidase-derived ROS have been implicated in MLC phosphorylation and the increase in endothelial permeability in response to hypoxia [401]. Furthermore, in an in vitro stroke model, the endothelial hyperpermeability induced by the Rac1/NADPH oxidase pathway was attributed to activation of RhoA [402]. The RhoA/Rho kinase may increase endothelial permeability via either phosphorylation (and internalization) of junctional proteins [247] or MLC-induced actomyosin contractions [402], or both. ROS can also activate TRP channels to allow intracellular entry of Ca²⁺ [403], which would also facilitate actomyosin contraction via activation of MLCK and phosphorylation of MLC [325].

Activation of endothelial cells by GPCR ligation or nonreceptor mechanisms (e.g., H_2O_2) leads to Ca^{2+} entry into endothelial cells and binding of Ca^{2+} to calmodulin, as well as activation of various kinases (e.g., PKC, PI3-kinase) [6,314,404,405]. Under basal conditions, eNOS is concentrated in caveolae, where its activity is inhibited by interaction with caveolin-1 and constitutive phosphorylation at Thr95. Upon agonist activation, caveolin-1 is displaced from eNOS and the Thr95 residue is dephosphorylated, whereas Ser1177 is phosphorylated. These events facilitate $Ca^{2+}/calmodulin$ interactions with eNOS resulting in NO generation. The major downstream signaling target is sGC, which increases intracellular levels of cGMP.

The NO/cGMP pathway has been implicated as both a negative and a positive modulator of endothelial barrier integrity [325,406,407]. With respect to its role in hyperpermeability responses, scission of eNOS-containing caveolae and delivery of eNOS to the endothelial cytoplasm via endocytosis seem to be a prerequisite [369,385]. NO can activate MLCK and induce actomyosin contraction via the sGC/cGMP/PKG pathway [325]. In addition, NO can nitrosylate β -catenin (cGMP independent), leading to its disassociation from VE-cadherin and IEJ disruption [408]. By contrast, the eNOS/cGMP pathway has been implicated in the attenuation of the cytokine-induced increase in endothelial monolayer permeability [409]. The NO/cGMP pathway can also ameliorate the thrombin-induced hyperpermeability response by increasing cAMP levels [389]. One possible explanation for the discordant effects of the NO/cGMP pathway is centered on the dose-dependent effects of cGMP on cAMP-hydrolyzing phosphodiesterases, PDE2 and PDE3 [389]. In this scenario, low levels of NO and cGMP inhibit PDE3 activity and increase cAMP levels (enhance barrier stability), whereas high concentrations of NO and cGMP stimulate PDE2 activity and decrease cAMP levels (diminish barrier stability). In addition, it was proposed that any other mediators (e.g., TNF) with the potential to influence endothelial PDE activity could alter the cell's responses to activation of the NO/cGMP pathway.

An issue that has been understudied is the precise linkage between NO and the Rho GTPases. One specific point worth pursuing is the linkage between specific Rho GTPases and isoforms of NOS. There is evidence for a relationship between Rac1 and eNOS [409,410], as well as RhoA (Rho kinase) and iNOS [381]. Additional work is warranted to resolve this issue relative to the endothelial cell signaling mechanisms that regulate endothelial barrier function.

Angiogenesis

One of the consequences of inflammation is tissue hypoxia due to a combination of increased metabolic demand of infiltrating cells (both pathogenic and immune) and decreased O₂ delivery (vascular damage and edema) [411]. Hypoxia is the primary stimulus for the formation of new blood vessels (angiogenesis) to restore O₂ delivery to the affected site [412]. Angiogenesis can be broadly divided into two phases: an initial destabilization of the microvasculature, followed by a restabilization of newly formed microvessels. Destabilization of the endothelium (e.g., pericyte dropout and weakening of the interendothelial junctions) allows for migration of endothelial cells to create new blood vessels. A selected endothelial cell (referred to as a "tip cell") begins to extend lamellipodia and degrade the basement membrane and ECM. The tip cell is followed by proliferating and migrating endothelial cells (referred to as "stalk cells"). Initially, the endothelial sprout of tip and stalk cells is relatively leaky and does not restrict the movement of macromolecules. During the restabilization phase, mural cells (e.g., pericytes) are recruited to the developing sprout and the interendothelial junctions of these neovessels become strengthened and less permeable to macromolecules.

Sentinel cells: signaling pathways (Fig. 4A)

The macrophage is also a pivotal effector cell in the initiation of angiogenesis [122,413,414]. Two major signaling pathways have been implicated in macrophage-induced angiogenesis: the hypoxia-inducible factor (HIF) and NF- κ B pathways [415,416]. Both the HIFa and the NF- κ B pathways can be activated by either inflammatory cytokines or hypoxia, or both [414,416,417]. The HIFa pathway leads to the transcriptional generation of the proangiogenic agent VEGF. The NF- κ B pathway leads to transcription of various cytokines (e.g., TNFa, IL-1 β). Of particular relevance to angiogenesis, NF- κ B binding sites are present on the HIFa promoter and NF- κ B can increase HIFa levels [416]. Thus, activation of both pathways within macrophages can lead to the generation of VEGF and the initiation of angiogenesis.

In the NF- κ B pathway, the IKK complex phosphorylates I κ B and targets it for proteasomal degradation, thereby freeing NF- κ B to translocate to the nucleus and transcribe relevant genes. It is readily accepted that inflammatory mediators (e.g., TNF α , IL-1 β) can activate NF- κ B via the IKK-dependent "canonical pathway" by phosphorylation of serine residues (Ser32 and Ser36) on I κ B [418]. On the other hand, hypoxia can activate NF- κ B via an IKK-independent "atypical pathway" via tyrosine kinase (e.g., c-Src kinase) phosphorylation of tyrosine 42 on I κ B [418].

In the HIF pathway, HIFa forms a heterodimer with HIF β , which can transactivate target genes, e.g., VEGF. The HIF pathway is regulated primarily by prolyl hydroxylase domaincontaining enzymes (PHDs) [415,419,420]. PHDs are members of the 2-oxyglutaratedependent ferrous iron (Fe²⁺) dioxygenase family that hydroxylate HIFa and target it for proteasomal degradation. The activity of PHDs is dependent on the O₂ concentration [415,420]. Under normoxic conditions PHDs are partially active, but any further decreases in local O₂ suppress PHD activity [415]. Thus, under normoxic conditions there is minimal, if any, active HIFa because of proteasomal degradation. However, under hypoxic conditions HIFa is stabilized (not degraded) and is free to interact with HIF β and transcribe relevant genes.

Interestingly, both the HIF and the NF- κ B pathways can be activated by decreasing the activity of PHDs. As mentioned above, PHDs hydroxylate HIFa, thereby routing it to the proteasome [415,419,420]. In the NF- κ B pathway, it has been proposed that PHDs hydroxylate the catalytic subunits of the IKK complex (at a target motif for hydroxylation similar to HIFa), thereby inhibiting I κ B phosphorylation and degradation [416,421]. In an inflammatory/hypoxic milieu, PHD activity is suppressed and both the HIF and the NF- κ B pathways are activated. Hypoxia, per se, is thought to be a stronger activator of the HIF pathway [416]. Thus, it is readily apparent that in an inflammatory/ hypoxic environment robust activation of both pathways would occur. As the inflammation resolves there would be a shift to a predominant role of the HIFa pathway favoring VEGF generation and angiogenesis.

Negative feedback loops exist for these two pathways; the HIF pathway leads to transcriptional increases in PHD levels and the NF- κ B pathway leads to increases in I κ B. Moreover, these two pathways are not mutually exclusive, but rather there is a great deal of cross talk at various levels of the two pathways [416,422–425]. There is also evidence that effective transcription of some genes may even require the binding of both transcription factors to their promoter elements [426]. Thus, although the salient features of the HIFa and NF- κ B pathways of gene expression are depicted in Fig. 4, their relative contributions, interactions, and modulation by additional signaling pathways during angiogenesis are clearly much more complex.

The macrophage is believed to be a key effector cell in the initiation of angiogenesis via generation of VEGF [413,414,427,428]. Macrophages are adapted to functioning in a hypoxic environment and hypoxia alone can result in the induction of an angiogenic phenotype in these cells, e.g., upregulation of the VEGF gene [427]. However, VEGF can be generated not only by macrophages, but also by endothelial cells [429], mural cells [430], or epithelial cells [429,431]. The likelihood that endothelial-derived VEGF via a HIF or NF-κB signaling pathway contributes to the initiation of angiogenesis seems unlikely, because endothelial cells with a PHD deficiency adopt a quiescent stable phenotype rather than the migratory phenotype required for angiogenesis [432]. An additional caveat to consider is whether resident or newly recruited monocytes/macrophages are the major contributors to neovascularization. During hypoxia or wounding, resident macrophage-like cells exposed to hypoxia can generate chemoattractants via the HIF pathway, which can recruit additional monocyte/macrophage populations to the affected site: these recruited accessory cells play an important role in new vessel formation [433,434]. Indeed, macrophages recruited to developing tumors are believed to be the initiators of angiogenesis in that environment [413,428].

Role of ROS and NO (Fig. 4A)

The inflammatory cytokines TNFa and IL-1 β and hypoxia can induce an intracellular oxidant stress via generation of ROS [219,435]. Current consensus holds that hypoxia or an oxidative stress induced by ROS activates the NF- κ B pathway via either the IKK-dependent or an IKK-independent pathway [435]. Similarly, although it is accepted that ROS play an important role in stabilizing HIFa, most probably via a MAP kinase pathway, the signaling pathways may be indirect (e.g., activation of NF- κ B pathway) [436–439].

An interesting integration of ROS-induced activation of both HIFa and NF- κ B centers on the ability of redox-sensitive PHDs to regulate both NF- κ B activation and HIFa stabilization [416,421]. Because inflammation and/or hypoxia can lead to oxidant stress, it is not surprising that ROS have been implicated in the regulation of PHD activity. Mitochondria-derived ROS appear to decrease PHD activity and increase HIFa availability in a variety of cell types [439,440], including macrophages [438]. NADPH oxidase-derived ROS have also been implicated in stabilizing HIFa via inhibition of PHDs [414,441]. Of note, mitochondrial ROS generation may be a prerequisite for NADPH oxidase activation [442–445]. A similar mechanism of action centered on regulation of PHDs has been proposed for the activation of the NF- κ B pathway [416].

As mentioned above PHDs are members of the 2-oxyglutaratedependent Fe²⁺ dioxygenase family, which are highly sensitive to prevailing O₂ [414,415,420]. It is generally believed that ROS exert their effects on PHD activity by oxidizing ferrous to ferric (Fe³⁺) iron [440,441]. Alternatively, ROS can reduce ascorbate levels, thereby increasing the ratio of Fe³⁺/Fe²⁺ [415]. Both of these ROS-mediated effects would render the enzyme inactive and promote stabilization of HIFa and activation of NF- κ B.

NO can also modulate both the HIFα and the NF- κ B pathways [446,447]. Most of the studies addressing this issue have made use of various NO donors. Generally, NO-mediated signaling involves chemical interactions with either transition metals (e.g., Fe-heme of sGC leading to increased cGMP) or protein thiols (e.g., cysteine nitrosylation) [446,448,449]. The latter mechanism seems to be operative in the HIFα and NF- κ B pathway in the context of hypoxia/inflammation. S-nitrosylation is reversible within cells by "denitrosylases" such as *S*-nitrosoglutathione reductase or thioredoxin, thereby making this signaling system analogous to the kinase/phosphorylation system [448,449]. It has been proposed that S-nitrosylation is favored (over sGC activation) at high NO concentrations (micromolar) [446,450]. However, in the presence of an intracellular pool of chelatable iron, S-

nitrosylation reactions have been demonstrated in macrophages even at low concentrations of NO (nanomolar), especially under hypoxic conditions ($O_2 < 1 \mu M$) [451].

NO donors inhibit NF-kB binding to DNA via S-nitrosylation of Cys62 in the p50 subunit of NF- κ B in both cell-free systems and various cells [447,452–454]. In addition, NO can nitrosylate Cys179 in IKK β and, thereby, prevent phosphorylation and subsequent degradation of I κ B [455]. Finally, NO has been shown to directly inhibit NF- κ B activity by nitration of the tyrosine residues (Tyr66 and Tyr152) on p65 [456]. Thus, the available evidence favors inhibition of the NF-kB pathway by NO-mediated S-nitrosylation reactions. With respect to the HIF pathway, S-nitrosylation chemistry favors HIF stabilization. Under normoxic conditions NO can S-nitrosylate HIFa and promote stabilization of HIFa (i.e., Cys533 and possibly others) [457,458] as well as enhancing its transcriptional activity (Cys800) [459]. Finally, NO donors can stabilize hypoxia-induced HIFa by preventing HIF ubiquitination [460], perhaps via S-nitrosylation of the ubiquitin ligase [448]. Of particular relevance to angiogenesis is the observation that S-nitrosylated HIFa binds more avidly to the VEGF promoter region in vitro and increases capillary density in vivo under normoxic conditions [461]. Collectively, the S-nitrosylation chemistry of NO is stimulatory in the HIFa pathway, as opposed to playing an inhibitory role in the NF-kB pathway. The net result of increased NO generation is a general shift in macrophage signaling from pathways supporting inflammation to those favoring angiogenesis.

Another mechanism by which NO can modulate the HIFa pathway is via interactions with mitochondria and subsequent regulation of PHD activity [420,442,446]. In this scenario, NO competes with O₂ for cytochrome c oxidase, leaving more O₂ available for use by PHD and resulting in destabilization of HIF α , particularly under hypoxic conditions [462]. However, the use of NO donors has led to equivocal results, their effects being O2, time, and dose dependent [414,420]. For example, when NO is generated at high concentrations (micromolar) it promotes stabilization of HIFa under both hypoxic and normoxic conditions; when NO is produced at low concentrations (nanomolar) it promotes destruction of HIFa only under hypoxic conditions [419,420,447]. A working hypothesis has been proposed that attempts to integrate the confounding effects of NO donors on the PHD/HIFa pathway [420,463]. The pharmacologic release of large amounts of NO would directly inhibit PHD activity and stabilize HIFa. The inhibitory effect of NO donors on PHD mimics hypoxia, i.e., NO competes with O₂ binding to the Fe²⁺ of PHD. The resultant decrease in proteasomal degradation of HIFa allows for an increased HIFa/HIFβ-mediated transcriptional generation of PHD. When the pharmacologic release of NO wanes the increased pool of PHDs rapidly shuttles HIFa to the degradation pathway, particularly under hypoxic conditions. Thus, because of the negative feedback loop, transient pharmacologic generation of NO initially increases HIF α , followed, at a later time point, by a decrease in HIFa.

Functional destabilization of microvessels (Fig. 4B)

In the adult, angiogenesis usually occurs by sprouting of new vessels from existing functional capillaries or microvessels [432,464,465]. This requires destabilization of the existing microvascular structure, i.e., pericyte dropout, alterations in endothelial cell phenotype, and breakdown of the basement membrane. Initially, an endothelial cell (tip cell) leaves the endothelial lining, penetrates the basement membrane, and invades the interstitium. The tip cell has a specialized phenotype, such as lamellipodia and filopodia to detect environmental cues for appropriately directed migration. The tip cell is followed by migratory/proliferative cells (trunk or stalk cells), which allow for extension of the sprouting vessel and lumen formation [465,466]. VEGF, generated by the HIF pathway, is believed to be the major growth factor involved in the initiation of angiogenic sprouting [467–472]. In

addition, the cytokines TNFa and IL-1 β , generated by the NF- κ B pathway, seem to play an important accessory role in angiogenesis [473,474].

Secreted VEGF can bind to the ECM, thereby creating a gradient of VEGF from the source (e.g., macrophage) to the endothelium [471]. Although many endothelial cells are subjected to the VEGF gradient, only a minor proportion of them develop a tip cell phenotype. The selection process seems to be dependent on Delta-like 4 (Dll4)/Notch signaling in some of the endothelial cells [464,465,471]. In some cells, VEGFR (primarily VEGFR2 or KDR/ Flk1) signaling increases VEGFR expression and induces the expression of Dll4; these cells will develop into tip cells. The enhanced Dll4/Notch signaling to adjacent endothelial cells suppresses VEGFR signaling in the tip cell favors lamellipodia and filopodia development and migration, whereas the increased Dll4/Notch signaling in adjacent endothelial cells inhibits filopodia development and drives the cells into a stalk phenotype [465,471].

Angiogenesis is facilitated by the expression of MMPs, which are capable of local degradation of basement membrane and ECM components [465,475]. MMP activity is limited to the tips of the sprouting neovessels [476]. The degradation of the ECM is highly regulated (both spatially and temporally), because, in addition to providing an impediment to cell migration, various ECM components serve as a scaffold for endothelial cell migration and regulate growth factor bioavailability [475,477,478].

VEGF also influences the dynamics of endothelial cell–cell interactions in the angiogenic sprout. VEGFR is closely associated with VEcadherin and upon VEGFR ligation VE-cadherin is phosphorylated and internalized [272,479,480]. Loss of VE-cadherin from the AJ contributes to destabilization of endothelial cell–cell contacts, thus accounting for the high permeability of angiogenic sprouts [481–484].

Macrophage-derived VEGF may also contribute to the angiogenic process by increasing mobilization of myeloid cells and endothelial progenitor cells (EPCs) to the site of angiogenesis [434,474,485,486]. Most of the myeloid cells (e.g., macrophages or macrophage precursors) seem to be derived from the bone marrow [474], whereas the EPCs are recruited primarily from local sites (e.g., blood vessel wall) [474,486]. Of interest is the observation that a subpopulation of macrophages (M2; involved in wound healing and angiogenesis) can accumulate in the affected area via proliferation rather than recruitment from the vasculature [487]. The localization of VEGF-recruited accessory cells to the affected area is believed to be due to perivascular expression of SDF1 (CXCL12) to attract CXCR4-bearing myeloid and endothelial progenitor cells [434,488]. There is evidence to indicate that these recruited accessory cells are the major driving force for angiogenesis [434,474]. For example, macrophage-induced angiogenesis is dependent on VEGF secretion by EPCs [474]. The issue of whether recruited EPCs actually become incorporated into the angiogenic stalk remains controversial [434,486,488].

Activated macrophages also release the cytokines TNFa and IL-1 β , which contribute to angiogenesis in the inflammatory/hypoxic setting. TNFa is believed to "prime" endothelial tip cells during the inflammatory phase for eventual migration induced by VEGF during the resolution phase [473]. IL-1 β plays an important role in the recruitment of accessory cells, e.g., EPCs and myeloid cells [474].

Role of ROS and NO (Fig. 4B)

VEGF-induced angiogenesis is associated with increased endothelial cell production of both ROS [219,489,490] and NO [491–493]. Genetic blockade approaches indicate that a major source of ROS is NADPH oxidase (NOX) [219], whereas eNOS is the source of NO [494].

Kvietys and Granger

At the receptor level, VEGFR signaling induces activation and translocation of the small GTPase Rac1 into the plasma membrane where it activates NOX [490]. The ROS generated by NOX, in turn, oxidize PTP (shifting the PTK/PTP balance in favor of PTK), which allows for autophosphorylation (activation) of VEGFR. Thus, a positive feedback system is initiated that amplifies VEGFR signaling via NOX. VEGFR signaling also activates eNOS via a PI3-kinase/Akt or AMPK pathway [493], although the role of the latter pathway has been questioned [495]. VEGF-induced increases in Rac1 activity can enhance eNOS activity as well as NADPH oxidase activity; both NO and superoxide production can occur concurrently [410,496]. Interestingly, VEGFR, eNOS, and NOX are compartmentalized within endothelial lipid rafts or caveolae [68,405,480,497]. This compartmentalization may provide an important signaling platform for efficient angiogenesis, because caveolae-deficient endothelial cells cannot migrate [491].

Endothelial tip cell migration toward a VEGF gradient involves penetration of the ECM, a process facilitated by endothelial MMPs [465,475,491]. MT1-MMP is localized within caveolae, and caveolamediated trafficking of the MMP within the endothelial cells ensures appropriate localization and activity [498]. eNOS-derived NO promotes MT1-MMP clustering and activation at the migrating front of endothelial cells, a critical event in endothelial migration and tube formation [499]. At low concentrations, NO can inhibit tissue inhibitor of metalloproteinase (TIMP) via a cGMP pathway, thereby altering the TIMP/ MMP balance in favor of MMP activity [500]. At high concentrations, NO may also interact with the cysteine-zinc moiety of MMPs, thereby enhancing their activity [500]. Endothelial cell migration requires a cycling of adhesive interactions to support directed movement, a process requiring ROS signaling [490]. As mentioned previously, NOX-derived ROS can alter the PTK/PTP balance in favor of PTK (oxidation of critical residues in PTPs), which results in phosphorylation of VE-cadherin/β-catenin and IEJ disassembly [246,398]. In addition, NOX is localized to the lamellipodia of migrating endothelial cells by polarized migration of lipid rafts [501]. Specific scaffolding proteins localize NOX signaling to focal adhesion complexes of lamellipodia and the leading "ruffled" edge of the migrating endothelium [502]. A ROS-induced PTK/PTP imbalance is believed to modulate focal complex turnover [501,502].

VEGF-induced destabilization of microvessels renders the developing microvasculature more permeable to macromolecules (vascular hyperpermeability) [368,481,483]. The VEGF-induced generation of NO and ROS is believed to play an important role in vascular hyperpermeability [368,503]. NOX-derived ROS have been implicated in the VEGFinduced hyperpermeability via phosphorylation of VEcadherin [368], leading to VEcadherin disassociation from the junctions and internalization [484]. The phosphorylation of VE-cadherin has been attributed to the nonreceptor tyrosine kinases of the Src family [483,484]. VEGF-induced activation of eNOS also results in endothelial production of NO and increased endothelial monolayer permeability [503]. The NO-mediated hyperpermeability requires eNOS endocytosis to subcellular compartments [504] and may involve the sGC/cGMP pathway [505]. Alternatively, NO-mediated S-nitrosylation of β catenin (Cys619) and AJ disassembly can contribute to the VEGF-induced hyperpermeability [408].

Functional restabilization of microvessels

As angiogenic sprouting progresses, cell-specific signals ensure appropriate maturation of the nascent blood vessel, i.e., vessel wall maturation/stabilization and lumen formation. Endothelial sprout maturation involves the recruitment of mural cells and strengthening of the barrier function of the neovessel [466,506,507]. Platelet-derived growth factor β (PDGF β) secreted by tip cells is primarily responsible for the recruitment of mural cells, i.e., pericytes and vascular smooth muscle cells [467,468]. PDGF β binds to local ECM

components, thereby creating a gradient for recruitment of mural cells. Mural cells express the PDGF β receptor, whereas endothelial cells do not [468]. PDGF β -dependent recruitment of mural cells is complex and involves stimulation of mural cell proliferation and migration to and adhesive interactions with the sprouting endothelium [506].

The newly recruited mural cells play a pivotal role in stabilization of the neovessel by generating angiopoietin-1 (Ang-1), which interacts with Tie2 receptors on endothelial cells [466]. Ang-1 plays a role in ensuring that MMP activity is confined to the tip cells and inhibits MMP activity in the stalk cells [476]. Ang-1/Tie2 signaling serves to stabilize the endothelial lining by strengthening cell-cell interactions [481,506,508]. Ang-1 ensures Tie2 localization to cell-cell contacts, where it forms complexes with VE-PTP and serves to stabilize IEJs, presumably via prevention of VE-cadherin phosphorylation [508]. Ang-1/ Tie2 signaling also inhibits VEGFR-mediated activation of Src kinase and subsequent VEcadherin phosphorylation and internalization [509]. This may represent a major mechanism by which Ang-1/Tie2 signaling antagonizes VEGF-induced hyperpermeability [507]. As a caveat, Ang-1/Tie2 signaling can also contribute to the destabilization phase of angiogenesis, an effect observed primarily when cell-cell contacts are disrupted and involve endothelial cell- ECM Ang-1/Tie2 complex formation [508,509]. Ang-2, which is generated by endothelial cells in response to VEGF, can serve as an antagonist of Tie2 activation and support the destabilization phase [466,509]. Collectively, the available information indicates that Ang-1 serves to stabilize the VEGF-induced angiogenic sprout, thereby contributing to the development of functional neovessels [510].

The mechanisms involved in lumen formation are poorly understood. It has been proposed that it occurs either by extensive coalescence of endothelial vesicles forming a space between adjacent endothelial cells or by invagination of endothelial cells around a preexisting space [465,511,512]. The Ang-1/Tie2 signaling pathway has been implicated in increasing the caliber of the sprouting vessels via apelin interactions with its receptor APJ [511,513].

Transforming growth factor β (TGF β) and S1P signaling has also been implicated in the final maturation of neovessels. TGFB secreted by both endothelial and mural cells is present in a latent form in the ECM [478]. When activated by MMP degradation of ECM components, TGFβ induces differentiation and maturation of mural cells [506], inhibits endothelial cell proliferation [514], and contributes to endothelial cell synthesis of basement membrane components (e.g., collagen IV, fibronectin) [471]. Coculture systems highlight the cooperative interaction between endothelial cells and mural cells in (1) activating latent TGF β [514,515] and (2) appropriate development of an endothelial basement membrane via regulation of the TIMP/MMP balance [516]. S1P, derived primarily from blood cells (red blood cells, platelets) [354,466,506] and also generated by endothelial cells [481], interacts with its receptor (S1P1) on endothelial cells to further strengthen both endothelial cell adhesive interactions (VE-cadherin) and endothelial cell-mural cell adhesive interactions (N-cadherin) [481]. In addition, S1P derived from recruited progenitor cells has been shown to be critical for endothelial adherens junction assembly [517]. There may also be significant cross talk between S1P and TGFB, i.e., SIP can activate TGFB receptors and activation of TGF^β receptors can generate S1P [518].

Role of ROS and NO

Although NADPH oxidase-derived ROS have been shown to play a role in PDGF β -induced smooth muscle chemotaxis [519,520], a direct link to mural cell coverage of sprouting endothelium is lacking. Moreover, there is relatively little information regarding the role of ROS in other phases of blood vessel stabilization, e.g., strengthening of inter-endothelial cell junctions. Indeed, based on the development of the vasculature in mice with endothelial-

specific deletion of Rac1, it has been proposed that NADPH oxidase is more important in the initiation of angiogenesis than in the development of mature vessels [521].

NO derived from eNOS has been shown to play an important role in vessel maturation and mural cell coverage of developing vessels in several models of angiogenesis. Mural cell recruitment is impaired in eNOS-deficient mice [522] and the adenoviral delivery of eNOS promotes pericyte coverage of neovessels [523]. However, the precise signaling mechanisms have not been addressed.

Ang-1 can reverse the VEGF-induced hyperpermeability of endothelial cell monolayers by phosphorylating the inhibitory Thr497 residue on eNOS, thereby decreasing NO production [503]. Alternatively, it has been proposed that Ang-1/Tie2 signaling can increase the expression of eNOS (via KLF2) and enhance IEJ stability [481]. The apelin/APJ pathway can also increase eNOS via the KLF2 pathway [524], but a direct linkage between NO generation and neovessel development has not been established.

Pharmacologic blockade of NO has been shown to enhance TGF β production in wounds, implying a role for NO in limiting TGF β production and excessive deposition of ECM (fibrosis) [525]. However, neither the source/isoform of NOS nor the mechanisms involved in eliciting this response are clear at present. S1P can also lead to eNOS activation [526], but whether NO is involved in the S1Pinduced strengthening of endothelial barrier function has not been directly assessed.

In summary, there is a paucity of information on the roles of ROS and NO in the stabilization phase of angiogenesis and further work in this understudied area is warranted.

Coagulation and thrombosis

It is now well recognized that inflammation and coagulation are intimately linked processes that can elicit a vicious cycle because activation of one process leads to activation of the other [527–529]. This interdependence of inflammation and coagulation is evident in a variety of acute (e.g., sepsis) and chronic (e.g., colitis) inflammatory diseases wherein changes in hemostatic biomarkers suggest subclinical activation of the coagulation system and there is an increased incidence of thromboembolic events [10,530–537]. Whereas the chemical mediators that link inflammation to coagulation and thrombosis remain poorly understood, the activation products of the various cell populations (platelets, leukocytes, endothelial cells, macrophages) that drive the inflammatory response have received the most attention. Reactive oxygen and nitrogen species are considered likely candidates because of their ability to alter platelet function and to influence the balance between procoagulant, anticoagulant, and fibrinolytic systems [538]. This section summarizes evidence that implicates ROS and reactive nitric oxide species (RNOS) as potential mediators of the procoagulant, prothrombotic phenotype that is assumed during inflammation.

The coagulation cascade (Fig. 5)

The antithrombogenic properties of the vascular system can be attributed to normal endothelial cell function [539–541]. These cells produce (e.g., prostacyclin and nitric oxide) or limit the accumulation of (e.g., the ectonucleotidase CD39 degrades ATP and ADP) substances that inhibit the adhesion and aggregation of platelets [542,543]. Endothelial cells also facilitate the production of anticoagulants (activated protein C, or APC) that prevent the activation/ deposition of coagulation factors on the vessel wall. Another important function of endothelial cells is the formation of a barrier that physically separates blood elements from the subendothelial matrix [1]. When the endothelial lining is disrupted and the underlying matrix is accessible to blood, platelets will rapidly adhere to exposed collagen

and von Willebrand factor in the subendothelial space (via an interaction between $\alpha 2\beta 1$ and glycoprotein (GP) Ib/IX). This is followed by platelet aggregation (via fibrinogen and GPIIb/IIIa) and the expression of cell-surface phospholipids, on which clotting factors in blood can assemble. During inflammation, the blood vessel wall does not exhibit endothelial damage or exposure of collagen and matrix material. In this setting, endothelial cell activation can result in the binding of platelets, with or without attached leukocytes, which can also result in platelet aggregation, activation of the coagulation system, and thrombus development [1]. Tissue factor (TF), which is expressed by endothelial cells, monocytes, platelets, and other cells in response to inflammatory mediators (e.g., TNFa, IL-1β, CD40L), initiates coagulation when exposed to blood [544]. TF triggers coagulation by binding to activated factor VII, which ultimately leads to activation of other coagulation factors and the conversion of prothrombin to thrombin. Thrombin cleaves fibrinogen to generate fibrin monomers, which polymerize to form a stable clot. Thrombin also amplifies the coagulation process by inducing TF expression on the vessel wall, via feedback activation of cofactors VIII and V, and by activating platelets through engagement of thrombin receptors [545]. Fibrinolysis (proteolytic degradation of fibrin), which prevents excess thrombus growth, is mediated by plasminogen and its activators (t-PA and u-PA) and controlled by plasminogen activator inhibitor-1 (PAI-1), which inhibits t-PA and u-PA [546].

Endogenous anticoagulant mechanisms, including tissue factor pathway inhibitor (TFPI), heparin–antithrombin, and the protein C pathway, serve to either inhibit the generation of thrombin or inactivate it [527,547]. TFPI inactivates factor VIIa bound to TF, whereas the heparin–antithrombin mechanism neutralizes factor Xa, thrombin, and factor IXa, as well as factor VIIa bound to TF [527,547]. The protein C pathway is initiated by the binding of thrombin to thrombomodulin on the surface of endothelial cells. The thrombin–thrombomodulin complex activates protein C, but this activation proceeds more efficiently when protein C is bound to the endothelial protein C receptor (EPCR). Although APC retains its affinity to EPCR, upon dissociation from EPCR it binds to protein S, which enables it to inactivate factors Va and VIIIa. Collectively, the anticoagulant mechanisms serve to effectively shut off coagulation; however, chemical signals that downregulate these mechanisms can initiate coagulation [527,528,539,541].

ROS and coagulation

The tight control of the coagulation system observed in healthy tissue can be disrupted by ROS, which can alter the expression and/or activity of key components of the coagulation cascade [534]. Redox-sensitive regulatory mechanisms have been implicated in the production and activation of tissue factor. The gene encoding TF is redox sensitive [548]. Exposure of endothelial cells or monocytes to ROS (e.g., H2O2 or xanthine-xanthine oxidase) results in increased TF mRNA and protein expression as well as enhanced TF procoagulant activity [549,550]. The action of ROS on endothelial cell TF expression is probably amplified by a corresponding inhibitory effect of ROS on tissue factor pathway inhibitor production by endothelial cells [551]. Endogenous ROS exert a similar influence on TF expression because the increased TF expression elicited in endothelial cells in response to cytokines (e.g., TNFa) and on vascular smooth muscle or platelets activated by thrombin is blunted by antioxidant treatment [552-555]. The redox-sensitive expression of TF has also been demonstrated in the heart after exposure to I/R. The I/R-induced TF expression was abolished by treatment with oxygen radical scavengers [549]. Several studies have linked ROS-dependent induction/activation of TF to NADPH oxidase in endothelial cells, platelets, and neutrophils [553,556]. A role for NADPH oxidase-derived ROS in thrombin-induced TF expression is supported by studies employing p47phox^{-/-}
Other components of the coagulation pathway are also influenced by ROS to promote a procoagulant environment. The protein C pathway is altered in several ways by ROS. The shedding of EPCR by endothelial cells can be induced by ROS and antioxidants have been shown to blunt the EPCR shedding elicited by cytokines [557]. APC, a serine protease, can be directly inactivated by oxidants, which induce changes in the structure of the active site of APC, in part via the oxidation of methionine [558]. Thrombomodulin, which also participates in the activation of protein C, is inactivated by neutrophil-derived oxidants via a mechanism that involves the oxidation of specific methionine residues [559]. Whereas ROS can act in multiple ways to curtail the generation of APC, it is interesting to note that this potent endogenous anticoagulant has been shown to possess intrinsic antioxidant properties [560], which has led to the proposal that the APC may exert its anti-inflammatory actions via this mechanism.

Thrombin and the heparin–antithrombin complex are also vulnerable to ROS-mediated functional alterations. Neutrophil-derived HOCl, but not H_2O_2 , has been shown to oxidize and functionally impair human thrombin [561]. The oxidized thrombin exhibits a diminished capacity to interact with thrombomodulin, protein C, and the antithrombin III–heparin complex, as well as a reduction in its ability to activate platelets. Oxidation of antithrombin by hydrogen peroxide appears to minimally alter thrombin-binding activity and heparin binding of the anticoagulant [562]. However, lipid peroxides have been shown to significantly reduce antithrombin activity by targeting the heparin-binding site [563].

The procoagulant, prothrombotic effects of ROS are also the result of effects on key components of the fibrinolytic system. Oxidatively modified fibrinogen forms fibrin at an accelerated rate and has a reduced capacity to stimulate tissue plasminogen activator to convert plasminogen to plasmin, compared to nonoxidized fibrinogen [564]. ROS also enhance fibrinolysis by inhibiting plasminogen activator [565] and by increasing the transcription of PAI-1 in endothelial cells, vascular smooth muscle, and other cells [566,567]. The contention that PAI-1 expression is redox sensitive is supported by reports describing the inhibition of PAI-1 expression in vascular smooth muscle cells by treatment with antioxidants or depletion of the p22 subunit of NADPH oxidase [568]. The oxidant-sensitive transcription factor AP-1 has been implicated in ROS-induced upregulation of PAI-1 [569].

NO and coagulation

Whereas superoxide and hydrogen peroxide tend to activate the coagulation cascade, nitric oxide exerts an opposite effect. Endogenous NO appears to prevent the development of a prothrombotic phenotype in endothelial cells. L-Arginine- but not D-arginine-supplemented endothelial cells exhibit a blunted expression of TF in response to challenge with either LPS or interleukin-1 β [570]. A similar effect on endothelial cell TF expression (induced by LPS) is noted on human umbilical vein endothelial cell monolayers exposed to the NO donor SNAP, whereas the NO synthase inhibitor L-NAME enhances the expression of TF [571]. The role of NO as a regulator of TF expression on endothelial cells has also been demonstrated in murine models of eNOS deficiency or overexpression [572].

Although endothelial cells have received considerable attention regarding the modulatory influence of NO on TF expression, there is also evidence implicating NO as a regulator of TF expression on monocytes. For example, the endogenous NO synthase inhibitor ADMA has been shown to elicit a dose-dependent increase in TF expression on the monocytic cell line THP-1 [573]. The effect of ADMA is NF- κ B dependent, suggesting that NO mediates

TF expression at the level of transcription. This differs mechanistically from the direct actions of peroxynitrite (ONOO), a product of the reaction of NO with superoxide, on TF activity. The extracellular domain of TF is rich in tyrosine, an amino acid that is vulnerable to attack by ONOO. Consequently, recombinant TF exposed to ONOO has more nitrotyrosine residues and exhibits reduced procoagulant activity [574]. The effects of ONOO on coagulation do not seem to be limited to TF. Exposure of human plasma to ONOO in vitro impairs hemostatic function and this is accompanied by decreased activities of factors VII and X and the factor VIII complex, as well as antithrombin and protein C activity [575].

The fibrinolytic system is also a target for NO and its products. In vitro studies indicate that NO donors decrease the expression and release of PAI-1 [576], whereas NOS inhibitors induce vascular PAI-1 expression [577,578]. The actions of NO on PAI-1 seem to be mediated via a cGMP-dependent mechanism. The net effect of NO on fibrinolytic balance remains unclear because the gaseous oxide also appears to reduce the expression and release of t-PA [579]. The influence of NO on fibrinolysis is evident in NO-deficient mice, which are characterized by increased t-PA levels, enhanced fibrinolysis, and decelerated thrombosis. The enhanced fibrinolysis has been attributed to a lack of NO-mediated Weibel-Palade release of t-PA (which is stored in these endothelial cell granules) [580].

Nitrosative stress has been implicated in the modulation of fibrinolysis. Some reports describe time- and dose-dependent plasminogen inactivation after exposure to the peroxynitrite donor 3-morpholinosydnonimine [581]. ONOO has also been reported to decrease fibrinogen function and inactivate t-PA [582]. It has been suggested that nitrating oxidants significantly accelerate clot formation and factor VIII cross-linking, whereas nonnitrating oxidants decelerate clot formation [583]. The results of this study also suggest the clots formed with fibrinogen exposed to nitrating oxidants may be more readily deformed by mechanical stress, which may have some bearing on the process of atherothrombosis.

Platelets: a source of ROS/RNOS

The activation, adhesion, and aggregation of platelets, a critical component of thrombus formation, represent the primary response to vascular injury (with activation of the coagulation cascade and fibrin formation as secondary responses). Upon exposure to activating signals (e.g., collagen), platelets rapidly adhere to the vessel wall, release the contents of their granules, undergo shape changes that allow them to spread and flatten along the vessel surface, and interact (aggregate) with one another to form a plug that seals the injured vessel surface [540,541,544]. Another consequence of platelet activation and aggregation is an enhanced ROS production, which is accompanied by increased oxygen consumption and elevated glutathione disulfide levels [584,585]. Stimulated platelets have been shown to produce superoxide, hydrogen peroxide, and hydroxyl radicals [586–589]. The quantity (flux) of superoxide produced by activated platelets is in the nanomolar range and comparable to that generated by endothelial cells, but is a small fraction of the amount (micromolar range) generated by activated neutrophils [590,591]. Consequently, it has been suggested that platelet-derived ROS probably play a more important role in autocrine or paracrine signaling, rather than a phagocytic function [586,592].

Although platelets have the potential to produce ROS from several enzymatic sources, most reports have attributed platelet-derived ROS to NADPH oxidase [586,592,593]. The isoform of NADPH oxidase expressed by platelets is similar to that found in neutrophils, with subunits for gp91phox, p47phox, and p22phox [590,594,595]. The limited capacity of platelets from patients with gp91phox deficiency to produce superoxide further highlights the importance of this enzymatic source [596]. Platelet agonists (e.g., collagen) increase

NADPH oxidase-dependent ROS production via a mechanism that involves PI3-kinase- and PKCζ-mediated translocation of the p47phox subunit to the plasma membrane [597]. Although collagen-induced GPIIb/IIIa-mediated platelet aggregation is associated with enhanced superoxide production via NADPH oxidase [538], the time course of superoxide generation lags behind the aggregation response, suggesting that superoxide is not critical for the initiation of aggregation [592]. Nonetheless, both NADPH oxidase inhibitors and superoxide scavengers have been reported to blunt platelet aggregation and thrombus formation on collagen [598].

Platelets and megakaryocytes also express the same isoform (NOSIII) of nitric oxide synthase that is detected in endothelial cells [599]. Platelet-associated NOS normally generates nitric oxide, but when cofactor (e.g., BH₄) levels are limited, the enzyme can also generate superoxide [8]. "Uncoupled" NOS has been implicated in platelet superoxide production because platelets derived from eNOS-deficient mice exhibit a marked attenuation of superoxide flux [600]. More attention has been devoted, however, to the nitric oxidegenerating function of platelet-associated NOS. Both resting and aggregating platelets generate NO, but the levels of NO produced by platelets is much lower than reported for endothelial cells [601,602]. As described for endothelial cells, the activity of plateletassociated NOS is regulated by Ca²⁺/calmodulin and requires NADPH [601,603]. The dependence of platelet NO production on NOS is evidenced by the absence of NO generation in wild-type platelets treated with NOS inhibitors and in eNOS-deficient platelets [604]. Platelet agonists, such as collagen and ADP, activate platelet NOS, which leads to increased intracellular levels of cGMP [599]. The effect of collagen on platelet NOS is mediated through engagement of the GPVI receptor and involves both the PI3-kinase/Akt pathway and protein kinase C [600]. CD40L, which is largely derived from platelets, has been shown to mediate agonist-dependent production of NO as well as superoxide in platelets through activation of the Akt and MAP kinase signaling pathways [605].

Platelets: a target for ROS/RNOS

Both superoxide and nitric oxide have been implicated in the regulation of platelet function, with the former inducing cell activation and aggregation and the latter exerting an inhibitory effect. A variety of mechanisms have been implicated in the proaggregation actions of ROS, including the inactivation of NO [534,606], inhibition of redox-sensitive ecto-ADPases [534,590,606], and enhanced reactivity of platelets to agonists, such as thrombin, ADP, and collagen [607]. Activation of platelet NADPH oxidase results in an increased release of CD40L from platelets (a indicator of platelet activation) [608] and elicits a concomitant activation of glycoprotein IIb/IIIa and α IIb β 3 integrin, which mediate platelet aggregation and adhesion [594], suggesting that the low levels of ROS produced by platelets are sufficient to modulate platelet function. Whereas both intra- and extracellular O₂⁻ appear to modulate CD40L expression and release, studies performed with extracellular antioxidants suggest that P-selectin expression is regulated by extracellular O₂⁻, and α IIb β 3 integrin activation is regulated by intracellular superoxide [609].

Although there is evidence that exogenous ROS can alter platelet function, the responses of platelets to various ROS species have been inconsistent and seem to result from differences in experimental protocols, including whether ROS exposure occurs in the presence of plasma. For example, superoxide is known to enhance platelet aggregation [610], but not in the presence of plasma [611–613]. The effect of superoxide is generally attributed to inactivation of NO; however, it has also been proposed that superoxide may indirectly promote platelet aggregation by inactivating PAF-acetylhydrolase, the enzyme that degrades PAF, a potent platelet agonist [614]. H_2O_2 can either stimulate or inhibit platelet aggregation [612], even in the presence of plasma [615–617]. The inhibitory effect of H_2O_2

seems to result from stimulation of guanylate cyclase [611], whereas the stimulatory effect has been attributed to Ca^{2+} release from the sarcoplasmic reticulum as a result of oxidation of sulfhydryl groups in the sarcoplasmic/endoplasmic reticulum Ca^{2+} ATPase and enhanced Ca^{2+} release from mitochondria [618]. Oxidized low-density lipoprotein (oxLDL) can also stimulate or inhibit platelet function in a concentration-dependent manner [619]. Whether the levels of H₂O₂ and oxLDL shown to alter platelet function are pathophysiologically relevant remains unclear.

It is now widely accepted that NO is an important regulator of platelet function, with NO derived from either endothelial cells or platelets exerting an inhibitory influence on platelet adhesion and aggregation [593,599]. Nitric oxide is also able to disaggregate preformed platelet aggregates [620] and to inhibit the recruitment of platelets into aggregates [621]. NO is considered to have an autoregulatory function in modulating platelet reactivity, because the NO produced by platelets upon activation serves to limit further adhesion and aggregation [621]. The inhibitory effects of NO on platelet function are counteracted by superoxide, which inactivates NO, and stabilized by superoxide dismutase. NOS inhibitors promote platelet adhesion and aggregation in vivo, suggesting that the basal level of NO produced by endothelial cells and platelets is sufficient to prevent these responses. NO donors or L-arginine inhibit platelet adhesion and aggregation in vivo [622,623]. The effects of NO on platelet function are largely related to stimulation of sGC and the generation of cyclic GMP. Several signaling pathways have been implicated in NO-mediated, cGMPdependent modulation of platelet function, including cyclic nucleotide-gated cation channels, cGMP-dependent protein receptors, cGMP-regulated phosphodiesterases, and cGMP-dependent protein kinases [624]. The net effect of cGMP-mediated activation of these pathways is a reduction in intracellular calcium, which inhibits the conformation change of GPIIb/IIIa into its active form [625]. Some actions of NO on platelets have also been attributed to a cGMP-independent mechanism [625]. For example, S-nitrosylation of N-ethylmaleimide-sensitive factor has been implicated in NO-mediated inhibition of platelet granule exocytosis and this effect is not prevented by sGC inhibition [626].

Thrombus formation: role of ROS and RNOS

Based on the known actions of ROS and NO on the coagulation cascade and platelet function, it can be predicted that ROS promote thrombus development and NO is antithrombotic. Studies that directly examine thrombus formation after endothelial injury in macroscopic or microscopic blood vessels are generally consistent with this prediction. Superoxide dismutase, catalase, and less specific free radical scavengers have been shown to blunt thrombus development in a variety of in vivo models [627,628]. The protection against thrombosis afforded by SOD has been attributed to preservation of NO bioavailability [629]. Iron chelation with desferrioxamine has also proven effective in blunting thrombosis [630], whereas chronic iron administration appears to accelerate arterial thrombosis [631], suggesting that iron-catalyzed radicals derived from superoxide and hydrogen peroxide contribute to thrombogenesis. There is evidence implicating both NADPH oxidase [598] and xanthine oxidase [632] as sources of the ROS involved in thrombus development.

The contribution of endogenous NO to thrombus formation is less clear. Pharmacological [633] inhibition of NOS appears to enhance thrombus development, an effect that can be reversed by coadministration of L-arginine or an NO donor [627,634,635]. This protective effect of endogenous NO is more evident in venules than in arterioles in otherwise healthy animals [635]. The prothrombotic effects of hypercholesterolemia and hyperhomocysteinemia have also been linked to reduced NO bioavailability [531,636]. Hypercholesterolemia elicits a more profound acceleration of thrombosis in arterioles than in venules, and the arteriolar response is reversed by L-arginine treatment [531]. eNOS-deficient mice, on the other hand, do not exhibit spontaneous thrombosis and some studies

reveal a minor role for NO in the inhibition of arterial thrombosis [604,633]. Other studies, however, indicate that eNOS deficiency potentiates thrombus development and the hemodynamic abnormalities associated with thrombosis [637,638]. The inconsistent results generated in eNOS-deficient mice have been attributed to compensatory mechanisms that oppose the effects of eNOS gene deletion [580] as well as the observation that the platelet phenotype of eNOS-deficient mice is apparent only after moderate to high thrombotic stimulation [637]. The relative contributions of platelets, endothelial cells, and other cells (e.g., leukocytes, macrophages) to the ROS and NO that modulate thrombus formation remain unknown.

Epilogue

The literature is generally consistent with the view that alterations in ROS and NO contribute to the five characteristic microvascular responses to inflammation, i.e., vasomotor dysfunction, leukocyte recruitment, increased vascular permeability, angiogenesis, and thrombosis. Although the relative contributions of ROS and NO to the inflammationinduced changes in vascular function and structure remain unclear, it is evident that the highly reactive nature of NO and ROS with specific cellular and molecular targets within and surrounding the blood vessel wall enables these mediators to elicit the diverse microvascular responses to inflammation. Of equal importance is the role of endothelial cells, not only as a source and target of ROS and NO, but also as critical players in each of the five vascular responses to inflammation. Endothelial cells seem to serve as gatekeepers in the modulation of vascular smooth muscle tone and for the adhesion and transmigration of leukocytes and the proliferation of blood vessels, and they provide the interface for the activation and deposition of coagulation factors and the binding of platelets during thrombogenesis. These unique properties of endothelial cells justify the inordinate attention that has been devoted to studying their production of, and responses to, reactive oxygen and nitrogen species.

Although much has been learned about the roles of NO and ROS in regulating the growth and function of blood vessels in health and disease, there are several areas of uncertainty and controversy in this field that warrant more attention. These include the interplay between the ROS- and the NO-generating systems, compartmentalization of ROS and NO generation, and the contribution of the endothelial glycocalyx as a source and target of ROS and NO.

A classic example of the interplay between ROS- and NO-generating systems is the uncoupling of eNOS by ROS, which converts eNOS from an NO-generating enzyme to one that generates superoxide [65]. The increased generation of ROS (e.g., NADPH oxidase) can result in BH₄ oxidation [64] and the depletion of NADPH [216], which results in the uncoupling of eNOS. However, it is also becoming apparent that, under certain conditions, endogenous ROS (e.g., NADPH oxidase-derived H₂O₂) can prime/activate eNOS to generate increased levels of NO [639–641]. This phenomenon highlights the unpredictable nature of the interplay between ROS and NO and should serve as a warning about data interpretation when there is a potential for both systems to be activated.

The multiple sources of NO (NOS isoforms) and ROS (mitochondria, oxidases), as well as the proximity of these sources to their molecular targets, necessitate a mechanism to ensure specificity of signaling. It has been proposed that signaling specificity of ROS and NO is achieved by subcellular compartmentalization of the generating systems and the relevant effector molecules [97,369,405,641–643]. In endothelial cells, NOS has been localized to the Golgi apparatus, cytosol, cytoskeleton, or lipid rafts/caveolae of the plasma membrane. Similarly, NADPH oxidase has been localized to the perinuclear endoplasmic reticulum, cytoskeletal elements, and lipid rafts/caveolae of the plasma membrane. This differential

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localization of ROS- and NO-generating systems should allow for selective activation of the NO- and ROS-producing enzymes and localized production of these reactive species. Furthermore, the localization of the ROS- and NO-generating systems is not fixed; both eNOS and NADPH oxidase can be translocated to other compartments to induce functionally distinct microvascular responses. For example, acetylcholine-induced vasodilation is associated with translocation of eNOS from the plasma membrane to the Golgi, whereas PAF-induced increases in vascular permeability are associated with translocation of eNOS to a cytosolic compartment [369,644]. During angiogenesis NADPH oxidase is translocated to the lamellipodia of the migrating endothelial cells [97,502]. The mobile nature of the intracellular compartments housing NO and ROS generators is an exciting aspect of ROS and NO signaling that deserves more attention.

An interesting and potentially important revelation of our examination of this literature is the possible contribution of the endothelial glycocalyx to the diverse microvascular responses to inflammation. The endothelial glycocalyx is a dynamic lining of the luminal surface of endothelial cells whose composition varies based on the rate of shedding and biosynthesis of its components [75,76]. This network of carbohydrate-rich proteoglycans, glycoproteins, and associated GAGs exhibits a high binding affinity for a variety of endothelial-cell-derived and circulating molecules that act as generators (xanthine oxidase, myeloperoxidase) or detoxifiers (ecSOD) of ROS [75]. The net effect of this binding property of the glycocalyx is to concentrate molecules on the endothelial cell surface that can alter cell function (e.g., chemokines) and/or the quality and density of the glycocalyx itself (e.g., ROS generators). It is not unexpected that the glycocalyx exerts an influence on the microvascular response to inflammation because the partial dissolution of the glycocalyx that accompanies this condition can: (1) affect the attachment of transmembrane components of the glycocalyx to caveolae, thereby disrupting its function as a mechanosensor for shear-induced vasodilation [75,76]; (2) promote leukocyte-endothelial cell adhesive interactions; (3) diminish the charge selectivity of the endothelial barrier, thereby promoting protein extravasation; and (4) reduce the capacity of the glycocalyx to bind anticoagulant molecules (e.g., anti-thrombin, TFPI) and diminish the thrombo-resistance of the endothelial cell surface [75]. The functional relevance of the glycocalyx to inflammation-induced angiogenesis remains unclear and warrants exploration.

Future development of technology that provides a minimally invasive means for monitoring vascular dysfunction and for the detection, quantification, and selective ablation of ROS and NO in the living animal is critical for advancement of this field of inquiry. In this regard, substantial progress is evident, in terms of both the approaches being applied currently and those with potential for future applications. For example, a fluorescent H_2O_2 sensor (HyPer) is available to monitor H_2O_2 levels in vitro [640,645] and in vivo [194]. It is specific for H₂O₂ (does not respond to superoxide or NO) and allows for compartmental quantification of H_2O_2 in real time, e.g., up to hours after injury and for a distance of 100–200 mm from the site of injury [194]. The burgeoning area of posttranslational regulation of gene products by microRNAs (miR's) also holds promise in the field of ROS and NO research. Several miR's have been identified that can regulate the inflammatory response [646–648], including miR's that can exert an influence, either directly or indirectly, on endothelial NADPH oxidase [649] and NOS [650,651]. Future advances in this area should lead to the availability of specific miR's and antagomirs to more comprehensively assess the roles of ROS and NO in the microcirculatory responses to inflammation. Epigenetics [652] and gastrointestinal biota [653] are two of many additional factors that warrant consideration in future studies that address the roles of ROS and NO in mediating the microvascular dysfunction that is associated with inflammation.

The quantitative significance of the balance between NO and ROS to the microvascular dysfunction in inflammation remains uncertain. Hence, there is a clear need for more research on, and an improved understanding of, the involvement of reactive oxygen and nitrogen species in the vascular responses that accompany different acute and chronic inflammatory diseases. Future development of technology that provides a noninvasive means for monitoring vascular dysfunction and the detection, quantification, and selective ablation of ROS and NO in the living animal is critical for advancement of this field of inquiry. The information that would result from this effort should provide novel and potentially useful insights into the signals that drive inflammation and other microvascular diseases (e.g., diabetes) and ultimately lead to the development of novel therapeutic strategies that diminish the morbidity and mortality associated with these diseases.

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Fig. 1.

Representative endothelial-dependent relaxing factors (EDRFs) and endothelial-dependent hypopolarizing factors (EDHFs). Endothelial activation by shear stress or ligands of Gprotein-coupled receptors (GPCRs) increases intracellular levels of Ca^{2+} , which is the initial event in the generation of EDRFs and EDHFs. EDRFs (red pathways): elevated endothelial Ca^{2+} levels activate at least two enzymes that generate EDRFs, nitric oxide synthase (eNOS) and cyclo-oxygenase (COX). NO (or HNO) derived from eNOS diffuses to smooth muscle and activates cGMP. The prostanoid PGI₂ derived from COX interacts with its receptor (IPR) and activates cAMP. These second-messenger systems, in turn, activate Ca^{2+} dependent K⁺ channels (BK_{Ca++}) in smooth muscle resulting in inhibition of voltage-gated Ca^{2+} channels ($Ca^{++}V$). The resultant decrease in intracellular Ca^{2+} levels also result in smooth muscle hyperpolarization by either activating Ca^{2+} -dependent K⁺ channels or generating H₂O₂. Increased K⁺ efflux (via SK_{Ca++} or IK_{Ca++}) results in endothelial membrane hyperpolarization (MHP), which can be transmitted to smooth muscle via myoendothelial gap junctions. Alternatively, the K⁺ ions entering the internal elastic lamina can cause

smooth muscle hyperpolarization. In either case, $Ca^{++}V$ are inhibited and smooth muscle Ca^{2+} levels decrease, resulting in smooth muscle relaxation. Finally, various oxidases (e.g., NADPH oxidase, COX) as well as eNOS (e.g., uncoupled) can generate superoxide, which is rapidly converted to H_2O_2 by Cu,Zn-SOD. H_2O_2 can then diffuse to the smooth muscle where it activates BK_{Ca++} and inhibits $Ca^{++}V$.



Fig. 2.

Leukocyte recruitment to sites of injury/infection: dominant role of ROS. (A) Activation of macrophages by infection/injury. PAMPs (e.g., LPS) generated by infection and DAMPs (e.g., HMGB1) generated by stressed/necrotic cells serve as ligands for TLR and RAGE on interstitial sentinel cells (e.g., macrophages). Ligation of these receptors results in activation of NADPH oxidase (NOX), which generates ROS (superoxide and its dismutation product H_2O_2) (red pathway). ROS can be (1) exported into the interstitium to affect adjacent endothelial cells and (2) further activate NOX (e.g., via an NF- κ B pathway) leading to the generation of additional ROS (feed-forward mechanism). Activation of NF- κ B can also increase iNOS levels, which results in NO generation (blue pathway). NO can dampen the

inflammatory response by interacting with superoxide within macrophages or adjacent cells. (B) Rapid and delayed phases of endothelial activation. Endothelial cells are activated by the proinflammatory milieu (chemokines, cytokines, ROS, LPS, HMGB1). Rapid activation (NF- κ B independent) of endothelial cells by chemokines and ROS results in further ROS generation via NOX (red pathways). Endothelial ROS contribute to adhesion molecule expression (P-selectin), which facilitates leukocyte rolling. ROS have also been implicated in endothelial cell generation of leukocyte activators (e.g., PAF and CXCL8), which are sequestered within the glycocalyx and facilitate leukocyte adhesion to the endothelian. Leukocyte adhesion to endothelium results in the clustering of endothelial adhesion molecules (docking structures). The resultant cell signaling disrupts adherens junctions (AJ) via NOX-derived ROS and facilitates leukocyte TEM. Delayed activation (NF- κ B dependent) of endothelium reinforces the leukocyte–endothelial adhesive interactions via continued and amplified generation of ROS, chemokines, and cytokines via the NF- κ B pathway.

А

GPCR

cAMP





Fig. 3.

Role of Rac1/RhoA balance in endothelial barrier integrity: impact of ROS and NO. (A) Basal permeability. In quiescent endothelial cells, cytosolic Rac1 is dominant and inhibits RhoA activation. Rac1/Cdc42 stabilizes both the cortical actin and the adherens junction components (VE-cadherin/ β -catenin). Basal eNOS activity generates NO (blue pathway), which dampens any superoxide production within endothelial cells. In addition, low levels of NO increase cAMP, whereas high levels decrease cAMP. Agents that tend to strengthen barrier integrity can either increase cAMP, leading to an increase in the Rac1/RhoA ratio (e.g., ANP), or increase eNOS activity (e.g., S1P). (B) Increased permeability. In response to proinflammatory mediators (e.g., VEGF, thrombin, histamine), RhoA becomes dominant

and inhibits cytosolic Rac1. Activation of GPCRs increases intracellular Ca^{2+} levels (via TRP channels) and activates RhoA. RhoA (via Rho kinase) (1) increases the PTK/PTP ratio, leading to disorganization of adherens junctions, and (2) activates MLCK and inhibits MLCP, leading to actomyosin-mediated contraction. Paradoxically Rho kinase activates Rac1 at the membrane, leading to activation of NADPH oxidase and superoxide production (red pathway), which in turn increases the PTK/PTP ratio. The increased intracellular Ca^{2+} also activates eNOS. Although low levels of NO (not shown) tend to stabilize adherens junctions, higher levels of NO (shown) favor disruption of the junctions as well as increasing MLCK activity (blue pathways).





Fig. 4.

Cooperative role of ROS and NO in angiogenesis. (A) Generation of VEGF by macrophages. Both ROS (red pathways) and NO (blue pathways) generated during hypoxia/ inflammation can decrease PHD activity, which activates the HIF pathway as well as the NF- κ B pathway. The HIF pathway leads to transcription of VEGF. The NF- κ B pathway leads to transcription of inflammatory cytokines. During the inflammatory response the NF- κ B predominates, and during the resolution of the inflammatory response the HIF pathway predominates. NO modulates both pathways at several points, being stimulatory in the HIF pathway and inhibitory in the NF- κ B pathway. (B) Initiation of angiogenesis: endothelial destabilization. VEGF derived from macrophages or recruited endothelial progenitor cells

(EPC) interacts with VEGF receptor (VEGFR) on selected endothelial "tip" cells to initiate cell migration. VEGFR ligation activates both eNOS and NOX to generate NO and superoxide, respectively. NO (blue pathways) is involved in (1) loosening of the adherens junctions and (2) MMP activation and TIMP inhibition to facilitate degradation of the ECM. Superoxide (red pathways) is involved in (1) formation of lamellipodia and (2) cyclic alterations in the strength of focal adhesion complexes (FACs). The migratory tip cells are followed by hyperpermeable (destabilized junctions) stalk cells.



Fig. 5.

Role of ROS and NO in coagulation and platelet aggregation. Thrombus formation involves the adhesion, activation, and aggregation of platelets as well as activation of the coagulation cascade. With vessel injury, platelets bind to exposed collagen and von Willebrand factor (vWF). Upon activation, platelets bind to one another using fibrinogen and GPIIb/IIIa to form an aggregate. Tissue factor (TF) triggers coagulation by binding to activated factor VII (fVII), which ultimately leads to activation of other coagulation factors and the conversion of prothrombin to thrombin. Thrombin cleaves fibrinogen to generate fibrin monomers, which polymerize to form a stable clot. Fibrinolysis (proteolytic degradation of fibrin), which prevents excess thrombus growth, is mediated by plasminogen and its activators (t-PA and u-PA) and controlled by plasminogen activator inhibitor-1 (PAI-1), which inhibits t-PA and u-PA. ROS and NO are known to modulate the coagulation pathway and fibrinolysis by interacting with multiple components of this cascade. ROS (red pathways) can promote the initiation of coagulation by targeting the TF-fVII complex as well as tissue factor protein inhibitor (TFPI). ROS also promote coagulation and thrombus formation by inhibiting the production of activated protein C (APC), enhancing the conversion of fibrinogen to thrombin and enhancing PAI-1 activity. NO (blue pathways) tends to exert an opposite effect on the coagulation cascade. In addition, NO targets the platelets to inhibit aggregation, whereas ROS promote this process. NADPH oxidase (Nox) seems to be a major endothelial cell source of the ROS that modulate platelet aggregation.