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The Intersection of Maternal Metabolic Syndrome, Adverse Pregnancy Outcomes, and Future Metabolic Health for the Mother and Offspring

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Abstract

The prevalence of obesity is ~40% in the United States, and the prepregnancy prevalence of obesity in females is ~30%. This has in part fueled an increase in metabolic syndrome (MetS) among females who are currently pregnant, have been pregnant, or are planning to become pregnant. Importantly, MetS in pregnancy is associated with increased pregnancy complications. Moreover, MetS in pregnancy may have long-lasting adverse cardiovascular and metabolic health implications for the mother and her offspring. To complicate matters, many adverse pregnancy outcomes seem to increase the risk of MetS in the mother after pregnancy. Herein, we describe the potential mechanisms behind the intersection of MetS, adverse pregnancy outcomes, and subsequent long-term disease in the mother and offspring. Because MetS is a cluster of coexisting conditions, it is challenging to identify mediators that can serve as biomarkers for early diagnosis and targets for MetS prevention and therapy.

Keywords: metabolic syndrome, pregnancy, pre-eclampsia, gestational diabetes, placenta

METABOLIC SYNDROME (MetS) is defined as a cluster of metabolic abnormalities such as obesity, insulin resistance, hypertension, dyslipidemia, and diabetes; it is characterized by chronic low-grade systemic inflammation and directly promotes the development of atherosclerotic cardiovascular disease (CVD) and type 2 diabetes (T2D).¹ A significant increase in MetS across the globe and among females who are currently pregnant, have been pregnant, or are planning to become pregnant has been reported,² and MetS has been associated with adverse pregnancy outcomes such as miscarriages, preterm birth (PTB), pre-eclampsia (PE), gestational diabetes (GDM), and stillbirth.^{3–6}

That MetS has consistently been associated with increased risk for CVD is not surprising; however, pregnancy complications such as PE and GDM, frequently associated with MetS, have also been associated with increased maternal risk for CVD.^{7,8}

Because MetS is a cluster of coexisting conditions, it is challenging to understand its pathophysiology. In addition, the metabolic adaptations that occur during pregnancy com-

PLICATE even more the identification of mediators that can serve as biomarkers for early diagnosis and targets for MetS prevention and therapy. Importantly, the consequences of MetS in pregnancy affect both the mother and the offspring; a transgenerational inheritance of MetS has also been postulated. Therefore, early diagnosis of MetS may be useful to screen women at higher risk for pregnancy complications, which may additionally associate with future risk of CVD and adverse health outcomes in the offspring.

Metabolic Adaptations During Pregnancy May Increase the Risk of MetS

Pregnancy is associated with important physical, metabolic, hormonal, vascular, and immunological changes leading to a pro-inflammatory, prothrombotic, highly insulin resistant, and hyperlipidemic state.³ These adaptations, necessary for the development of the fetus and to prepare the mother for delivery and lactation, can also increase the susceptibility to develop MetS. In the early stages of pregnancy, the

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increased levels of estrogen, progesterone, and cortisol favors lipogenesis and accumulation of fat.⁹ Then, during mid-pregnancy, the diabetogenic effect of pregnancy on metabolism is noticeable when placental-derived hormones lead to an insulin-resistant state.⁹

Maternal MetS and Adverse Pregnancy Outcomes: A Reciprocal Relationship

It has been widely reported that a higher incidence of pregnancy complications, such as GDM and PE, occur in women with MetS.³⁻⁶ In addition, obese pregnant females have an ~25% higher risk of stillbirth and a higher risk of developing PE compared with nonobese pregnant females.^{3,5,7,8} Although increased serum glucose, triglyceride, and low-density lipoprotein-cholesterol levels and lower high-density lipoprotein-cholesterol levels have been observed in pre-eclamptic patients with MetS compared with pregnant individuals without MetS, it is not clear whether MetS is secondary to PE or whether PE plays a role in the pathogenesis of MetS. An increased risk of GDM has also been observed among women who are overweight or obese and the risk of developing GDM increases proportionally to the degree of obesity.¹⁰

In addition, overweight women are at a significantly higher risk of experiencing recurrent miscarriages compared with those of average weight.⁴ Similar to those in the non-pregnant state, maternal obesity is related to metabolic and cardiovascular complications of pregnancy, including PE and GDM.³ PE and CVD risk factors (including hypertension) are associated not only by common pathophysiology but also epidemiology.

For example, high blood pressure before pregnancy is a risk factor for PE and PE is in turn a risk factor for hypertension and CVD after pregnancy.¹¹ Moreover, an increased risk of PTB, especially extremely preterm delivery, has been reported among obese and overweight females.¹² Furthermore, up to 70% of females with GDM will develop T2D after pregnancy.¹³ Women with GDM also have a 2.4-fold higher risk of developing MetS after delivery compared with women with a normal pregnancy.¹³

Although pregnancies affected by MetS are associated with adverse pregnancy outcomes, an increased incidence of

MetS has been reported in women and their offspring after a complicated pregnancy; in particular, pregnancies in which an exacerbated pro-inflammatory state has been identified, such as PE and PTB.^{14,15}

MetS in pregnancy is associated with subsequent metabolic and CVD in the mother and offspring.¹⁶ Obesity is characterized by an increased inflammatory response, and fetuses of overweight and obese pregnant females also have shown a higher risk of MetS after birth.¹⁷

Potential Mechanisms

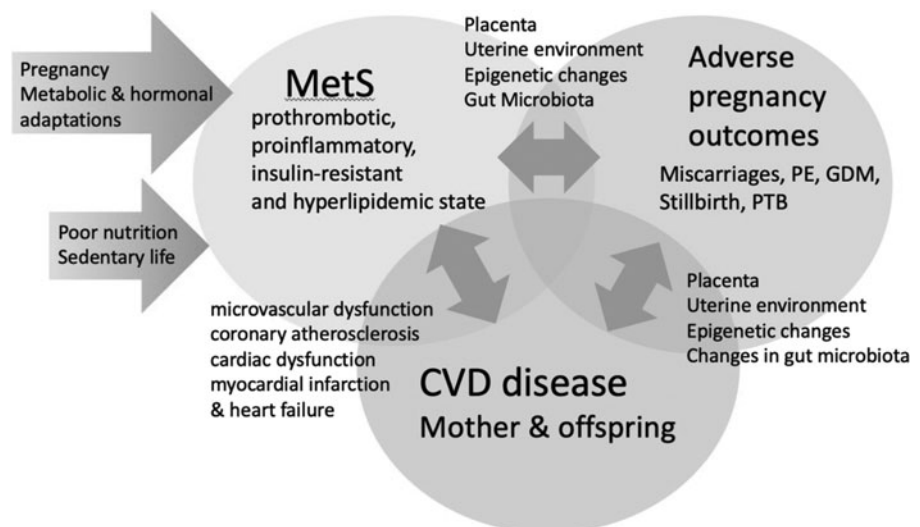
The mechanisms behind the cross-associations between MetS, CVD, and adverse pregnancy outcomes are difficult to differentiate due to the common risk factors. Roles for the intrauterine environment, epigenetic changes, the placenta, and changes in the gut microbiota have been postulated (Fig. 1).

Although a substantial body of evidence demonstrates that environmental exposures such as poor nutrition and sedentary lifestyle contribute significantly to the risk of MetS, there is growing evidence about the contribution of the intrauterine environment.¹⁸ That MetS is more frequently observed in those that were born from complicated pregnancies further supports the importance of exposures *in utero* to the development of MetS.^{14,15} Nutritional stress/stimuli during critical periods of early development can also impact metabolic programming¹⁹ contributing the etiology of chronic diseases such as obesity, T2D, and CVD later in life. Furthermore, it has been suggested that these effects may also affect subsequent generations, perpetuating MetS and CVD.

An adverse intrauterine environment can result in epigenetic programming, a critical underlying mechanism of fetal metabolic programming.¹⁹ Epigenetic variations established *in utero* can further be passed through multiple generations representing a plausible link between the *in utero* environment and later disease susceptibility.¹⁹

MicroRNAs regulate gene expression and may affect the development and functioning of the cardiovascular and endocrine systems. Interestingly, a postpartum profile of microRNAs associated with diabetes/cardiovascular/cerebrovascular diseases has been identified in women that

FIG. 1. Suggested mechanisms in the cross-association between MetS, adverse pregnancy outcomes (miscarriages, PE, GDM, stillbirth and PTB, and CVD), cardiovascular disease; GDM, gestational diabetes; MetS, metabolic syndrome; PE, pre-eclampsia; PTB, preterm birth.



experienced pregnancy complications such as gestational hypertension, PE, intrauterine growth restriction, and PTB.^{20,21} MiRNAs also play important regulatory roles in adipocyte differentiation, metabolic integration, insulin resistance, and appetite regulation, suggesting the potential use of miRNAs as novel biomarkers and therapeutic targets for MetS.²²

The placenta also plays an important role in the maternal metabolic adaptation of pregnancy and the modulation of the intrauterine environment. A negative effect of maternal obesity on placental development and function leading to impaired offspring programming and adverse health outcomes has been described. Specifically, maternal prepregnancy obesity is associated with a dysregulated placental transcriptome, particularly in pathways relating to inflammation, immune responses, glucocorticoid signaling, and angiogenesis.²³ Increased mRNA expression of pro-inflammatory cytokines and increased density of pro-inflammatory macrophages have also been found in placentas of obese females.²⁴

Placental dysfunction is commonly seen in females with obesity, chronic hypertension, diabetes, and dyslipidemia, all major features of MetS.²⁵ Maternal placental syndromes, such as PE and placenta abruption or infarction, are also more prevalent in females with insulin resistance, diabetes, and MetS.²⁵ Furthermore, placental insufficiency observed in PE has been associated with the postpartum development of MetS in females without prepregnancy metabolic disorders.²⁵

Animal models of placental insufficiency have shown an adverse metabolic and cardiovascular phenotype in the offspring.^{26,27} In one of these models, a role for soluble bioactive factors secreted by the placenta, vascular function modulators, markers of metabolic disease, vasoconstrictors, and pro-inflammatory mediators has been described in the metabolic and CVD observed in the mother and offspring.²⁶

Changes in maternal gut microbiota might also contribute to metabolic diseases, including GDM and obesity. The maternal microbiome during pregnancy may alter not only the developing neonatal microbiota but also intestinal absorption of nutrients and increased lipid deposition, potentially leading to MetS.²⁸ Changes in the gut microbiome composition and the metabolic hormonal environment has been observed in overweight and obese pregnant females and women with GDM.²⁹ Several bacteria have also been correlated with plasma glucose and blood lipid levels in females with GDM and hyperlipidemia.³⁰

Increased gut permeability may explain the role of the microbiota in the pathogenesis of GDM and MetS.³¹ Specifically, certain mucin-degrading bacteria that increase gut permeability have been detected in GDM,³¹ facilitating not only the increased absorption of nutrients rich in calories but also the movement of inflammatory mediators from the gut into the maternal circulation, potentially contributing to the development of MetS.

Conclusions

There are connections among maternal MetS, adverse pregnancy outcomes, and future metabolic health for the mother and offspring. Women with MetS during pregnancy are at higher risk for adverse short- and long-term outcomes; conversely, pregnancy complications associated with an increased pro-inflammatory state have been associated with an increased risk of MetS after pregnancy in mother and offspring.

Maternal lifestyle, the intrauterine environment, placental-derived bioactive molecules, microRNAs, the microbiome, and epigenetics may all play a role in the pathogenesis of MetS, the associated long-term health complications to the mother and offspring, and transgenerational inheritance of the syndrome. Future studies are required to address the overlapping/common mechanisms among MetS, adverse pregnancy outcomes, and the risk of future MetS and CVD in mother and offspring.

Disclaimer

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