Short Communication

Down-Regulation of Angiopoietin-1 Expression in Menorrhagia

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Angiogenesis is an essential component of endometrial repair and regeneration following menses. Perturbation of this process is associated with menorrhagia, a common gynecological disorder that results in excessive menstrual bleeding. Angiopoietin-1 (Ang-1) promotes vascular maturation via the Tie-2 receptor, while angiopoietin-2 (Ang-2) is its natural antagonist that destabilizes vessels and initiates neovascularization in the presence of vascular endothelial growth factor. To test the hypothesis that menorrhagia arises as a result of poor signal for vascular maturation, we have examined the expression of Ang-1, Ang-2, and Tie-2 in endometrium throughout the menstrual cycle from 30 normal women and 28 patients with menorrhagia. Ribonuclease protection assay and Western blot analysis showed Ang-2 expression was consistently higher than Ang-1 in normal endometrium throughout the cycle. However, with menorrhagia Ang-1 mRNA and protein were not detected or downregulated, while Ang-2 was observed at similar levels in both normal and menorrhagic endometrium resulting in a greater than a 50% decrease in the ratio of Ang-1 to Ang-2 protein. In situ hybridization and immunohistochemical studies supported these findings and revealed cyclical changes in the expression of Ang-1 and Ang-2. These results suggest that the angiopoietin/Tie-2 system promotes vascular remodeling in endometrium and loss of normal Ang-1 expression may contribute to the excessive blood loss observed in menorrhagia. (Am J Pathol, :773-780)

During menstruation, shedding of the functional layer of the endometrium occurs followed by intense vasoconstriction of the remaining basal arteriolar fragments, which prevents excessive blood loss until the damaged surrounding tissues and blood vessels are repaired and regenerated.¹ Excessive menstrual bleeding (>80 ml) is a common gynecological problem in women of reproductive age, accounting for over 20% of outpatient clinic visits, which may lead to iron deficiency anemia and require hysterectomy. Although commonly associated with the presence of conditions such as fibroids and carcinoma, the finding that approximately 50% of menorrhagia cases occur in the absence of any uterine pathology² suggests a defect in the cellular processes of menstruation. The mechanisms controlling menstruation are regulated by local factors within the endometrium that include vasoregulators,^{3,4} angiogenic growth factors,^{3,5,6} and matrix metalloproteinases.⁷

A large number of angiogenic mediators may contribute to the repair and regeneration of the endometrium although none have been clearly demonstrated to play a role in vivo.⁶ The angiopoietin family of growth factors are ligands for the largely endothelial-restricted Tie-2 receptor tyrosine kinase which is essential for vascular development.^{8–12} Mice deficient in *tie*-2 die *in utero* by embryological day E10.5 indicating that Tie-2 is required for the remodeling and maintenance of the primary vascular network during development.^{9,13–15} Ang-1 binds and induces autophosphorylation of Tie-2,10 promoting endothelial cell migration, ¹⁶ sprouting, ^{17,18} and survival ^{19,20} in vitro; and increases new vessel growth, branching, maturation, and integrity in vivo.21,22 Ang-1 secreted by perivascular cells is thought to promote the production of smooth muscle cell mitogens such as platelet-derived growth factor and heparin-binding epidermal growth factor in the endothelium.²³ In contrast, Ang-2 acts as a natural antagonist of Tie-2 signaling leading to vessel destabilization and neovascularization in the presence of vascular endothelial cell growth factor (VEGF), or vascular regression in its absence.²³ Supporting their opposing activities, disruption of the ang-1¹¹ or over-expression of ang-2¹² in transgenic mice results in death in utero due to

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a broad failure of vascular morphogenesis with a similar phenotype to *tie*-2 ^(-/-) knock-out mice.^{9,13–15} Thus the angiopoietins appear to play a key role in coordinating new blood vessel growth sending permissive signals via Tie-2 to either promote vascular maturation/integrity, or destabilize vessels leading to angiogenesis or vascular regression depending on the molecular context in which they act.²³

It has been suggested that disturbances of the normal processes of angiogenesis occurring in the endometrium during menstruation may result in menorrhagia.5,24,25 Although there is little direct evidence to support this hypothesis, increased endothelial cell turnover and multiple abnormalities in endometrial microvascular morphology have been noted in patients with menorrhagia.^{24,25} Moreover, any perturbation in the development of endometrial spiral arterioles (through which the majority of menstrual blood loss occurs) might result in inadequate vasoconstriction and repair leading to menorrhagia. In this study we examined the expression and distribution of Ang-1, Ang-2, and Tie-2 at a mRNA and protein level in both normal and menorrhagic endometrium during the proliferative and secretory phases of the menstrual cycle. We found that the normal pattern of Ang-1 expression is down-regulated in the endometrium of women with menorrhagia. Our observations indicate that Ang-1 may be involved in endometrial regeneration by stabilizing newly formed blood vessels and that loss of normal Ang-1 expression may lead to inadequate vascular remodeling/ maturation and excessive blood loss observed in menorrhagia.

Materials and Methods

Reagents

Rabbit polyclonal antibodies against human Ang-1, Ang-2, and Tie-2 cDNA constructs were kindly provided by Dr. George Yancopoulos, (Regeneron Pharmaceuticals, Tarrytown, NY). Rabbit anti-Tie-2 polyclonal antibodies (sc-324) were purchased from Santa Cruz Biotechnology, Inc. (Autogenbioclear, Wilts, UK). All other reagents were obtained from Sigma (Poole, Dorset, UK) unless stated otherwise.

Tissue Collection

Endometrium was obtained from patients 25 to 48 years of age under ethical approval from the South Birmingham Ethical Committee. Normal endometrium was obtained from 30 patients undergoing hysterectomy for benign conditions or dilatation and curettage for laparoscopic sterilization. Endometrium was collected by Pipelle biopsy from 28 patients who were referred from their general practitioners complaining of menorrhagia as described previously.⁵ These patients had regular menstrual cycles, had not had recent pregnancies, or received any hormonal treatments for the 3 months preceding endometrial biopsy. Tissues were rinsed in sterile saline and immediately snap frozen in liquid nitrogen.

Endometrial samples were dated from the last menstrual period and the stage of cycle confirmed by independent histological analysis using the Fox and Buckley criteria²⁶ as described previously.⁵

Ribonuclease Protection Assay

Total RNA was isolated from snap-frozen endometrial tissue following homogenization by the method of Chomczynski and Sacchi.²⁷ Ang-1 (570 bases), Ang-2 (388 bases), and Tie-2 riboprobes were transcribed from cDNA templates in the presence of $\left[\alpha^{-32}P\right]$ -UTP (Amersham Pharmacia Biotech, High Wycombe, Bucks, UK) as described previously.²⁸ Ang-1 and Ang-2 (1.0 to 5.0 \times 10^5 cpm) probes and a 127 base β -actin (1.0 to 5.0 \times 10⁴ cpm) control probe (Ambion, Austin, TX) were combined with 10 μ g of total RNA and co-precipitated. Ribonuclease protection assays were then performed using the RPA II kit according to the manufacturer's instructions (Ambion) and the protected fragments resolved by 6% denaturing polyacrylamide gel electrophoresis (PAGE). The Ang-1, Ang-2, and Tie-2 protected fragments were quantified from autoradiographs using the UVP Gelbase densitometry software and the results normalized to β -actin or 28s RNA respectively.

Western Blotting

Endometrial tissue lysates were prepared and subjected to Western blot analysis as described previously.^{5,29} Briefly, samples (50 μ g) were separated by 8% sodium dodecyl sulfate (SDS) polyacrylamide gel electrophoresis (SDS-PAGE) and semidry electroblotted onto Hybond-C nitrocellulose (Amersham, Pharmacia Biotech). The blots were blocked in 1% (w/v) milk protein and incubated with a 1:1000 dilution of either anti-Ang-1, Ang-2, or Tie-2 polyclonal antibodies overnight at 4°C. Antibody binding was detected using a goat anti-rabbitperoxidase conjugate and enhanced-chemiluminescence (ECL detection kit; Amersham Pharmacia Biotech), and quantified by laser densitometry.

In Situ Hybridization

Plasmids containing human Ang-1, Ang-2, and Tie-2 cDNAs were linearized and used as a templates to generate digoxigenin-11-UTP-labeled riboprobes using the RNA color kit (Amersham Pharmacia Biotech) and *in situ* hybridization performed as described previously.²⁸ Briefly, sections were hybridized with digoxigenin-labeled riboprobes at 55°C overnight. The sections were incubated with sheep anti-digoxigenin antibody for 3 hours and probe-binding detected using 5-bromo-4-chloro-3 indoyl phosphate/nitro blue tetrazolium (BCIP/NBT) at 4°C for 24 to 48 hours.

Immunohistochemistry

Immunohistochemistry was performed as described previously.³⁰ Serial $3-\mu$ m formalin-fixed, wax-embedded

sections were incubated with a 1:100 dilution of either anti-Ang-1, Ang-2, or Tie-2 polyclonal antibodies for 1 hour. Antibody binding was detected using biotinylated goat anti-rabbit secondary antibody, streptavidin-biotinperoxidase complex (ABC Kit; DAKO Ltd., Bucks, UK), and diaminobenzidine, and the sections were then counterstained with hematoxylin. In control sections the primary antibody was replaced with non-immune rabbit immunoglobulin or omitted.

Results

Quantitation of Ang-1, Ang-2, and Tie-2 mRNA in Normal and Menorrhagic Endometrium

Ribonuclease protection analysis was used to quantify total ang-1, ang-2, and tie-2 expression in endometrial samples isolated from normal and menorrhagic endometrium. The levels of Ang-1, Ang-2, and Tie-2 mRNA detected in endometrial tissue relative to *B*-actin mRNA or 28s rRNA controls, respectively, and typical results from these assays are shown in Figure 1, A-D. Although there were variations in ang-1 and ang-2 expression between tissues isolated from different individuals, levels of Ang-1 mRNA were found to be consistently lower than Ang-2 mRNA detected in normal cyclic endometrium (Figure 1, A and B). Tie-2 mRNA was detected in the majority of endometrial samples analyzed and similarly showed interindividual variation (Figure 1, C and D). In contrast to normal endometrium, total ang-1 expression was broadly down-regulated in samples of menorrhagic endometrium from both the proliferative and secretory phases of the menstrual cycle, while Ang-2 mRNA was detected at similar levels to those observed in normal endometrium. Conversely, there was generally elevated Tie-2 mRNA expression in endometrial samples from menorrhagia patients (Figure 1, C and D).

Quantitation of Ang-1, Ang-2, and Tie-2 Protein in Normal and Menorrhagic Endometrium

Endometrial tissue was pooled from five normal or menorrhagic women for each phase of the menstrual cycle and 50 μ g of total protein subjected to Western blot analysis for Ang-1, Ang-2, and Tie-2. In normal endometrium, Ang-1 and Ang-2 were detected as \sim 70 kd bands in all phases of the cycle examined with maximal levels detected in early secretory endometrium (Figure 1, E and F). Consistent with the observed decrease in Ang-1 transcripts (Figure 1, A and B), Ang-1 protein was significantly reduced in menorrhagic compared with normal endometrium in both the proliferative (P < 0.05) and secretory (P < 0.01) phases of the cycle (Figure 1, E and F). Whereas an increase of total Ang-2 protein in proliferative endometrium was evident in menorrhagia compared with normal cyclic endometrium (Figure 1E). Moreover, the overall ratio of Ang-1 to Ang-2 was markedly reduced in menorrhagic endometrial tissue in all phases of the cycle (Figure 1F).

Tie-2 was detected as two bands, at ~135 kd and ~140 kd, which probably correspond to cytoplasmic precussor and fully mature forms³¹ of the receptor respectively as previously reported with this antibody³² (Figure 1E). Tie-2 was expressed throughout the proliferative and secretory phases of the cycle. In contrast to the observed general increase in Tie-2 mRNA levels the relative level of Tie-2 receptor detected in menorrhagic endometrium was reduced in the proliferative and late secretory phases of the cycle in comparison with normal endometrium (Figure 1E).

Localization of Ang-1, Ang-2, and Tie-2 mRNA in Endometrium during the Menstrual Cycle

In situ hybridization was used to determine the distribution of mRNA in human endometrium. Ang-1, Ang-2, and Tie-2 expression showed cyclical changes during the menstrual cycle (Figure 2). Ang-1 mRNA was most abundant in the endometrial stroma and glandular epithelium of the early and mid-proliferative menstrual phases (Figure 2A) and decreased in the late proliferative phase of the cycle. Although we did not detect significant Ang-1 mRNA in the blood vessels during the proliferative phases it was predominantly expressed in the stroma surrounding the blood vessels (Figure 2B). In addition to the endometrial stroma, *ang*-1 expression was also present at lower levels in the glandular epithelium during the secretory phases (Figure 2B).

Ang-2 mRNA was detected in normal endometrial tissue throughout the menstrual cycle. In the proliferative phases moderate staining was detected in the glands and stroma (Figure 2E). In secretory endometrium, Ang-2 mRNA was also observed in the glands and stroma with increased expression in mid- to late secretory phases of the cycle (Figure 2F).

Tie-2 expression was detected mainly in the blood vessels and glandular epithelium of normal endometrium. In the early proliferative phase (Figure 2G) of the cycle, intense *tie-2* expression was observed in the glandular epithelium which decreased in mid- and late proliferative phases. Moderate Tie-2 mRNA hybridization signal was observed in the glands during the early secretory phase of the cycle. While in the early and mid-secretory phases *tie-2* was strongly expressed in endothelium and glandular epithelium (Figure 2), reduced *tie-2* hybridization signal was detected in the late secretory phase of the cycle.

In menorrhagic endometrium *ang*-1 expression was notably reduced or absent from the proliferative and early secretory phases and absent in the majority of specimens examined from mid- and late secretory phases of the menstrual cycle (Figure 2C). In contrast there were no consistent observable differences in the overall distribution and expression of Tie-2 and Ang-2 mRNA in the samples of menorrhagic endometrium examined. No hybridization was detected in control sections hybridized with sense riboprobes to Ang-1, Ang-2, and Tie-2 (Figure 2D).



Figure 1. Quantitative analysis of Ang-1, Ang-2, and Tie-2 expression in normal and menorrhagic endometrium. **A:** A representative RNase protection autoradiograph showing the relative expression of Ang-1 and Ang-2 mRNA expression in normal⁹ and menorrhagic¹¹ endometrium samples. Endometrial samples were collected at the early proliferative (EP), mid-proliferative (MP), late proliferative (LP), early secretory (ES), mid-secretory (MS) and late secretory (IS) phases of the menstrual cycle. Total RNA was extracted from endometrial tissue and analyzed by RNase protection for the presence of Ang-1, Ang-2, and β -actin (control) mRNA. **B:** Densitometric analysis was performed on the autoradiographs and the densities of the Ang-1 and Ang-2 protected fragments normalized to the internal β -actin control fragments and represented graphically as relative density units. **C:** *Tie-2* expression in endometrial samples by Western blot analysis. Protein extracts (50 µg) pooled from endometria tissue from 5 individuals taken at early (E), mid- (M), and late (L) proliferative (*pro.*) and secretary (*sec.*) phases of the menstrual cycle were subjected to Western blotting and enhanced chemiluminescent detection with antibodies against Ang-1, Ang-2, and Tie-2. **F:** Densitometric analysis of Ang-1 and Ang-2 expression was performed on the luminographs and the results expressed as the ratio of Ang-1, Ang-2, and Tie-2. **F:** Densitometric endometrial tassue from 5 function and the results expressed as the ratio of Ang-1, Ang-2, and Tie-2. **F:** Densitometric endometric endometric analysis of Ang-1 and Ang-2 expression was performed on the luminographs and the results expressed as the ratio of Ang-1, Ang-2 for normal and menorrhagic endometric metric endometric endometric the cycle.

Immunolocalization of Ang-1, Ang-2, and Tie-2 during the Menstrual Cycle

We examined the distribution of Ang-1, Ang-2, and Tie-2 receptor in normal endometrium during the menstrual

cycle and compared this with staining in endometrial samples from menorrhagic women (Figure 3). Perinuclear staining for Ang-1 was detected in the stromal cells throughout the menstrual cycle. Ang-1 was also detected in the glandular epithelium and blood vessels and, in



Figure 2. Localization of Ang-1, Ang-2, and Tie-2 transcripts in human endometrium from 30 normal (**A, B, D–H**) and 28 menorrhagic (**C**) individuals. *In situ* hybridization was performed on endometrial tissue sections using digoxigenin-labeled antisense riboprobes for Ang-1, Ang-2, and Tie-2. **A:** Strong Ang-1 hybridization signal was detected in the stroma (S) and glandular epithelium (G) of early proliferative phase endometrium (magnification, $\times 200$). **B:** *Ang*-1 was highly expressed in the stroma surrounding blood vessels (bv) in secretory phase normal endometrium (magnification, $\times 100$). **C:** Weak/absent Ang-1 hybridization signal in secretory phase menorrhagic endometrium (magnification, $\times 100$). **D:** No hybridization was observed on control sections hybridized with sense Ang-1 riboprobe (magnification, $\times 200$). **E:** Detection of Ang-2 mRNA in the glandular epithelium and stroma in proliferative phase endometrium (magnification, $\times 100$). **C:** Weak/absent Ang-1 (S) strong expression of *tie-2* was present in glandular epithelium and blood vessels of mid-secretory endometrium (magnification, $\times 200$). **G:** Strong expression of *tie-2* was present in glandular epithelium and blood vessels in late secretory (magnification, $\times 100$) and **H,** mid-secretory endometrium (magnification, $\times 200$).

particular, the endothelium of small venules (Figure 3A). In the secretory phase there was strong expression of Ang-1 in the perivascular stroma with low level expression in the glandular epithelium (Figure 3B). Intense Ang-2 expression was detected in the glandular epithelium and weaker staining of the blood vessels and stroma in proliferative endometrium (Figure 3F). Ang-2 exhibited a similar biphasic pattern of staining to that observed for Ang-1 with reduced Ang-2 staining of the decidualized stroma surrounding the spiral arterioles during the secretory phase.

Immunostaining for Tie-2 was present in the endothelium of blood vessels and also in the glandular epithelium throughout the menstrual cycle. Generally the distribution



Figure 3. Immunohistochemical localization of Ang-1, Ang-2, and Tie-2 in 30 normal (**A**, **B**, **F–H**) and 28 menorrhagic (**C–E**) endometrium specimens taken throughout the menstrual cycle. **A:** In proliferative endometrium, Ang-1 staining was detected in the glandular epithelium (G), stroma (S) and blood vessels (magnification, $\times 200$). **B:** Intense Ang-1 staining was observed in the stroma surrounding the blood vessels (bv) although it was very low or absent in the glandular epithelium of normal secretory endometrium (magnification, $\times 100$). **C** and **D:** The presence of low levels of Ang-1 in glandular epithelium and endothelium in the secretory menorrhagic endometrium of two patients (magnification, $\times 200$). **E:** Control section demonstrating a lack of background staining in sections in which the primary anti-Ang-1 antibody was replaced with rabbit serum. **F:** Strong Ang-2 immunoreactivity was detected in the glandular epithelium of proliferative endometrium (magnification, $\times 100$). **G:** Moderate Tie-2 staining was detected in glandular epithelium with weak staining of some blood vessels in proliferative endometrium (magnification, $\times 100$). **H:** In mid-secretory endometrium strong Tie-2 staining was present in the glandular and squamous epithelium and blood vessels with some staining of stroma (magnification, $\times 100$).

of Tie-2 produced an inverse pattern of staining for the Ang-1 with pronounced staining of the endothelium and epithelium (Figure 3G). There was clear membranous staining of the glands which became concentrated in the apical portions of the glandular epithelial cells during the secretory phase (Figure 3H).

In menorrhagia Ang-1 staining was markedly reduced compared with normal endometrium. Weak staining for Ang-1 was observed in the apical portions of the glandular and squamous epithelium with reduced stromal staining during the secretory phase of the menstrual cycle compared with normal endometrium (Figure 3B) (Figure 3, C and D). Ang-1 was also detected on the endothelium of blood vessels during the secretory phase of the cycle (Figure 3B).

Discussion

Normal menstruation is characterized by rapid regression of endometrial thickness, followed by intense vasoconstriction of the spiral arterioles and subsequent bleeding from focal points. A rapid and highly coordinated vascular response is required for endometrial repair and regeneration following menstruation. In this study, we have examined the expression Ang-1, Ang-2, and their cognate receptor, Tie-2, in normal and menorrhagic endometrium during the menstrual cycle. In endometrium of patients suffering from menorrhagia there was a marked down-regulation of both Ang-1 mRNA and protein while the expression of Ang-2 increased or remained at similar levels resulting in a large decrease in the ratio of Ang-1 to Ang-2.

Tie-2 and its ligands were expressed in a coordinated manner in endometrial tissue throughout the menstrual cycle with marked differences in the cellular distribution of Ang-1 and Ang-2. Consistent with its largely endothelial-restricted expression and recent studies of endometrium,33,34 Tie-2 was detected in the endometrial blood vessels. However, we also observed Tie-2 mRNA and protein in the glandular epithelium of the endometrium. Similarly, Tie-2 expression was recently reported in the epithelium of the thyroid.³² Tie-2 and its activating ligand, Ang-1, were highly expressed in the blood vessels and associated stroma of early proliferative phase endometrium, where the levels of Ang-2 were relatively low. The expression of Ang-1 mRNA predominantly in the stroma surrounding blood vessels and detection of Ang-1 protein on the endothelium of these blood vessels supports the paracrine mode of action of this ligand-receptor system. 10,11,21,23 Moreover, it suggests that they play a role in the neovascularization of early proliferative endometrium, which coincides with the initial distinct phase of angiogenesis in human endometrium.35 Many angiogenic factors implicated in endometrial development such as basic fibroblast growth factor (bFGF), VEGF, and platelet activating factor are reported to be highly expressed during the late proliferative stage.⁶ However, we found that Ang-1, Ang-2, and Tie-2 decreased in the late proliferative phase of the cycle. It is suggested that the major mechanism of new vessel growth in the endometrium is intussuception and/or vessel elongation rather than classical sprouting angiogenesis.⁶ The critical involvement of the Tie-2 receptor in the process of intussusception has been demonstrated in *tie*- $2(^{-/-})$ knockout mice.14

The angiopoietins are thought to coordinate angiogenesis by providing permissive signals that result in the stabilization of newly formed blood vessels by mediating perivascular cell (pericyte and smooth muscle cell) migration and the formation of basement membranes.^{11,23} That Ang-2 mRNA is only readily detected in the uterus, ovary, and placenta, tissues that are undergoing vascular remodeling in the adult, highlights its importance in blood vessel development.¹² In agreement with Li and colleagues³³ we found Ang-2 mRNA expression to be consistently greater than Ang-1 throughout the cycle in normal endometrium. However, we detected Ang-2 mRNA and protein in the glandular epithelium, blood vessels, and associated stroma, whereas Li and colleagues³³ only detected Ang-2 mRNA in natural killer cells.

The observed increase in Ang-1 expression in the perivascular stroma during the early secretory phase indicates that the angiopoietin/Tie-2 system is important for endometrial vessel development during the post ovulatory phase when the spiral arterioles are developing.^{6,36} Human endometrium expresses angiogenic factors, including bFGF and VEGF,3,5,6 that are reported to upregulate Tie-2 expression in cultured endothelial cells.³⁷ The fundamental role of angiogenesis in the development of the endometrium was demonstrated with the angiogenesis inhibitor, AGM-1470, which suppressed endometrial maturation in mice.³⁸ Normal endometrial angiogenesis is perturbed in menorrhagia^{24,25} and there is an increase in the number of small venules in the deep endometrium and inner myometrium in women with menorrhagia.²⁵ In the endometrium of women with menorrhagia, we observed decreased levels of Ang-1 and Tie-2 receptor that may contribute to disturbed endometrial vascular remodeling in this condition. Endothelial cell proliferation is reported to be significantly greater in menorrhagia without significant change in the overall endothelial cell density²⁴ implying a continual process vascular remodeling in menorrhagic endometrium throughout the menstrual cycle. This is consistent with our findings of a greater than 50% decrease in the ratio of Ang-1 to Ang-2 in menorrhagia and the proposed function of Ang-2 as a promoter of vessel destabilization.^{12,23}

In menorrhagia, the rate of blood loss increases significantly and menstrual bleeding persists on average 24 hours longer suggesting that repair of the endometrial blood vessels is compromised.^{2,36} The majority of menstrual blood loss occurs through the spiral arterioles that develop from endometrial arterioles during the secretory phase of the cycle.^{6,36} There appears to be no significant difference in the overall number of spiral arterioles in menorrhagic compared with normal endometrium.³⁶ However. recent evidence suggests that the phenotype of the smooth muscle cells in the spiral arterioles is altered in idiopathic menorrhagia without observable difference in the overall number of smooth muscle cells associated with these vessels. The proliferation of spiral arteriole smooth muscle cells is significantly reduced in the mid- to late secretory phases in menorrhagic endometrium.³⁶ It is therefore possible that the reported change in smooth muscle cell phenotype is a direct result of the altered balance of Ang-1 and Ang-2 levels leading to destabilization of the spiral arterioles in menorrhagia. It is clear, however, that the demonstration of ang-1 gene expression in the human endometrium and its consistent down-regulation in menorrhagia establishes this family of angiogenic growth factors as important regulators of uterine function.

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