Prenatal Nutritional High Fat Environment and the Evolving Concept of Intergenerational Cardio-Metabolic Disease Risk

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**Editorial**

The most rapid and dramatic consequence in the course of human history with the advent of the global economy are the changes in eating patterns, though undoubtedly fuelled by urbanization. Radical dietary shifts in many developed and developing nations are supplanting traditional patterns of eating with Western diet high in animal products, refined carbohydrates and low in whole grains, fruits, and vegetables. For example, in China, consumption of animal products increased by nearly 40% between 1989 and 1997 [1] and high fat (HF) food sales more than doubled between 1999 and 2005 [1]. The term “Coca-colonization,” a reference to the ubiquitous presence of Coca-Cola, Pepsi, and McDonald’s, describes a world that is moving toward a common diet, one accompanied by the more sedentary lifestyles associated with increased risk of chronic disease [2,3].

According to the World Health Organization's global database, countries like China, India and Pakistan have a preschool childhood obesity prevalence of about 1% [4]. Childhood obesity increases the risk of obesity in adulthood and parental obesity interacts quite strongly to alter this risk. There are several other interactive factors as well contributing to the increased prevalence of obesity in childhood. Societies like China, India and Pakistan, which are rapidly urbanising, demonstrate increases in energy intake, dramatic increases in fat intake along with increased levels of sedentarianism. Food balance data from the Food and Agriculture Organization show that the change in energy intake in Asian countries has been small, but there have been large changes in consumption of animal products, sugars and fats [5]. The net effect has been a marked shift in the diet with energy from fat (both animal and vegetable) increasing each year [5]. Data from India show that higher-income groups consumed a diet with 32% of the energy from fat while the lower-income groups consumed only 17% energy from fat [6-9].

Therefore, nutritional history of current populations becomes an important determinant of their health. There is a growing recognition of the importance of environmental factors acting on the genotype throughout the life cycle of an individual to progressively modify its phenotype. The periconceptional and intrauterine period seem to be the most crucial, when a small change in environment could have a large effect on the phenotype. The periconceptional and intrauterine period seem to be the most crucial, when a small change in environment could have a large effect on the phenotype. The periconceptional and intrauterine period seem to be the most crucial, when a small change in environment could have a large effect on the phenotype.

The basic principle of prevention of cardio-metabolic syndrome events consists of the control of classical risk factors with specific interventions. This clinical ‘high-risk’ approach has led to an improvement in prognosis, although still far below expectations. In other words, the mere control of hypertension, hypercholesterolemia, or obesity, though definitely beneficial, leaves a residual risk, which is still greater than that reduced by
treatment. Thus, it is necessary to identify windows of time in the lifecycle when the genome is highly susceptible to such influences. Such knowledge could lead to the discovery of different treatment targets that may have a bigger impact upon CVD burden. Any preventive intervention will therefore have to start in utero, and improving the health of young girls will be a very important aspect of such an approach.

The cardio-metabolic programming effect of maternal hypercholesterolemia and the observation of strikingly high foetal cholesterol levels at the end of 2nd trimester raise interest in maternal–fetal cholesterol transport. It is speculated that maternal cholesterol might cross the placental barrier such as in Smith Lemli Opitz syndrome where survival of foetuses depends on maternal cholesterol [15]. Identification of active transport mechanisms on the maternal side of the placenta in rabbits [16] and hamsters [17] has provided evidence for maternal–foetal cholesterol transport. The trans-placental cholesterol transport mechanism mediated by endothelial cells lining the placental villi into the lumen of foetal microvessels has recently been elucidated [18]. Results indicate that regulation of these mechanisms by increased foetal demand for cholesterol differs from that induced by maternal hypercholesterolemia [19]. Increased cholesterol transport, at least during parts of gestation may therefore constitute one mechanism by which maternal hypercholesterolemia affects foetal programming and hence requires further investigations.

Yet, epigenetic changes associated with increased disease susceptibility such as obesity, type-2 diabetes and cardiovascular dysfunction are emerging as critical components of enduring effects induced by early life experiences. The available data are now beginning to provide a molecular basis for involvement of oxidative stress in modifying biological properties of DNA methylation and other epigenetic effects [20]. It is reported that in utero conditions result in specific epigenetic change [21,22]. Developing organisms seem to have a wide window of susceptibility to epigenetic changes but the periconceptional period is particularly important towards sensitivity to suboptimal nutrition during this developmental stage in which widespread reprogramming of the epigenome occurs [23-25]. It is speculated that HF over-nutrition in pregnancy and/or postnatal could also lead to metabolic dysfunction in major organs such as kidneys and liver later on in life [26-30]. These epigenetic changes within suboptimal nutrition could have long-lasting effects on epigenomic methylation patterns in specific gene promotors. Moreover, the study of epigenetics will further guide us in understanding the hormonal regulation of enzymes controlling acetylation and methylation and sex differences across the epigenome.

Therefore, from all the discussion, it will be ideal to demonstrate (1) adverse effects of impaired early-life maternal HF nutrition associated epigenetic changes in offspring and (2) the mechanisms/impact of early interventions such as statins in mothers on HF diet for prevention or reversal of such epigenetic consequences during intrauterine development. Today, we have to recognize that it may be still too early to test these hypotheses in the clinical setting, yet the animal models have established cause-effect relationships for some maternal factors and led to the identification of specific questions in a high fat maternal environments. Despite this, mechanistic studies, both in animals and in humans, are needed in order to achieve a better understanding of the role of maternal high fat diet in the pathophysiology of cardio-metabolic syndrome. In addition, development of methods that would allow better characterization of this cardio-metabolic milieu in the humans is necessary. Only a more in-depth knowledge of these important pathophysiological mechanisms can allow the identification and testing of novel therapeutic strategies aimed at targeting the early developmental origins of cardiovascular disease in the populations.

So far from our own work [11,13], we have to some extent demonstrated in animal models the effects of long term maternal high fat diet and statin treatment on the offspring predisposition to early cardio-metabolic metabolic disease. However, much of our understanding of the disorder is far from complete. Future research originating from present work could focus on the translation of unexplored signalling pathways to answer the central question such as, how do such maternal high fat diets affect the long term cardiovascular health of offspring.

Throughout foetal development, the main nutritional source for the foetus comes from the mother. As a result, the quality of the maternal diet affects growth and development of the foetus during this period [31] though the mechanisms involved are poorly understood. Nevertheless, experimental animal models whereby the maternal diet is manipulated have provided insight into the links between in-utero environments and susceptibility to disease in later life (Figure 1). Rodent offspring from dams that were fed a protein-restricted diet during pregnancy later developed hypertension in adulthood [32]. However, it is fair to suggest that developmental programming of cardio-metabolic syndrome in adulthood is not limited to maternal dietary restriction alone. Ours and other groups have shown that feeding dams a high fat diet during pregnancy and/or lactation can also leads to the offspring developing vascular dysfunction and hypertension [11,13,33,34] and that changes in the maternal diet during pregnancy are linked with an increase in cardiomyocytes and foetal left ventricular mass [35].

Cardiomyocytes go through terminal differentiation at or after birth and this is characterised by the transition from hyperplastic growth to hypertrophic growth [36-38]. Many genes are reported to be involved in cell cycle progression and are associated with cardiac hypertrophy [39]. It remains however to be determined whether the hypercholesterolemic condition during pregnancy leads to any long term changes in these genes in the adult offspring’s heart; and whether the cholesterol-lowering effect of statin given to pregnant dams consuming a high fat diet influence the expression of these genes.

Conclusion

Recent studies have provided insights into the relationship between maternal nutritional status and intergenerational risk of cardio-metabolic disease. Evidence derived from animal models has concluded that high fat diet coupled with sedentary lifestyles provokes a cardio-metabolic trait whose effect transcends at least two offspring generations. However, much knowledge is still considered hypothetical in the absence of randomised clinical trials in the human. Development of interventions is limited by the absence of research at molecular/cell levels to identify potential genetic targets and at the human levels where many questions still remain answered.
**Figure 1:** The hypothesized "Hypercholesterolemic-mediated Teratogenesis" model. This model demonstrates on one extreme (right side) the risk of adult disease and on the other extreme (left side) range of postnatal physiological settings for survival fitness. At the x-axis (bottom) five different qualities of prenatal environments from very poor to very rich are presented. The red line curve represents the environment that the foetus anticipates postnatally judged from the nutritional and related signals it receives from the mother through the placenta. Provided that the postnatal environment matches the range for which the foetus has set its postnatal physiology by the processes of predictive adaptive responses, the disease risk is low. However, when it lies outside that range then the postnatal environment turns bad (indicated by the red arrow) and the disease risk increases. The foetus perceives the more restricted environment and predicts a poorer postnatal survival. But maternal high-fat dietary stress worsens the situation. Over-nutritional high fat environment prenatally and postnatally shifts the postnatal survival associated with increased disease risk further to the upper limit (represented with curved black line). However, administration of statin to mothers on high fat diet, even at rich nutritional upper limit, can provide a good safeguard from the disease process and maintains the survival fitness even if increasing affluence raises the richness of the mature postnatal environment.

**References**


