

Prenatal origins of adult disease

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Purpose of review

Human epidemiological and animal studies show that many chronic adult conditions have their antecedents in compromised fetal and early postnatal development. Developmental programming is defined as the response by the developing mammalian organism to a specific challenge during a critical time window that alters the trajectory of development with resulting persistent effects on phenotype. Mammals pass more biological milestones before birth than any other time in their lives. Each individual's phenotype is influenced by the developmental environment as much as their genes. A better understanding is required of gene–environment interactions leading to adult disease.

Recent findings

During development, there are critical periods of vulnerability to suboptimal conditions when programming may permanently modify disease susceptibility. Programming involves structural changes in important organs; altered cell number, imbalance in distribution of different cell types within the organ, and altered blood supply or receptor numbers. Compensatory efforts by the fetus may carry a price. Effects of programming may pass across generations by mechanisms that do not necessarily involve structural gene changes. Programming often has different effects in males and females.

Summary

Developmental programming shows that epigenetic factors play major roles in development of phenotype and predisposition to disease in later life.

Keywords

animal studies, developmental programming, human epidemiology

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Introduction

In the last 10–15 years there has been a growth of interest in the concept that many chronic adult conditions have their antecedents in abnormal fetal and early postnatal development. This concept has been termed developmental programming. Developmental programming can be defined as the response by the developing mammalian organism to a specific challenge during a critical time window that alters the trajectory of development qualitatively and/or quantitatively with resulting persistent effects on phenotype.

Developing organisms pass more biological milestones before birth than at any other time in their lives. It should therefore be no surprise that significant alterations in the timing or nature of these developmental steps have consequences in terms of organ function in later life. It also needs to be recognized that the development of each individual's specific phenotype, although based on a specific genotype, is influenced to a varying extent by epigenetic/environmental factors. Although an extreme example, there can be little dispute that the genome of

the fetus exposed to repeated and excessive alcohol will function very differently throughout life from the way it would have done in the absence of that exposure. Fetal alcohol syndrome represents the epigenetic effects on the fetal genome of excessive exposure to alcohol during development. Thus it is vitally important to understand the gene–environment interactions that lead to adult disease.

Acceptance and understanding of developmental programming comes from human epidemiological studies and a wealth of carefully controlled animal investigations primarily in rodents and sheep. There are numerous reviews on the exposures, mechanisms and outcomes involved [1–15]. In the early days of the concept of developmental programming several principles rapidly became clear. These are laid out in Table 1 [16]. For this review we will restrict our discussion to six of these principles – 1, 2, 4, 7, 9 and 10 – and focus on controlled animal investigations demonstrating that maternal undernutrition, overnutrition and stress, and fetal hypoxia, are all environmental factors that support the general principles of developmental programming.

Table 1 Ten principles of developmental programming [16]

Principle 1	During development, there are critical periods of vulnerability to suboptimal conditions. Vulnerable periods occur at different times for different tissues. Cells dividing rapidly at the time of exposure are at greatest risk. Risk factors include too much of a normal chemical such as a hormone, critical nutrient or vitamin; deficiency of a normal chemical such as a hormone, critical nutrient or vitamin; abnormal chemicals such as alcohol or nicotine; abnormal physical forces, such as high blood pressure.
Principle 2	Programming has permanent effects that alter responses in later life and can modify susceptibility to disease.
Principle 3	Fetal development is dependent on fetal physical activity. Normal development is dependent on continuing normal activity. Each phase of development provides the required conditions for subsequent development.
Principle 4	Programming may involve structural changes in important organs. The absolute numbers of cells in the organ may increase or decrease; the relative proportions and distribution of different types of cell within the organ may be unbalanced; the normal blood supply to the organ may be compromised; too many or too few hormone receptors may form with a resultant resetting of feedback and other control mechanisms.
Principle 5	The placenta plays a key role in some forms of programming.
Principle 6	Compensation carries a price. In an unfavorable environment, the developing baby makes attempts to compensate for deficiencies. Following compensation, birth weight may be normal or only slightly decreased. However, the compensatory effort carries a price.
Principle 7	Attempts made after birth to reverse the consequences of programming may have their own unwanted consequences. When postnatal conditions prove to be other than those for which the fetus prepared, problems may arise.
Principle 8	Fetal cellular mechanisms often differ from adult processes. Fetuses react differently to suboptimal conditions than do newborn babies or adults.
Principle 9	The effects of programming may pass across generations by mechanisms that do not necessarily involve changes in the genes.
Principle 10	Programming often has different effects in males and females.

Principles 1 and 2

Principle 1 is that there are critical periods of vulnerability to suboptimal conditions that occur at different times for different systems; Principle 2 is that there are permanent effects that alter responses in later life and can modify susceptibility to disease. It is impossible to separate these two principles. The first refers to the timing of exposure and Principle 2 to the outcome resulting from that exposure. Perhaps the best human example comes from the study of individuals who survived the Dutch Hunger Winter of 1944–1945 [17–21]. Children who passed any period of their prenatal development during that time suffered from poor nutrition. Epidemiological studies of these individuals, now over 60 years old, have demonstrated that undernutrition of their mothers at different stages of pregnancy produced different outcomes. For example, undernutrition in the last third of gestation decreased the percentage of the babies who become obese as adults whereas undernutrition during the first two trimesters increased the prevalence of adult obesity [22,23].

The idea that there are critical time windows when developing systems are most vulnerable to challenge, and the fact that these vulnerable periods differ among systems and species, is clearly supported by work in animals. Changes in activity of a given system across development have been clearly demonstrated in numerous models, leading to the suggestion that suboptimal conditions will impact the same system differently depending on the timing of the insult. A well-known example comes from work in several laboratories that provides insights regarding the role of the renin–angiotensin system (RAS) in development of the heart

[24–29]. Angiotensin II receptor type 1 (AT1) mRNA levels remained constant across the late gestation period, while receptor type 2 (AT2) mRNA expression was much higher with a dramatic decrease soon after birth [26]. In sheep, AT1 blockade during gestation was shown to have chamber and gestational age-specific effects on AT1 mRNA expression [increased at 95 and 135 days of gestational age (dGA) in the right atrium and the left ventricle at 110 dGA] [26]. Similarly, other investigators have reported that angiotensin II signaling has gestational age-specific effects on fetal heart growth in sheep and pig [25,27]. Sundgren *et al.* [25] have shown that angiotensin II promotes hyperplastic growth during early gestation, whereas Beinlich *et al.* [27] have demonstrated its importance in hypertrophic growth during neonatal life in the pig.

In a now classical study, Barraclough and Gorski [30] demonstrated that when female rats receive a single dose of androgen in the first 5 days of postnatal life, they did not show normal reproductive cycling at puberty. If exposure to androgen occurred 15 days later, near weaning, there was no persistent effect on reproduction. More recent animal studies have focused on poor nutrition in fetal life (for example, isocaloric low-protein diets [31], global nutrient restriction [32,33], and reduced uterine blood flow [34]), maternal stress (in rhesus monkeys [35,36] and pigs [37]) and exposure to various pharmacological agents (such as antenatal glucocorticoids [38–41]). Other studies compare outcomes resulting from insults during fetal life with insults occurring during early postnatal life [42,43].

The study by Barraclough and Gorski [30] also clearly demonstrated the second principle of developmental

programming, namely that there are permanent effects of neonatal androgen exposure on structures in the brain, particularly the hypothalamus, that regulate female reproductive cycle, following exposure at a critical time in development. More recently various well-characterized animal models have shown long-term consequences for the offspring. In rats, guinea-pigs, sheep and humans, fetal growth restriction and reduction in either maternal protein or global caloric intake leads to hypertension [44–50], obesity [10,51–56], diabetes [9,57–61], altered endocrine function [62–67] and mood disorders [68–71]. Finally, in comparing conclusions based on data from different species, it is clear that the trajectory of development of different systems varies between species. The simplest examples are differences between altricial species (born in an immature state and in which much development occurs after birth) and precocial species (born at a more advanced state of maturation). Thus the periods of vulnerability of the developing reproductive system which are postnatal in rodents are generally prenatal in sheep and humans.

Principle 4

Principle 4 states that programming may involve structural changes in important organs. We can say with confidence that a suboptimal developmental environment alters organ growth. The absolute numbers of cells in an organ may increase or decrease as a result of nutrient insufficiency and altered cell growth and division. Organ growth restriction has been demonstrated in models of maternal nutrient restriction [72,73], decrease in uterine blood supply as a result of uterine artery ligation [74] or uterine or umbilical artery embolization [75,76]. In each model the fetal nutrient supply is compromised.

Growth restriction may manifest itself only within certain components of an organ. We have shown that maternal nutrient restriction during the first half of gestation in the sheep induces asymmetric growth restriction [73]. Regardless of the cause, deficient levels of nutrients and oxygen lead to alterations in blood flow to, and vascular development in, the fetal brain, heart, lungs, abdominal viscera and skeletal muscle [77,78]. Maternal low-protein diets given to rats in pregnancy result in a smaller number of blood vessels per unit area in the fetal rat pancreas [79]. This decrease in potential vascular perfusion of the pancreas is likely to limit pancreatic islet function and be one of the major factors in the predisposition of offspring of nutrient-restricted mothers to develop diabetes.

Growth of the fetal kidney has received considerable attention because of its importance both during fetal life and in the transition to an independent postnatal existence. Several studies have investigated aspects of the

renal structure following nutritional deficit during pregnancy. In humans, fetal growth restriction is associated with a decrease in kidney size [80,81] and post-mortem glomerulus number [82]. Hughson *et al.* [83] showed a direct relationship between total glomeruli number in adult humans and birth weight. Furthermore, mean glomerulus volume was inversely correlated with glomerulus number. In rodents and sheep a reduction in total caloric food intake [49,84] or total dietary protein [85–87] during pregnancy induced a nephron deficit [88] and an increase in glomerular size [87]. Recently, a maternal diet low in protein has been shown to alter these relationships by changing cell turnover and gene expression at the beginning of metanephrogenesis in rats [89].

Widespread change in receptor populations is a further refinement of the idea that structure is impacted by a suboptimal environment *in utero*. Glucose transport proteins are found in fetal tissues from early in gestation [90–92] and are altered by maternal under nutrition [93]. Gene and protein expression within the fetal kidney, for example, has recently been found to be sensitive to maternal nutrient status in the rat [89,94,95], sheep [96,97] and the baboon [98–100]. Recent reports from our group have shown that maternal nutrient restriction during the first half of gestation in the sheep induces asymmetric growth restriction [73], alters gene transcription in the fetal heart [101–104], downregulates fetal skeletal-muscle protein synthesis [105], and alters concentrations of amino acids in fetal fluids by mid-gestation [106].

Principle 7

Principle 7 states that when postnatal conditions prove to be other than those for which the fetus prepared, problems may arise. The livers of fetal rats undernourished *in utero* show dramatically altered function and structure [107]. Individual lobules have more cells with phosphoenolpyruvate carboxykinase (PEPCK) activity than fetal livers of rats whose mothers received adequate nutrition. PEPCK is the key gluconeogenic enzyme and this nutritionally induced change indicates the fetal need to increase gluconeogenesis in the face of decreased glucose availability. Forhead and colleagues [108] have shown in the chronically catheterized fetal sheep that an increase in PEPCK is one of the many changes the fetus normally makes in late development to prepare to perform gluconeogenesis postnatally. If the neonate has an imbalance in liver glucose metabolism tending towards increased glucose production, that adaptation will be of value if food shortage is experienced postnatally. This response to adverse prenatal conditions has been called a predictive adaptive response (PAR), indicating its value in helping the offspring survive [109]. However, if the postnatal dietary regimen is adequate – or even

overabundant, as in our overconsumption-orientated society – the PAR is maladaptive and may predispose to obesity.

The idea that it is important to match prenatal development with the postnatal environment has important implications for the management of growth-restricted human neonates. Animal models have a major role to play in understanding effects of altered diet at different times of development with subsequent postnatal changes. Some outcomes of developmental programming can be certainly considered as PARs (such as an increased tendency to gluconeogenesis, which will be beneficial to survival under certain circumstances). However, other outcomes are distinctly maladaptive responses (MARs). Outcomes that could be considered MARs with no apparent value to survival are the decreased muscle mass or numbers of glomeruli that result from maternal nutrient deficiencies and predispose directly and inevitably to suboptimal health in later life.

Several rodent studies have investigated outcomes when neonates that are growth restricted at birth have been fed postnatally to induce catch-up growth [42,110]. As discussed above if the postnatal environment differs from the one in which the fetus developed, the mismatch may lead to later health problems. Ozanne and Hales [42] fed pregnant mice either a 20% protein or an 8% low-protein diet to restrict fetal growth and cross fostered pups at birth so that the offspring of mothers fed on a low-protein diet during pregnancy were reared by normally fed dams, the catch-up group. In the low-protein group offspring of mothers fed the normal protein diet during pregnancy were reared by mothers fed on a low-protein diet. Offspring in the catch-up group showed rapid catch-up growth and died earlier than controls. Interestingly, mice that grew normally before birth but were fed by mothers on the low-protein diet lived 57% longer than the catch-up group, approximately equivalent to a human living 75 rather than 50 years [42].

Principle 9

Principle 9 is that programming may cross generations. The consequences of various challenges to which the developing organism is exposed can be passed transgenerationally from female offspring, challenged during their own development to their own offspring [111]. In the 1990s two independent groups of investigators demonstrated that the F₁ female diabetic offspring of F₀ rats treated with streptozotocin during pregnancy themselves have F₂ offspring with altered glucose and carbohydrate metabolism [112–114].

More recently, Zambrano and colleagues [110] determined whether, when F₀ female rats are exposed to

protein restriction during pregnancy and/or lactation, their female pups (the F₁ daughters) deliver offspring (the F₂, or granddaughters and grandsons) with evidence *in vivo* of altered glucose and insulin metabolism. The F₀ rats were fed a normal control 20% casein diet (C) or a restricted diet (R) of 10% casein during pregnancy. After delivery, the mothers received either C or R diet during lactation to provide four sets of offspring – groups CC, RR, CR, and RC – where the first letter represents the diet during pregnancy and the second the diet during lactation. All the female offspring were fed *ad libitum* with C diet after weaning and during their first pregnancy and lactation. As they grew the female offspring of RR and CR mothers had low body weight and food intake with increased sensitivity to insulin during a glucose tolerance test at 110 days of postnatal life. Grandsons of the CR mothers showed evidence of insulin resistance. In contrast, granddaughters of the RC mothers showed evidence of insulin resistance.

Principle 10

Principle 10 is that programming may affect males and females differently. It is fairly well accepted that men are at greater risk for cardiovascular and renal disease than women of similar ages [115–117]. Recent work has clearly shown sex-specific expression of genes that code for proteins involved in drug metabolism and osmoregulation in the murine kidney that may directly contribute to sex effects in renal development or function [118]. Similarly, genes encoding drug and steroid metabolism were found differentially expressed between the sexes in the liver [118], an organ in which gender-specific transcriptional regulation of numerous genes by growth hormone has been described [119–122].

The transgenerational study by Zambrano *et al.* [110] demonstrates that maternal protein restriction adversely impacts glucose and insulin metabolism of male and female second-generation offspring in a manner specific to gender and the developmental time window. There are many other examples now in the literature of differing responses of male and female offspring to a variety of challenges. Recent work in rats has shown gender differences in the programming of the myogenic vasodilatory response to nitric oxide and prostaglandins in the mesenteric vasculature in male offspring following hypoxia during gestation [123]. In another study using radiotelemetric recording methods in conscious rats, raised blood pressure in female but not male offspring of dams fed a diet rich in animal lard has been demonstrated [124].

Our work in sheep has shown that maternal global caloric restriction alters intrarenal immunoreactive AT₁, AT₂ and renin expression in ways specific to gestational age

and gender [97]. We have also evaluated the effect of 30% global restriction from 30 to 90 days of gestation in the baboon (term 180 days) on the intrarenal RAS in male and female fetal kidneys at 90 days. Steady-state mRNA and protein were evaluated using the Human Genechip U133A 2.0 or Western blot in six controls fed *ad libitum* (C; three males, three females) and six nutrient-restricted (NR; three males, three females) fetuses. AT1 mRNA was increased (92%) in NR males (NRM) compared with C males (CM), C females (CF), and NR females (NRF). Both diet and the interaction between diet and gender were significant. There was no diet or gender effect on AT2 mRNA expression. Renal AT1 protein expression exhibited both diet and gender effects, with immunoreactivity being increased in NRM compared with all other groups. Conversely, AT2 protein was decreased in NRM compared with CM, CF and NRF, resulting in an AT1/AT2 protein ratio that was increased in NRM compared with CM, CF and NRF. A diet-gender interaction was observed on angiotensin-converting enzyme (ACE) immunoreactivity. Finally, no group differences were found in maternal plasma cortisol concentrations at 90 days. We found that diet-induced alterations in mRNA and protein expression, and the AT1/AT2 protein ratio, occur in male fetuses rather than females and are independent of maternal cortisol levels, leading to the postulate that gender-based sensitivity to nutrient deficit likely reflects differences between trajectories of growth, development and caloric demand in male and female fetuses.

Conclusion

The mounting human epidemiological and controlled animal experimental data clearly demonstrate that adult health is determined by both an individual's genome and the environment in which that genome develops. Whereas there are genetic conditions such as Huntington's chorea and cystic fibrosis that have a dominant and inevitable effect on adult health, the phenotype that emerges as a result of the interaction of epigenetic influences on the genome is critical to good adult health in a much larger segment of the population. Thus understanding the prenatal (and immediate postnatal) influences on adult disease and developing preventive strategies for pregnancies at risk is one of the major challenges of biomedical research and the delivery of obstetric care.

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Additional references related to this topic can also be found in the Current World Literature section in this issue (pp. 196–197).

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