

Embryonic and Fetal Programming of Physiological Disorders in Adulthood

Christopher Lau* and John M. Rogers

In the past decade, data from numerous epidemiological studies have indicated strong inverse associations between birth weight and risk of coronary heart disease, hypertension, type 2-diabetes, and other diseases in adulthood. The "Barker hypothesis" thus postulates that a number of organ structures and functions undergo programming during embryonic and fetal life. This developmental programming determines the set points of physiological and metabolic responses in adult life. Alterations of nutrient availability during gestation may lead to developmental adaptations, via hormonal maneuvers by the embryo and fetus that readjust these set points. These adaptive measures have short-term benefits to the embryo and fetus, so that the newborn will be better prepared for the adverse environment (e.g., undernutrition). However, adequate nutritional support during postnatal life that enables catch-up growth may create metabolic conflicts that predispose the adult to aberrant physiological functions and, ultimately, increased risk of disease. It is plausible that other adverse in utero conditions, including exposure to developmental toxicants, may similarly alter adult disease susceptibility. This article provides an overview of the Barker hypothesis, its supporting evidence, the current advances in understanding the biological mechanisms underlying this phenomenon, and its implications for developmental toxicology. **Birth Defects Research (Part C) 72: 300–312, 2004. Published 2005 Wiley-Liss, Inc.†**

INTRODUCTION

Teratology is classically defined as the study of abnormal development during embryonic and fetal stages that leads to anatomical malformations, while more recently this has been extended to encompass functional deficits in the immature organism. Indeed, since the seminal works by Warkany and his colleagues in the 1940s (Warkany and Nelson, 1940; Warkany, 1943; Warkany and Schrafenberger, 1947), experimental teratology combined with clinical birth defect research

has evolved into a crosscutting discipline involving developmental biologists, geneticists, toxicologists, pediatricians, and epidemiologists. A majority of the extant teratology literature focuses on structural defects induced by an altered intrauterine environment during gestation. While overt structural defects are usually associated with functional deficiency, the reverse is not necessarily true. Biochemical changes induced by prenatal insults that lead to physiological deficits of organ function may not always be ac-

companied by *detectable* anatomical abnormalities. Hence, in recent decades, considerable attention has been drawn to *functional teratology*, an extension of the investigation beyond morphological examinations to include the evaluations of functional integrity of organ systems (for examples, see Kavlock and Grabowski, 1983; Lau and Kavlock, 1994; Herzyk et al., 2002). Nonetheless, a majority of these

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works has focused primarily on the perinatal and early postnatal periods, and long-term follow-up studies are seldom available.

In the early 1990s, a novel hypothesis was advanced by Barker and associates to link nutritional insults during embryonic and fetal development not only to impaired maturation of physiological functions, but also to latent diseases in adulthood (Barker, 1992a; Hales and Barker, 1992). The "Barker hypothesis" postulates that a number of organ structures and associated functions undergo programming during embryonic and fetal life, which determines the set point of physiological and metabolic responses that carry into adulthood. Hence, alterations in embryonic and fetal nutrition, as well as endocrine status during gestation, can result in developmental adaptations that produce permanent structural, physiological, and metabolic changes, thereby predisposing an individual to cardiovascular, metabolic, and endocrine diseases in adult life (Barker, 1992b). This "fetal origins" hypothesis has increasingly gained support in the past decade from both epidemiological studies as well as investigations in animal models. This article will provide a review of the Barker hypothesis, the current advances in understanding, and implications for future teratological evaluations.

MATERNAL NUTRITIONAL STATUS AND ADULT CARDIOVASCULAR DISEASES

In a series of retrospective studies, Barker and coworkers examined the incidence of coronary heart disease in middle and late-life men and women, and correlated that to the body measurements at birth. In one study, the record of 16,000 subjects born in Hertfordshire, England between 1911 and 1930 was traced. Using birth weight as a surrogate indicator of intrauterine growth, Osmond et al. (1993) reported a fall in death rates from coronary heart disease by almost a factor of two, between those who were born at the upper (4.3 kg) and

lower (2.5 kg) extremes of weight. Variations of birth weight seen at Hertfordshire did not appear to be associated with differences in social class. Similar results were obtained by Barker et al. (1993a) from another study conducted in Sheffield, England, with men born between 1907 and 1925. It was noted that the subjects who were small at birth (but born at full term), rather than those born prematurely, were at increased risk of coronary heart disease.

Since then, a robust body of epidemiological evidence supports the British findings. In the San Antonio Heart Study, Valdez et al. (1994) examined the prevalence of ischemic heart disease in young adults and compared that to the participants' birth weight. These investigators reported the odds of expressing the disease increased 1.72 times for each tertile decrease in birth weight. These findings were independent of sex, ethnicity, current socioeconomic status, or obesity. In the Caerphilly study, Frankel et al. (1996) followed a cohort of men in South Wales for 10 years, and reported a significant inverse relationship between birth weight and incidence of coronary heart disease. This association was not changed by adjustment for age, social class, marital status, and smoking history. Stein et al. (1996) studied a group of men and women who were born between 1934 and 1954 in and still residing in Mysore, South India, and noted that low birth weight, short birth length, and small head circumference at birth were associated with a raised prevalence of coronary heart disease. The prevalence fell from 11% in people whose birth weights were 2.5 kg or less, to 3% in those whose birth weights were more than 3.1 kg. In the Nurses' Health Study in the United States, Rich-Edwards et al. (1997) followed a large cohort (over 120,000 women) for 16 years, and indicated a moderate but consistent inverse relationship between birth weight and risk of nonfatal cardiovascular disease. Women who had birth weights of 2.5 kg or less had a significantly higher risk (by 23%) of nonfatal

the combination of small size at birth and during infancy, followed by accelerated weight gain from the ages of three to 11 years, predicted significantly higher cumulative incidence of coronary heart disease, type-2 diabetes, and hypertension.

cardiovascular disease compared with the rest of the cohort. Consistent with other studies, the association persisted despite adjustment for hypertension, diabetes, raised cholesterol level, height, body mass index, and childhood socioeconomic status. More recently, Barker et al. (2002) carried out a longitudinal study of Finnish men and women born in Helsinki between 1924 and 1944, and reported that the combination of small size at birth and during infancy, followed by accelerated weight gain from the ages of three to 11 years, predicted significantly higher cumulative incidence of coronary heart disease, type-2 diabetes, and hypertension.

Besides coronary heart disease, blood pressure in adults has also been found to be inversely related to birth weight. As early as 1988, Gennser et al. (1988) evaluated the hospital records of a small pool of Swedish army conscripts and discovered that the risk of increased diastolic blood pressure in early adult life was significantly higher among those who had been smaller at birth than those whose birth weight had been appropriate for gestational age (odds ratio 3.63). These results were confirmed by another Swedish study with a

larger cohort (Leon et al., 1996), where men who had lower birth weight (<3.25 kg) and were above median adult height were found to have the highest blood pressure. Law et al. (1993) examined the British records of men and women of various ages, and observed that those who had lower birth weight had higher systolic blood pressure. This relationship became stronger with increasing age, such that at ages 64–71 years, systolic pressure decreased by 5.2 mm of Hg for every kilogram increase in birth weight. Curhan et al. (1996) examined the records of over 160,000 women in the Nurses' Health Studies in the United States, and reported an age-adjusted odds ratio for hypertension in the low birth weight category (<2.3 kg) of 1.39–1.43 compared to the average birth weight category (3.9 kg). Correspondingly, a decrease of systolic blood pressure by 0.43 mm of Hg, and of diastolic blood pressure by 0.21 mm of Hg, was noted for each 0.45 kg increase in birth weight. The results of this study were consistent with the British data, although the magnitude of the effect was much smaller. In fact, Huxley et al. (2000) reviewed 80 studies from population cohorts around the world describing the relationship of blood pressure with birth weight, and noted a fall in blood pressure with increased birth weight, and meta-analysis demonstrated an average drop of 2 mm of Hg/kg. Interestingly, postnatal catch-up growth was positively associated with blood pressure, with the highest blood pressures occurring in individuals having low birth weight but high rates of postnatal growth. Results from more recent studies are consistent with these general observations (Eriksson et al., 2000; Adair and Cole, 2003; Singh and Hoy, 2003; Hardy et al., 2004; Fagerberg et al., 2004). In addition, IJzerman et al. (2003) suggested that the increase in blood pressure related to low birth weight might involve an increase in cardiac sympathetic activity.

In order to minimize the potential influence of genetic confounders in the association of fetal weight defi-

cits and adult hypertension, studies were conducted in twin pairs. Poulter et al. (1999) reported a graded inverse relation between differences in birth weight and adult blood pressure in a British cohort of twins. IJzerman et al. (2000) examined the birth records of a Dutch cohort of monozygotic and dizygotic twins, and determined their blood pressure at adolescent age. These investigators reported that intrapair differences in birth weight were negatively and significantly associated with differences in systolic blood pressure, with a regression coefficient of -5.7 mm of Hg/kg of birth weight in the dizygotic but not monozygotic twins, suggesting that genetic factors might play a role in the association between birth weight and blood pressure. In comparison, Christensen et al. (2001a) measured the blood pressure of adolescent twins in the Minnesota Twin Family Study and found a negative association between birth weight and systolic blood pressure in the overall samples, with a regression coefficient of -1.88 mm of Hg/kg. However, in contrast to the IJzerman et al. (2000) study, this effect was more pronounced in the monozygotic twins, with a regression coefficient of -2.44 mm Hg. Indeed, Phenekos (2001) examined prospective twin studies in the United States and Northern Europe, and failed to find evidence to support genetic factors that might predispose low birth weight and adult diseases. Loos et al. (2001) evaluated a women's cohort from the East Flanders Prospective Twin Survey and noted a similar decrease of systolic pressure (by 4.27 mm of Hg) and diastolic pressure (by 2.18 mm of Hg) per kilogram increase in birth weight. It is noteworthy that the values of regression coefficient revealed in these twin studies are quite similar to those reported from singleton adolescents (Huxley et al., 2000).

Epidemiological and clinical studies, therefore, suggest a link between nutritional deprivation in utero, reflected by low birth weight, and latent cardiovascular diseases in adulthood. However, among

these studies, the actual maternal and fetal nutritional status was always inferred. Thus, episodes of periodic famine, where the maternal nutritional intakes were better documented, offer a unique opportunity to test the Barker hypothesis. Generally speaking, results from these studies are supportive of the hypothesis in a broad sense, but far from definitive. For instance, study of the health consequences of adults born during the Dutch famine of 1944–1945, when the official daily rations dropped from about 1,800 kcal to 400–800 kcal, revealed that only those exposed to malnutrition during early (but not middle or late) gestation were found to have a higher prevalence of coronary heart disease when compared to better nourished subjects (those born the year before or conceived in the year after the famine), and this effect was independent of birth weight (Roseboom et al., 2000a). It is of interest that substantial effects of the famine on fetal growth (indicated by birth weight, length, and head circumference) were seen only with third trimester exposure (Stein et al., 2004), which may explain the lack of significant birth weight association with the early gestational effects of famine. Alternatively, Roseboom et al. (2000a) contended that because the famine ended abruptly, the women who were malnourished during early pregnancy received adequate nourishment during the later part of pregnancy, thus allowing the fetus to catch up on their growth. It is this subpopulation of adults who had the highest prevalence of coronary heart disease. These investigators surmised that the transition from nutritional deprivation in early gestation to nutritional adequacy later on led to metabolic conflicts that resulted in an increased risk of coronary heart disease. On the other hand, no effect of prenatal exposure to undernutrition on blood pressure was observed in the adult subjects (Roseboom et al., 1999), although a more atherogenic lipid profile was seen in those exposed to famine (Roseboom et al., 2000b).

The Leningrad famine in 1941–1944, where the average daily ration for most of the citizens provided around 300 cal and contained virtually no protein, afforded another case study. Surprisingly, comparison of adult subjects exposed to malnutrition in utero during the siege of Leningrad, to those who were born before the siege and those born outside of the affected area (unexposed groups), revealed no relationship between intrauterine growth and adult cardiovascular disease, blood pressure, or plasma lipid concentrations (Stanner et al., 1997). One plausible explanation for the different findings between these two studies may be the length of the famines (much longer in Leningrad) and the abrupt recovery of nutritional status with the Dutch famine (compared to the long periods of relative food shortage preceding and following the Leningrad siege). Hence, the fetuses and infants in the Netherlands were allowed catch-up growth that could account for the latent coronary heart disease, whereas those in Russia had to adapt to a sustained period of nutritional deprivation and, consequently, avoided any metabolic conflicts in adult life (Stanner and Yudkin, 2001).

INTRAUTERINE GROWTH DEFICIT AND ITS ASSOCIATION WITH TYPE 2 DIABETES AND METABOLIC SYNDROME: THE THRIFTY PHENOTYPE HYPOTHESIS

The association of poverty and malnutrition with infant mortality and disease development is not a novel concept (Forsdahl, 1977; Barker and Osmond, 1986), but it was not until the Hertfordshire study, in which Hales et al. (1991) evaluated the glucose tolerance of 64-year-old men and traced their birth records, that strong correlations between low birth weight and metabolic diseases were derived. These investigators found that those men born weighing less than 2.5 kg were almost seven times more likely to

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have either impaired glucose tolerance or diabetes than those born weighing more than 4.3 kg. Because a link between hypertension and abnormalities of carbohydrate and lipoprotein metabolism had been suggested previously (Reaven and Hoffman, 1989), Barker et al. (1993b) evaluated the same subjects in the Hertfordshire study and found that the prevalence of the metabolic syndrome fell progressively from those with the lowest birth weights to those with the highest. Indeed, when the current body mass index was adjusted for, the odds ratio for the metabolic syndrome in subjects weighing less than 2.5 kg at birth was 18 in comparison with those weighing more than 4.3 kg. Similar results were obtained in another study conducted with men and women in Preston, England that yielded an odds ratio of 14 (Barker et al., 1993b). By and large, these findings were confirmed by other studies with different populations. Two Swedish studies with elderly men reported an inverse relationship between the prevalence of glucose intolerance and birth weight (Lithell et al., 1996; McKeigue et al., 1998). Rich-Edwards et al. (1999) examined nearly 70,000 women in the Nurses' Health Study in the United States, and noted that birth weight was inversely associated with risk for type 2 diabetes during

adulthood; this association was not influenced by ethnicity, childhood socioeconomic status, or adult life style factors. Carlsson et al. (1999) conducted a population-based cross-sectional study of Swedish middle-aged men, and reported that birth weights of less than 3 kg were associated with an odds ratio of 2.3 for diabetes.

Similar observations were made with the Pima Indian children and young adults (Dabellea et al., 1999; Pettitt and Jovanovic, 2001; Stefan et al., 2004) and among school children in Taiwan (Wei et al., 2003). Interestingly, a U-shape relationship between birth weight and risk of type 2 diabetes was noted in these two case studies. Whether the low and high birth weight subjects may share a common etiology for development of diabetes remains to be clarified. Furthermore, to determine if the link between birth weight and diabetes might involve genetic factors, Iliadou et al. (2004) examined a large cohort of Swedish twins, and reported a significant increase in risk of developing type 2 diabetes for a 1-kg decrease in their mean birth weight (odds ratio 2.13), lending support to malnutrition being a possible factor underlying the in utero programming effect that led to a subsequent increased health risk.

Based on their findings, investigators from the Hertfordshire studies proposed a "thrifty phenotype" hypothesis that suggested a malnourished fetus made adaptive changes in glucose-insulin metabolism (including reduced capacity for insulin secretion and insulin resistance), which were designed to improve survival under conditions of nutritional deprivation (Hales and Barker, 1992). The deficient insulin secretion would not be detrimental to health, had the individuals continued to be malnourished (and remained thin); however, when the nutritional status became adequate or excessive during postnatal periods, physiological conflicts arose as glucose intolerance was triggered by a positive caloric balance. Several studies have confirmed an additive effect of fetal growth restric-

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tion and subsequent obesity in worsening glucose tolerance and insulin sensitivity (reviewed in Petry and Hales, 1999; Hales and Barker, 2001). Indeed, in the thrifty phenotype hypothesis, "catch-up" growth that leads to childhood obesity has been regarded as a key component in the manifestation of latent diseases in adulthood (Hales and Ozanne, 2003); hence, no increase in risk of diabetes for small size at birth was observed without excessive postnatal weight gain (Hypponen et al., 2003). Similarly, Cianfarani et al. (1999) has advanced a "catch-up growth" hypothesis with intrauterine growth retarded (IUGR) children. These researchers postulated a rearrangement of the endocrine system in the third trimester leading to low levels of insulin, insulin-like growth factor (IGF)-1, insulin-like growth factor binding protein (IGFBP)-3, and high

levels of growth hormone, IGFBP-1, and IGFBP-2. Shortly after birth, tissues that were chronically depleted of insulin and IGF-1 during fetal life and suddenly exposed to increased levels of these two hormones might counteract their additive insulin-like actions by developing insulin resistance as a metabolic defense mechanism to protect the organism from hypoglycemia. Thus, those IUGR children who showed early and complete recovery from intrauterine growth retardation would be at higher risk for metabolic disorders in adulthood. Recent models, based on multiple-regression analysis of the health outcomes on various birth weights and growth rates, also support the fetal programming effect and the significance of catch-up growth (Cole, 2004; Kuzawa, 2004). Moreover, using a mouse model, Ozanne and Hales (2004) demonstrated that the lifespan of the animal was considerably shortened if the postnatal period of growth was accelerated to make up for the growth deficits in utero; additionally, those mice that were exposed to an obesity-inducing diet after weaning were particularly susceptible to the adverse effects on longevity.

LOW BIRTH WEIGHT AND FUNCTIONAL DEFICIENCY OF OTHER ORGAN SYSTEMS IN ADULTHOOD

In addition to the cardiovascular diseases and diabetes described above, the Barker hypothesis has prompted other investigators to examine if functional deficiency of various organ systems can be similarly linked to impaired fetal growth. Most of these studies have been conducted fairly recently, and the findings are therefore less definitive and uniform. Hinchcliffe et al. (1992) examined the kidneys of stillborn IUGR infants and noted 35% fewer nephrons in this group, compared to the average infants at comparable gestational age. Mackenzie et al. (1996) and Manalich et al. (2000) reported strong correlations between glomerular number (direct) and size (inverse) with low birth weight in human neonates,

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and it was reasoned that the shortfall in nephron endowment could be the renal risk factor for hypertension in those of low birth weight. However, Marchand and Langley-Evans (2001) recently raised the possibility that reduced renal reserve and hypertension might merely be coincident, and not causally associated. Hoo et al. (2004) reported that forced expiratory volume of the lung was significantly reduced in the low birth weight infants in the first year of life. In the Hertfordshire study, Barker et al. (1991) noted that the mean forced expiratory volume and forced vital capacity, measurements of function in elderly men, rose with increase in birth weight, independently of smoking habit and social class. Evaluation of another cohort in South India provided very similar results, suggesting that adult lung function is programmed in fetal life (Stein et al., 1997). In addition, Lopushaa et al. (2000) detected a link between exposure to the Dutch famine in mid- and early gestation and prevalence of obstructive airways disease in adulthood.

McDade et al. (2001) followed the development of thymic function in a cohort from the Philippines and reported that prenatal undernutrition was significantly associated with reduced thymopoietin production, suggesting that the environments during fetal and early infant life

may have long-term implications for immunocompetence and adult disease risk. In contrast, Moore et al. (2001) reported that immune function in rural Gambian children was not related to birth size or prenatal high-energy supplementation, although the investigators pointed out the possibility that the defect related to low birth weight might be in immunologic memory, rather than in the early immune response that was measured in this study. Clark et al. (1996) investigated whether functional alterations of the hypothalamic-pituitary-adrenal (HPA) axis might be associated with cardiovascular diseases and diabetes in the low birth weight adults. They measured the urinary metabolites of glucocorticoid hormones and found an inverse association of metabolite excretion with birth weight. Phillips et al. (2000) provided supporting evidence for a relationship between low birth weight and elevated fasting plasma cortisol concentration (an indicator of HPA activity) in adults; furthermore, men with low birth weight were found to have significantly lower pituitary-adrenal responses to the dexamethasone/corticotrophin releasing hormone test (Ward et al., 2004). However, when Fall et al. (2002) examined the serum cortisol profiles of healthy elderly subjects, they did not find any association between low birth weight and continuously raised cortisol concentration in old age. These investigators deduced that the programming effects on the HPA axis might be on its reactivity, rather than the basal tone of hormone secretion.

The association between low birth weight and reproductive function is more subtle. Cresswell et al. (1997a) followed the elderly women in the Hertfordshire study and noted that the age at menopause was unrelated to birth weight, but it occurred earlier in women who had low weight at one year of age. Adair (2001) examined a cohort of young girls in the Philippines and found that although birth weight alone was not significantly related to age at menarche, girls who were relatively long and thin at

birth attained menarche an average of six months earlier than those who were short and thin; this effect was most pronounced among girls with greater than average growth increments in the first six months of life. Supporting evidence may be found in an animal study in which IUGR or food restriction during the preweaning period led to delayed onset of puberty in rats; these effects were not dependent on the achievement of a certain crucial weight of the pups at puberty (Engelbregt et al., 2000). On the other hand, Cresswell et al. (1997b) reported that women with polycystic ovaries were found to have above average birth weight and were born to overweight mothers. McCormack et al. (2003) followed a Swedish cohort of women (<50 years old), and found that size at birth was positively associated with rates of breast cancer in premenopausal women. These findings are in general agreement with another study examining a Danish cohort (Ahlgren et al., 2004), and suggest a link between high birth weight and cancer incidence. Together with the U-shape findings of diabetes (association with low and high birth weights) in the Pima Indian and Taiwanese children, the issue of macrosomia and long-term adverse health effects needs to be scrutinized more closely. Adult atherosclerosis (Palinski and Napoli, 2002) and high risk of the development of pre-eclampsia (Dempsey et al., 2003) have also been tied to low birth weight. Lastly, Thompson et al. (2001) examined an elderly cohort to determine if there was an association between birth weight and the risk of depressive disorders, and noted that the odds ratios for depression among men, but not women, rose incrementally with decreasing birth weight. These investigators suggested a neurodevelopmental etiology of depression and that fetal undernutrition might predispose men to depression through programming of the HPA axis. Additional research will be required to explore this possibility.

INTERGENERATIONAL EFFECTS OF FETAL PROGRAMMING

While the aforementioned epidemiological findings have revealed the fetal origins of latent diseases in adulthood, several provocative studies have extended this programming concept to span the next generation (reviewed in Reusens and Remacle, 2001; Drake and Walker, 2004). The notion of trans-generational effects of pregnancy outcomes is actually not new, and has been discussed previously in an excellent review (Emanuel, 1993). In toxicological studies, residual adverse effects derived from in utero exposure to chemicals (Csaba and Inczeffi-Gonda, 1999; Stoll et al., 2003; Newbold, 2004) and ionizing radiation (Dubrova, 2003) have been reported to be passed on to the next generation. To determine whether this phenomenon is extended to fetal programming, Skjærven et al. (1997) examined the birth records of Norwegian women, and reported that a mother's birth weight was strongly associated with the weight of her baby, and maternal birth weight was associated with perinatal survival of her baby for mothers whose birth weights were under 2 kg (odds ratio 2.3). In a follow-up study of men and women whose mother's or father's size at birth was recorded, Barker et al. (2000) reported that the children's birth weight was strongly related to their mother's birth weight, and blood pressure in the adults was lower with increasing mother's birth weight, whereas the father's birth size was not related to the adult offspring's blood pressure. Studying three generations of the British national birth cohort, Hypponen et al. (2004) indicated that the mother's birth weight was the strongest determinant of the offspring's birth weight. Interestingly, this intergenerational birth weight association was not observed for mothers born either very small or very large. Similar to the previous study, the father's birth weight or body mass index did not affect the offspring's birth weight. In comparison, Veena

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et al. (2004) reported that the birth weights of both parents could predict the offspring's birth weight equally well in a South Indian cohort, although the mother's body mass index had a greater effect than the father's on the offspring's birth weight. These observations are consistent with a study employing dexamethasone-programmed rats, a model in which fetal exposure to excess glucocorticoid resulted in low birth weight with subsequent adult hyperinsulinemia and hyperglycemia (Drake et al., 2004). These researchers reported that the male offspring of female rats that had been exposed to dexamethasone prenatally, also had reduced birth weight, glucose intolerance, and elevated hepatic gluconeogenic enzyme activity; these effects were resolved in a third generation. In contrast, Rogers et al. (2003) failed to detect any evidence for intergenerational reproductive effects due to prenatal and/or postnatal undernutrition in a mouse model. Despite catch-up growth in the IUGR pups (such that at weaning, their weights were no longer different from that of controls), no significant differences

were noted in the average F2 litter size or pup weight at birth. An account for these disparate findings remains to be established.

DISSENTING VIEWS

Despite the mounting evidence to support the Barker hypothesis, this provocative concept is not without its critics (Paneth and Susser, 1995; Huxley et al. 2002; Huxley and Neil, 2004) and discordant findings. Most of these were derived from twin studies, in which no significant correlations were noted between birth measurements and acute myocardial infarction (Hubinette et al., 2001), cardiovascular mortality (Christensen et al., 2001b), or blood pressure (Baird et al., 2001; Nowson et al., 2001; Zhang et al., 2001; Johansson-Kark et al., 2002). Barker (1995) argued that twins were a mixture of proportionately and disproportionately small babies (Leveno et al., 1979), and the prevalence of coronary heart disease might depend on the proportionality of these babies, rather than their size alone. However, de Geus et al. (2001) compared blood pressure of twins with their singleton siblings in the Netherlands, and did not observe any difference in blood pressure between twins and singletons, despite the lower birth weight in the twins. Williams and Poulton (2002) followed the birth size, growth, and blood pressure of a cohort in New Zealand, and found no evidence to suggest that children with a low birth weight who became overweight or obese had extra-high blood pressure. In a prospective longitudinal study, Falkner et al. (2004) also failed to detect significant correlation coefficients of birth weight with all blood pressure measurements.

Another issue of contention is the use of birth weight as a surrogate for intrauterine nutritional deficits that presumably led to fetal growth retardation, a key component linking to the latent diseases in adulthood in the Barker hypothesis. However, with the possible exception of the famine studies where food intake of the populations was

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recorded, the actual nutritional status of the expecting mothers in the epidemiological studies described above was typically unknown. Symonds et al. (2000), Wilcox (2001), and Morley et al. (2002), in particular, questioned whether birth weight alone was a reliable marker for gestational exposure that increased the risk of adult diseases. These investigators pointed out that low birth weight babies could be produced at high altitudes, but that did not lead to an increase in infant mortality. Babies born of Mexican-American mothers in the United States tended to have lower birth weight compared to non-Hispanic whites, but this subpopulation actually had lower infant mortality rates. Whether the adults born at high altitudes or to Mexican-American heritage are more prone to cardiovascular and metabolic diseases than the average populations, as predicted by the Barker hypothesis, remains to be investigated. Alternatively, other measures of birth size or proportionality may provide better markers of an adverse fetal environment and be more accurately correlated to adult health outcomes. For instance, fetal-placental weight ratio at birth has been suggested as a good predictor of adult diseases (Barker et al., 1990; Godfrey and Robinson, 1998).

In addition to the "thrifty phenotype" concept that relates adult diseases to fetal programming in re-

sponse to adverse intrauterine environment, a role for genetic susceptibility has been suggested by the association of low birth weight with type 2 diabetes and cardiovascular diseases (Hattersley and Tooke; 1999; Lindsay et al., 2000; Stern et al., 2000; Frayling and Hattersley, 2001). These investigators proposed that fetal polygenic genetic factors, involving defects such as mutation, duplication, etc., might lead to increased insulin resistance both in utero and in adult life, and this could produce two phenotypes—a small, thin baby and an adult with insulin resistance and increased risk of hypertension and atherosclerosis, particularly in the presence of obesity. To support this contention, it is generally presumed that birth weight and size would be influenced by both sets of inherited paternal and maternal genetic factors. Specifically, if genes shared by the father and the fetus are functional, then insulin resistance of the father should show an inverse correlation with his child's birth weight, measures of insulin-mediated growth and endothelial function. Polymorphisms of genes associated with insulin resistance should also be correlated to impaired fetal growth. This correlation should be evident both within populations and within families, whereby newborns inheriting an insulin-resistance-associated mutation would be expected to have lower birth weight than their siblings who did not inherit the mutation. Nonetheless, the validity of this alternative explanation for the association of low birth weight and metabolic syndrome is yet to be determined.

SUPPORTING FINDINGS FROM ANIMAL MODELS

Despite skepticism and alternative theories, the Barker hypothesis has slowly gained acceptance over the years (Gillman and Rich-Edwards, 2000; Jaquet et al., 2003). However, research on this novel concept largely based on epidemiological observations is not without its challenges (Lucas, 1998; Gillman, 2002). Among them, the veracity of

Petry et al. (1997) reported that rats exposed to protein restriction during gestation and early postnatal life, and subsequently fed a highly palatable diet to induce obesity, were glucose intolerant, hypertriacylglycerolemic, and hypertensive; these effects were additive,

maternal and fetal nutritional status, and the underlying biological mechanisms linking fetal programming to alterations of physiological functions and responses in adulthood are most prominent. Thus, laboratory animal models that are more amenable to experimental manipulations offer an avenue to further investigation of the Barker hypothesis. Indeed, various models involving dietary, surgical, genetic, and pharmacological interventions have recently been reviewed (Bertram and Hanson, 2001; Ozanne, 2001; Langley-Evans, 2001; Holemans et al., 2003).

Results from these experimental models, by and large, substantiated the epidemiological findings, and are described briefly here. Using a rat model, Langley and Jackson (1994) and Langley et al. (1994) reported increased systolic blood pressure and altered glucose tolerance in adults that were exposed to maternal low protein diets during fetal development. Petry et al. (1997) reported that rats exposed to protein restriction during gestation and early postnatal life, and subsequently fed a highly palatable diet to induce obesity were glucose intolerant, hyper-

triacylglycerolemic, and hypertensive; these effects were additive, suggesting that early protein restriction and later obesity were independent risk factors for the development of hypertension. Ozanne et al. (1998) noted ketosis resistance in rat offspring of protein-malnourished dams, a metabolic condition that has been similarly observed in some forms of human diabetes. Kwong et al. (2000) observed that maternal low protein diet in the rat, during only the preimplantation period (for the first 4.25 days after mating), was sufficient to produce blastocyst abnormalities, low birth weight, altered postnatal growth, and hypertension during adulthood. Lewis et al. (2001, 2002) examined the effects of maternal iron deficiency in the rat, and found that pup weight was reduced at birth and systolic pressure of the offspring was significantly elevated. Negative correlation of birth weight and systolic pressure in the adult offspring was obtained from guinea pigs that were exposed to a mild regimen of maternal undernutrition (85% of ad libitum food intake from four weeks before, and throughout pregnancy) (Kind et al., 2002). Similarly, using a sheep model, Gopalakrishnan et al. (2004) reported an elevation of blood pressure in three-year-old animals that were subjected to maternal undernutrition (50% of ad libitum nutrient intake) throughout pregnancy. Maternal protein restriction was reported to lead to hyperinsulinemia and reduced insulin signaling protein expression in adult rat offspring (Fernandez-Twinn et al., 2004). Prenatal stress was found to induce intrauterine growth restriction that led to glucose intolerance and altered feeding behaviors in the adult rat (Lesage et al., 2004). Without employing any experimental manipulation, Woods and Weeks (2004) selected rat pups whose birth weight was less than 90% of the mean pup weight for the litter, and noted that the blood pressure of the runts was significantly higher than their normal birth weight littermates, and that this relative hypertensive effect was not likely due to gross differences in renal function.

The possible biological mecha-

nisms underlying intrauterine programming of adult diseases have been discussed in recent reviews (Symonds et al., 2001; Bertram and Hanson, 2002; Fowden and Forhead, 2004). A majority of these studies focused on resetting of fetal endocrine homeostasis in response to fetal growth impairment brought forth by maternal nutritional deficits. Several hormones are known to regulate fetal growth and development and may also play a central role in intrauterine programming. These include anabolic hormones such as insulin, insulin-like growth factors (IGF-I and IGF-II), prolactin and thyroid hormones, as well as catabolic hormones such as the glucocorticoids. These hormones act as nutritional and maturational cues, and adapt fetal development to the prevailing intrauterine conditions, thereby maximizing the chances of survival in utero and at birth. However, alterations of these signals in response to an adverse intrauterine environment (such as a deficit of nutrients) will accordingly adjust the set point of these hormones in modulating their functional reactivity in the tissues, as well as regulatory controls of the organ systems. Such endocrine maneuvers may have short-term benefits to the well being of the fetus, but also permanently reset the endocrine systems so as to predispose the adult to aberrant physiological functions and ultimately, disease pathogenesis. Indeed, Plagemann (2004) described these hormones as "endogenous functional teratogens." Alterations of the glucocorticoid hormones and the HPA axis, in particular, have drawn considerable attention, and evidence of their involvement in fetal programming has been discussed in detail in two excellent reviews (Bertram and Hanson, 2002; Fowden and Forhead, 2004). Briefly, their concentrations in utero can be elevated by nutritional perturbation and other stressful insults that are known to have programming effects. These changes of hormone levels are paralleled by altered expression of glucocorticoid receptors (that confer hormonal actions) and the associated genes of

receptors, enzymes, ion channels, and transporters that are regulated by the glucocorticoids, as well as ontogenetic deviations of the HPA axis. Hence, these endocrine changes can be both the cause and the consequence of intrauterine programming.

Alternatively, an intriguing epigenetic hypothesis of regulating the intrauterine supply of and demand for nutrients by imprinted genes has been advanced to account for fetal programming (reviewed in Rakyan et al., 2001; Reik et al., 2003; Waterland and Jirtle 2004). Epigenetic control of gene expression involves modification of the genome not involving alterations of the DNA sequence, and is typically mediated by changing the DNA methylation pattern and/or modifications of chromatin packaging via changes in histone acetylation. These mechanisms may influence gene expression by transcriptional silencing of the modified allele (Rakyan et al., 2001) and, therefore, may affect the phenotype without changing the DNA sequence per se. These molecular events are established early in development and maintained throughout life, thus making them prime candidates to account for fetal programming. The seminal work by Waterland and Jirtle (2003), in particular, supports this hypothesis. In this study, the investigators were able to demonstrate that dietary supplementation can dramatically alter the heritable phenotypic changes in agouti mice. Furthermore, the transposon gene involved in this alteration was shown to be imprinted, such that only the maternal contribution was expressed. If the findings with the agouti mouse model are applicable to humans, they may provide a viable mechanism to explain fetal programming via metabolic imprinting that leads to chronic diseases in adult life (Waterland and Jirtle, 1999).

IMPLICATIONS TO DEVELOPMENTAL TOXICOLOGY

An increasing body of evidence from epidemiological observations and studies with various animal models has pointed toward an in-

verse correlation between low birth weight and latent diseases in adults. A majority, of these studies have presumed that intrauterine malnutrition is the cause of low birth weight. However, in addition to deprivation of nutrients, deficits in fetal growth can also be produced by chemical insults during pregnancy. In fact, the Guidelines for Developmental Toxicity Risk Assessment issued by the U.S. Environmental Protection Agency (1991) state that:

"A change in offspring body weight is a sensitive indicator of developmental toxicity, in part because it is a continuous variable. In some cases, offspring weight reduction may be the only indicator of developmental toxicity. While there is always a question as to whether weight reduction is a permanent or transitory effect, little is known about the long-term consequences of short-term fetal or neonatal weight changes. Therefore, when significant weight reduction effects are noted, they are used as a basis to establish the NOAEL (no-observed-adverse-effect-level)."

Indeed, several studies have attempted to link toxicant-induced reductions of fetal weight at term to increased incidence of developmental malformations and prenatal death (Ryan et al., 1991; Catalano et al., 1993; Gaylor and Chen, 1993). The question thus arises as to whether the fetal and birth weight deficits induced by developmental toxicants may similarly lead to reprogramming of set points for physiological functions and responses that ultimately lead to cardiovascular and metabolic diseases in adulthood. Naturally, maternal malnutrition itself can be associated with the chemical insult, typically indicated by a transient reduction of maternal food intake. Hence, it is not unreasonable to apply the Barker hypothesis to toxicological investigations. Furthermore, in most toxicological studies, fetal and neonatal weight deficits resulting from chemical exposure in

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The question thus arises as to whether the fetal and birth weight deficits induced by developmental toxicants may similarly lead to reprogramming of set points for physiological functions and responses that ultimately lead to cardiovascular and metabolic diseases in adulthood.

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utero are transitory, and therefore, have been argued to be innocuous. However, one can also view this "recovery" as catch-up growth during postnatal development, which is a key component to the manifestation of adult diseases in the Barker hypothesis. Does it mean that this population is, in fact, more susceptible to adult diseases and therefore assumes higher health risks? Indeed, the advent of the Barker hypothesis, if proven applicable to toxicant-induced fetal growth impairment, would require reconsideration of the risk assessment guidelines for developmental toxicity. Thus, teratologists may yet have to expand the scope of their research further still, not only to examine the anatomical structure of an organ and its functional maturation, but also to follow potential latent abnormalities of the system into adult age, given that the adult-onset diseases may have their origins in insults during fetal development.

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