Notwithstanding these important unresolved questions, the Zhu et al. (5) study should open up a new area of investigation as additional studies probe the connections between β1-adrenergic receptor–mediated signaling and CaMKII-dependent effects. If future studies support the proposal by Zhu et al., that one of the consequences of prolonged β1-adrenergic receptor stimulation is cardiomyocyte cell death via a CaMKII-mediated pathway, it will indeed be an important contribution to our understanding of the pathogenesis of heart failure that will suggest novel therapeutic targets.


See the related articles beginning on pages 649 and 707.

**Soluble VEGF receptor Flt1: the elusive preeclampsia factor discovered?**

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The occurrence of seizures (eclampsia, from the Greek “eklampsia,” sudden flashing) has been a long-known and feared complication of pregnancy, often killing both mother and child. Preeclampsia, or the condition preceding full-blown eclampsia, affects up to 5% of pregnant women and is diagnosed by the onset of hypertension and proteinuria in the second trimester (1). Preeclampsia may eventually progress to glomerular malfunction, thrombocytopenia, liver and brain edema, and associated life-threatening seizures (2) (Figure 1). Preeclampsia has sometimes termed the “disease of theories,” as several models for its pathogenesis have been proposed. But, as of today, no satisfactory unifying hypothesis has emerged (1). The restricted occurrence of preeclampsia to humans and primates and the lack of a suitable animal model have hampered the understanding of its pathogenesis (3). In this issue of the JCI, S.E. Maynard et al. (4) report the novel insight that circulating levels of two angiogenic growth factors, VEGF and placental growth factor (PIGF), may play a more important role than previously believed. In particular, the authors propose that, in pregnant women with preeclampsia, the placenta produces elevated levels of the soluble fms-like tyrosine kinase 1 (sFlt1) receptor, which captures free VEGF and PIGF. As a result, the normal vasculature in the kidney, brain, lungs, and other organs is deprived of essential survival and maintenance signals and becomes dysfunctional (Figure 1).

As the authors show in their rodent model, this may lead to the development of hypertension and renal disease, reminiscent of preeclampsia in humans. In another study in this issue, V. Eremina et al. (5) provide additional evidence for a critical role of VEGF in renal disease during preeclampsia. These authors demonstrate that mice lacking one VEGF allele in renal podocytes develop the typical renal pathology found in pregnant women with preeclampsia. These studies therefore shed unprecedented light on the pathogenesis of preeclampsia and offer novel therapeutic opportunities for this disease.

**sFlt1: a likely candidate preeclampsia factor**

For the fetus to develop normally, it must receive sufficient oxygen and nutrients (6). These are supplied via the maternal spiral arteries in the uterus. During normal pregnancy, cytotrophoblasts convert from an epithelial to an endothelial phenotype (a process termed pseudo-vasculogenesis) and invade maternal spiral arteries. This vascular remodeling increases the bulk flow and the supply of nutrients and oxygen to the fetus by the end of the first trimester (7, 8) (Figure 1). Vascular factors such as VEGF, angiopoietins, and ephrins have been implicated in this process (7). In preeclampsia, pseudo-vasculogenesis is defective, and the resultant placental ischemia has been proposed to trigger the release of unknown placenta-derived factors. The latter would induce systemic endothelial dysfunction and thereby contribute to the renal, cardiovascular, and neurological defects of preeclampsia (Figure 1). Despite intensive efforts, the precise nature of the placenta-derived...
preeclampsia factors has remained enigmatic for years.

In their study, Maynard et al. (4) may have discovered a likely candidate preeclampsia factor. Indeed, having used gene profiling, these authors report that the placenta of pregnant women with preeclampsia produced increased levels of sFlt1. Flt1, binding VEGF and its homologue PlGF, exists in two forms: a membrane-bound receptor tyrosine kinase which transmits angiogenic signals (Flt1), and a soluble secreted ectodomain which only captures VEGF and PlGF (sFlt1). As sFlt1 lacks a cytosolic domain, its function is restricted to regulating (reducing) the levels of free VEGF and PlGF available to signal via intact Flt1 and fetal liver kinase-1. Even though sFlt1 may be one of the few, if not the only, inhibitor of VEGF known today, it remained unknown whether it might contribute to any sort of human disease.

Altered angiogenic balance causes general endothelial dysfunction
How then can elevated circulating levels of sFlt1 contribute to preeclampsia? Whereas dynamic surges of high VEGF levels mediate angiogenesis in the embryo and in the adult during disease, continuous low levels of VEGF are required for endothelial cells to survive prolonged periods and function properly. Thus, when sFlt1 plasma levels rise, they may reduce the circulating VEGF and PlGF levels below a critical threshold required for maintenance of the established vasculature in the adult. The resultant endothelial dysfunction may disrupt the blood-brain barrier and cause intracranial hypertension, result in edema in the liver, and affect glomerular function. Indeed, Eremina et al. (5) demonstrate that glomerular capillary function is under the strict gene-dosage–dependent control of VEGF. That is, when VEGF levels in the renal prodocytes drop 50%, glomerular endothelial cells swell, capillary loops collapse, and proteinuria develops—as in preeclampsia. We previously demonstrated that absence of the VEGF164 and VEGF188 isoforms impairs glomerular filtration (9). Endothelial dysfunction may also deregulate hemostasis and trigger thrombocytopenia. By inducing vasoconstriction, VEGF also induces hypotension, and thus lower circulating VEGF levels will cause elevated blood pressure, another hallmark of preeclampsia. Thus, the sFlt1 hypothesis allows the proposal of a unifying model, explaining, perhaps not all, but at least several, of the hallmark symptoms of preeclampsia.

As insightful as these studies are, they do not address the question as to what upregulates sFlt1 expression in the placenta in preeclampsia. Hypoxia upregulates Flt1 expression and, since the placenta is hypoxic in pregnant women with preeclampsia, could be responsible (Figure 1). It also remains to be determined whether the elevated sFlt1 levels in the placenta would not further aggravate the disease by impairing VEGF-dependent pseudo-vasculogenesis and maternal spiral artery remodeling.

What medical implications do these studies suggest? First, strategies designed to normalize circulating free VEGF and PlGF levels might be expected to halt progression of the disease. Second, chronic administration of VEGF inhibitors, currently being evaluated for their therapeutic potential to inhibit cancer, retinal neovascularization, and chronic inflammation, may pose a risk of endothelial dysfunction, renal insufficiency, and hypertension. Results from ongoing clinical trials support this notion. Third, renal parameters and blood pressure might be used as surrogate markers of the success of anti-angiogenesis therapy—the angiogenesis field desperately needs such markers. At the very least, these studies raise new hopes that novel strategies may be designed to combat preeclampsia and thereby improve the chances of the mother and child for a healthy future.
Over 65 years ago, A. Randall demonstrated that interstitial crystals located in the papillary interstitium, the site of initial solid phase, the site of initial crystallization would account for growth into larger, clinically significant, calculi. Randall’s plaques, were common in most cases of nephrolithiasis and are associated with the relatively common metabolic abnormality of idiopathic hypercalciuria (1). These patients generally absorb an excess amount of dietary calcium leading to increased urine calcium excretion and supersaturation with respect to calcium oxalate and calcium phosphate; they subsequently form stones. Other patients with nephrolithiasis, who have had an intestinal bypass procedure, absorb oxalate in excess leading to increased urine oxalate excretion and supersaturation with respect to calcium oxalate; they also subsequently form stones. In these and other causes of nephrolithiasis, the site of the initial solid phase has long been the subject of debate. Over 65 years ago, A. Randall demonstrated that interstitial crystals located at, or adjacent to, the papillary tip, Randall’s plaques, were common in most cases of nephrolithiasis. He found that these crystals were composed not of calcium oxalate, the most common solid phase found in patients with nephrolithiasis, but of calcium phosphate (3). He believed that the calcium phosphate crystals formed in the papillary interstitium and then eroded into the urinary space, serving as a heterogeneous nucleation surface for calcium oxalate. B. Finlayson later argued that, due to rapid flow of the renal ultrafiltrate through the tubule, there was insufficient time for formation of a lumen-obstructing solid phase (4), which also suggested that an intratubular site of stone formation was unlikely. However, other investigators found that calcium oxalate crystals adhered to cultured tubular cells (5), where they could either be endocytosed or remain on the cell surface, serving as a nidus for growth into larger, clinically significant, calculi.

Site of the initial solid phase

Where is the site of initial crystallization — the interstitium, the tubular lumen, or perhaps the renal calyx, where supersaturated fluid awaits excretion into the ureter? Knowing the site of initial crystallization would improve understanding of the pathogenesis of stone formation and allow investigators to propose and test more focused hypotheses. This would help them to devise effective therapy aimed at preventing recurrent nephrolithiasis, which affects approximately 50% of stone formers within five years of the initial stone (6). Yet until the elegant study by A.P. Evan et al. reported in this issue of the JCI (7), we did not have an answer to this rather elementary question. These investigators performed kidney biopsies on stone-forming patients to determine the anatomical site and composition of the initial solid phase. They sampled areas adjacent to Randall’s plaques in patients undergoing percutaneous nephrolithotomy. In hypercalciuric calcium oxalate stone formers, they found initial calcium phosphate (apatite) crystallization in the basement membrane of the thin limbs of the loop of Henle (Figure 1) with subsequent extension to the vasa recta, then to the interstitial tissue surrounding the terminal collecting ducts, and finally, in the most severe cases, to the papillae. Erosion of this solid phase into the urinary space, which is supersaturated with respect to calcium oxalate, may have promoted heterogeneous nucleation and formation of kidney stones. In patients with hyperoxaluria resulting from intestinal bypass, the initial crystals were again a calcium phosphate complex, but these arose within the tubule lumens of terminal collecting ducts (Figure 2). Contact of these crystals with urine, supersaturated with respect to calcium oxalate, may have promoted heterogeneous nucleation and formation of kidney stones. Non-stone formers, subjected to nephrectomy, had neither plaque nor crystals. Thus there are different sites of initial

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See the related article beginning on page 607.

Nephrolithiasis: site of the initial solid phase

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