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## **What a general paediatrician needs to know about early life programming**

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**Summary:** The process whereby early exposure to an adverse environment has an influence on later life outcomes has been called ‘early life programming’. Whilst epidemiological evidence for this has been available for decades, only in recent years have the mechanisms, in particular epigenetic modifications, for this process begun to be elucidated. We discuss the evidence for early life programming, the possible mechanisms, how effects may be transmitted across generations, and conclude by looking at some examples relevant to general paediatrics.

### **Childhood and later disease risk**

The fact that early exposures might have an influence on health outcomes later in life has been recognised since the first half of the twentieth century. In 1933 Kermack and colleagues [1] analysed historic death rates data for England, Scotland and Sweden and noted that “the figures behave as if the expectation of life was determined by the conditions which existed during the child's earlier years”. They further speculated that “improvement in infantile mortality is dependent in large measure on improvement in maternal health”. In 1977 Forsdahl [2] correlated higher infant mortality in Norway with a later increased risk of death from cardiovascular causes. He proposed poverty in childhood and adolescence followed by prosperity as a risk factor for cardiovascular disease (CVD), and hypothesised that “some form of permanent damage caused by a nutritional deficit” might be involved.

### **The Barker hypothesis**

Ongoing epidemiological work continued to show an association between low birthweight and a higher risk of CVD, stroke, the metabolic syndrome and osteoporosis in later life. Barker and colleagues, in a series of papers (e.g.[3]) argued that a fetus faced with undernutrition slows its growth rate to reduce its nutritional requirements, but this period of undernutrition might also lead to reduced function in key organs, altered metabolic and endocrine feedback loops, and an increased vulnerability to adverse environmental stressors. Over time these ideas have developed into the DOHaD (Developmental Origins of Health and Disease) concept, whereby early life exposures are thought to lead to “programming” of cardiovascular, neuroendocrine and metabolic systems, predisposing the individual to later life non-communicable diseases (NCD).

Some authors have put this concept of programming within an evolutionary paradigm with the idea of the “predictive adaptive response” [4]. They argue that these stereotyped responses to an adverse early life environment are adaptive in the short-term, and particularly when individuals continue to live in a resource poor environment represent the best way to guarantee they reach reproductive age themselves. However, in a resource rich postnatal environment such as that of the developed world, these programmed changes might have the (unanticipated) effect of predisposing affected individuals to an increased risk of NCDs in adulthood (Figure 1). Regardless of the validity of the idea of a

“predictive adaptive response”, in recent years focus has shifted from extremes of birthweight to how programming might occur across all pregnancies and in individuals with birthweights within the normal range.

### **Adverse environments *in utero***

A large number of human cohort studies have demonstrated a link between lower birthweight (suggesting *in utero* exposure to an adverse environment) and a higher risk of CVD, stroke, insulin resistance and type 2 diabetes in adulthood, in a variety of settings in the developed and developing world,[4] and these findings have been replicated extensively in animal studies.[5] In addition to cardiometabolic sequelae, low birthweight has also been related to increased risk of death from infectious causes,[6] altered immune function, an increased risk of asthma and atopic dermatitis,[7] and neurodevelopmental disorders such as attention deficit hyperactivity disorder (ADHD) [8] and schizophrenia.[9] The importance of the early life environment in influencing later disease risk has been highlighted by a recent analysis showing that the greater the number of adverse early-life risk factors an individual is exposed to, the greater the risk of overweight and obesity in childhood.[10]

Although many of the original studies focussed on poor maternal nutrition as a major contributor to low birthweight, a wide variety of endogenous and exogenous factors are now recognised to influence cardiovascular, respiratory, metabolic and neurodevelopmental outcomes in offspring. Animal models using uterine artery ligation to create *in utero* hypoxia have shown that offspring are at risk of cardiovascular and metabolic complications.[5] Similarly, human studies suggest that maternal hypertension [11] and cigarette smoking,[12] which both lead to *in utero* hypoxia, also increase the risk of CVD. Although the original epidemiological studies did not distinguish between intra-uterine growth restriction (IUGR) and prematurity as a cause of low birthweight, it is increasingly realised that prematurity itself is a major risk factor for the development of NCDs (reviewed in [13, 14]). Interestingly, maternal obesity/overnutrition during pregnancy, which commonly leads to *increased* birthweight, also associates with adverse offspring health outcomes which, perhaps surprisingly, are similar to those seen with undernutrition. In animal and human studies maternal overnutrition is linked with an increased predisposition to obesity, hypertension, hyperinsulinaemia, hyperglycaemia, and increased plasma triglycerides, cholesterol and leptin in offspring.[15] In humans, maternal obesity has also been linked to an increased risk of ADHD and problems with emotional regulation [16] and with premature mortality from cardiovascular events.[17]

Prenatal glucocorticoid overexposure is also associated with programmed effects. Epidemiological studies show that pregnant mothers exposed to a significant life event (death of a loved one, exposure to terrorism or a natural disaster) give birth to infants with a lower birthweight,[18] who have an increased risk of impaired cognition.[19] Maternal stress is also associated with effects on

neurodevelopment in neonates, manifesting as lower scores on neonatal assessment,[20] behavioural and emotional problems at the age of four,[21] decreased grey matter density [22] and lower cognitive and language abilities in childhood.[19] The children of mothers treated with glucocorticoids because of a risk that the unborn child had congenital adrenal hyperplasia appear to be at risk of worse cognitive function than controls.[23] Studies in rodent, sheep and non-human primate models show an association between antenatal glucocorticoid exposure (exogenous and endogenous), raised blood pressure (BP) and altered glucose-insulin homeostasis, neuro-endocrine function and behaviour.[24] Finally, a variety of other adverse environmental factors have been shown to impact on neurodevelopmental outcomes: maternal infection, alcohol consumption, recreational drug use, treatment with certain medications (e.g. sodium valproate) and prenatal exposure to toxins such as arsenic and lead have all been associated with an increased risk of adverse neurodevelopmental outcomes including schizophrenia and autism [reviewed in 25].

### **Postnatal factors which influence the risk of developing NCD**

The long-term consequences of early exposures are modulated by the postnatal environment. Early postnatal growth patterns influence disease risk, for example, in a trial involving a cohort of small for gestational age (SGA) infants, those randomised to a high protein formula who had greater weight gain had significantly higher BP at 6-8 years [26] and conditional gains in abdominal circumference also associate with higher childhood BP.[27] Three large cohort studies have shown that excessive weight gain in infancy is associated with an increased risk of greater total fat mass and percentage body fat, lower insulin sensitivity and higher systolic BP in childhood.[28] These relationships are complex however; extensive data from the Helsinki birth cohort study and the Hertfordshire cohort show that low weight at one year of age but an early ‘adiposity rebound’ associates with a higher risk of CVD and type 2 diabetes and that boys who were born small but were tall at school entry had a 6-year reduction in lifespan.[29, 30] These data are supported by animal studies, for example in mice, poor fetal growth resulting from maternal protein restriction, followed by rapid postnatal growth results in reduced life span.[31]

Because of phenotypic similarities between adults born SGA and prematurely, some researchers have postulated that whilst in low birthweight infants the adverse environment is experienced *in utero*, in preterm infants these environmental challenges occur postnatally.[32] Preterm birth, regardless of birthweight relative to gestation (i.e. without evidence of *in utero* growth restriction), has in itself has been associated with a reduction in insulin sensitivity,[32, 33] changes in the endocrine regulation of childhood growth,[34] and increased adiposity.[35] Similar to SGA infants, rapid early weight gain may also be detrimental for preterm babies: in a randomised controlled trial which allocated preterm infants to high or lower nutrient diet, those with the most rapid weight gain in the first two weeks of life showed evidence of insulin resistance in adolescence [36] and other studies report that upward

centile crossing for weight in infancy [37] and childhood [38] are associated with insulin resistance and higher BP.

However, it may not only be low birthweight and preterm infants who are susceptible to the influences of postnatal diet. Maternal obesity and overnutrition are also risk factors for childhood obesity, and this is exacerbated by a high energy diet in infancy.[39] In rodent models of maternal overfeeding, offspring are predisposed to obesity and metabolic abnormalities, and this effect is amplified when the offspring are exposed to high-fat diets following weaning.[40] Finally, there may be an important influence of childhood exposure to stressful experiences; severe stress during childhood is a well-known risk factor for both mental health disorders and CVD.[41]

### **Mechanisms**

A consistent criticism of the DOHaD concept has been the difficulty in disentangling the putative effects of programming from the shared genetic and environmental influences affecting offspring. Nevertheless, studies in humans and in animal models have attempted to address a number of potential mechanisms.

Programming may exercise its long-term effects via structural changes in organs. Human infants born after IUGR have reduced numbers of nephrons, [42] increasing the risk of hypertension in adulthood. Prematurity has also been associated with a reduction in capillary density, thought to be related to an increased risk of hypertension.[43] In animal models, antenatal glucocorticoid exposure leads to reduced nephron number and changes in cardiac noradrenergic innervation [24] together with a reduction in pancreatic beta-cell growth, a risk factor for type 2 diabetes.[44] Maternal obesity impacts on offspring body fat and muscle composition, which may contribute towards the development of insulin resistance.[15] Antenatal glucocorticoids affect hippocampal growth, and associate with delayed maturation of neurons, myelination, glia and vasculature.[24]

Programming may also lead to longer term hormonal changes. In rodents, prenatal glucocorticoid excess affects the renin-angiotensin-aldosterone system and both undernutrition and glucocorticoid overexposure affect glucose-insulin homeostasis in a variety of animal models.[24] Maternal obesity/overnutrition during pregnancy leads to programmed alterations in the brain, in particular the hypothalamus, which may impact on appetite control.[15] Importantly, nutrient deficiencies, low birthweight and maternal stress affect hypothalamic-pituitary-adrenal (HPA) axis development [45] and this may be a key mechanism linking the early life environment with later disease risk. Exposure to an adverse *in utero* environment is associated with altered HPA axis activity in childhood and adulthood [46, 47] which may relate to the higher risk of neurodevelopmental problems [48] and changes in HPA axis activity are also a risk factor for CVD in later life.[46] Finally, re-setting of the

HPA axis affects the development of key glucocorticoid responsive organs such as the kidneys, adipose tissue and pancreas, [45] linking structural and hormonal changes in the programming of later life disease.

There has been much interest in recent years in the role of epigenetic modifications in early life programming. Epigenetic modifications lead to changes in gene expression that are not explained by changes in DNA sequence, and during normal development, key developmental stages are characterised by epigenetic modifications that have the potential to be altered/disrupted by environmental cues. Epigenetic modifications include DNA methylation, histone marks and small, non-coding RNAs.

DNA methylation is crucial for normal development and is involved in cell differentiation, genomic imprinting and X chromosome silencing. A growing number of studies have described alterations in DNA methylation and gene expression in association with early life exposures. Adults exposed *in utero* to severe calorie restriction during the Dutch Hunger Winter of 1944/5 have reduced methylation at regions controlling the expression of the insulin-like growth factor 2 (IGF2), a key hormone in growth and development [49] and individuals exposed to an unbalanced diet *in utero* showed differential methylation at 11 $\beta$ -hydroxysteroid dehydrogenase type 2 (important in the regulation of BP), the glucocorticoid receptor (GR) and IGF2, in association with a risk of obesity and hypertension.[50] In rats, undernutrition and/or prenatal glucocorticoid overexposure influence DNA methylation of GR and IGF2 [51, 52] suggesting a mechanism by which adverse early life environments may affect hormonal homeostasis in offspring. Studies also suggest an association between maternal anxiety/depression and altered DNA methylation at genes important in HPA axis regulation and mood in the offspring.[53] Such effects are also seen with postnatal exposures: experience of childhood abuse associates with altered HPA and autonomic responses to stress and with altered DNA methylation in both peripheral blood and brain.[54-56]

DNA is organised around histones which can also be modified to control gene transcription. The development of glucose-insulin dyshomeostasis in rats after IUGR is associated with alterations in histone modifications at the *Pdx1* gene, a transcription factor regulating pancreatic development and  $\beta$ -cell differentiation, and additionally at the hepatic transcription factor *Hnf4a*, which associates with insulin resistance.[57] Gene expression can also be modulated by non-coding RNAs which are posited to regulate gene expression by transcriptional and post-transcriptional gene silencing. Studies in adipose tissue of rats exposed to a low protein diet *in utero* and in humans born at low birthweight have demonstrated persistent changes in a specific micro-RNA which may influence adipocyte differentiation and maturation.[58]

Nevertheless, although studies have shown epigenetic changes in association with early life exposures, the degree to which such changes are adaptive within the life course, occur as a consequence of the induced disease state or represent a pathophysiological response to adverse exposures remains to be determined as the mechanism(s) of these changes are elucidated. Recent studies suggesting that a significant proportion of neonatal DNA methylation changes are determined by genotype rather than environment [59] highlight the difficulties in distinguishing genetic from epigenetic inheritance.

### **Intergenerational transmission of programmed effects**

In addition to the effects of early life exposures on individual outcomes, there is increasing evidence that the effects of early life exposures may be transmitted non-genomically to subsequent generations.[60] Three human cohort studies suggest possible instances of the transmission of programmed effects, two through the female line [61, 62] and one patrilineally.[63] In this latter study, Kaati et al used detailed historical information on cohorts from Sweden which showed that food availability during the grandparents' childhood influenced the risk of CVD and diabetes in their grand-offspring. Intergenerational effects of early life environmental interventions have also been shown in a number of animal models [reviewed in 60]. Potential explanations for the transmission of programming effects across generations include persisting adverse environmental conditions, adverse *in utero* experiences resulting in programming effects in a female which then influence her own pregnancy or effects that are transmissible through gametes, such as DNA methylation changes or non-coding RNAs [60] (Figure 2). Finally, the experience of maltreatment during childhood may result in subsequent harsh and/or neglectful parenting in adulthood and this could be perpetuated across a number of generations.[64] These studies raise the possibility that antenatal, infant and childhood exposures may have consequences for multiple future generations. Indeed, intergenerational effects of the early life environment have been proposed as one mechanism to explain persisting health disparities amongst disadvantaged populations, potentially through effects on the HPA axis.[65, 66]

### **Significance for paediatricians**

An understanding of early life programming and its consequences is of clear importance for paediatricians who are ideally placed to identify those most at risk of later disease and to facilitate the development and implementation of interventions. Given that adverse early life environments may affect not only children's later life outcomes, but that of their own offspring, a paediatrician's role takes on important public health aspects.

Firstly, we can influence care from the beginning of life by emphasising the importance of good maternal health and antenatal care in optimising child and adult health. Addressing modifiable risk factors, including maternal obesity, excess gestational weight gain, maternal smoking and vitamin D levels and breastfeeding duration [10] could make a significant contribution to child health, and by implication, improve adult health. Indeed, paediatricians have been involved in the development of the National Institute for Health and Care Excellence evidence-based guidelines for the antenatal and postnatal care of women and babies ([www.nice.org.uk](http://www.nice.org.uk)) and these have been endorsed by the Royal College of Paediatrics and Child Health.

Since any effects of the *in utero* environment may be amplified as a consequence of early growth patterns, optimising early nutrition may be a key way in which paediatricians can influence later health. In preterm babies, studies are ongoing to develop nutritional strategies which optimise neurodevelopment and prevent extrauterine growth restriction without promoting the development of longer term metabolic complications,[67] however there is much less evidence for the optimal nutritional management of the SGA infant born at, or near term,[68] or in the management of the offspring of obese women. Overweight and obese women are less likely than lean women to exclusively breastfeed at 2 months of age and are more likely to introduce early weaning foods.[69] Breastfeeding may be protective against childhood obesity as a consequence of reduced protein content compared to formula, the presence of active hormones such as leptin, ghrelin and adiponectin which may influence appetite control, and poorer regulation of satiety with bottle feeding.[39] Thus, the provision of additional support for breastfeeding and improving the food choices of these women has the potential to reduce childhood overweight and obesity, with obvious consequences for the next generation.[69]

There is ongoing research (at present mainly in animal models) investigating therapeutic options to reverse or prevent the effects of *in utero* programming. Studies are ongoing to evaluate strategies for the prevention of CVD in individuals born with IUGR.[70] In animal studies, therapies which have been examined include micronutrient supplementation (eg. folate, glycine and choline) given during pregnancy to mitigate the effects of undernutrition, or statins for hypercholesterolaemia in pregnancy which may protect offspring against the conditioning effect of a high fat diet.[4] Similarly, there may be a role for leptin and statin therapy in those born SGA to modify the long-term hormonal and cardiovascular effects of an adverse antenatal environment.[4]

Since programming effects may also occur as a consequence of experiences in infancy and childhood, targeted interventions during infancy and childhood may improve later health. For example a randomised controlled trial of the effects of additional support for single, poor, deprived mothers during pregnancy and first two years of life resulted in improved educational performance in

childhood, less substance abuse at 12 years, and less criminal behaviour at 19 years in their offspring.[71] Importantly, evidence suggests that such interventions also have the potential to improve health outcomes across generations.[65, 66] Paediatricians could have a key role in identifying the factors which lead to the perpetuation of effects and in designing and implementing interventions to interrupt these intergenerational cycles [60] (Figure 3).

In conclusion, paediatricians are ideally placed to undertake research leading to an increased understanding of the mechanisms underpinning early life programming, to develop strategies for the early identification of disease risk and finally, to design and implement therapeutic strategies, with consequences that may improve health not only for the children we care for, but also for future generations.

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**Figure 1: Early life programming and later disease risk.** Exposure to an adverse early life environment results in changes which may maximise short- and long-term survival but which can result in an increased risk of disease in later life. Such programmed effects may also include strategies to maximise reproductive success and population/species survival.

**Figure 2: Potential mechanisms accounting for the transgenerational transmission of disease risk.** A) Persistence of an adverse environment leads to the re-induction of programmed effects in each subsequent generation. B) Maternal effects: the induction of programmed effects in the F1 offspring following *in utero* exposure lead to the induction of programmed effects on the developing F2 fetus and so on. C) Exposure to an adverse environment affects the developing F1 fetus AND has direct effects on the germ cells which will form the F2 generation and these changes are maintained in the germ cells for a number of subsequent generations.

**Figure 3: Intergenerational cycle of disease risk.** Early life programming may result in effects which persist across generations. Paediatricians are well-placed to identify those at risk and to develop and implement interventions at many points during the life-cycle.

Figure 1

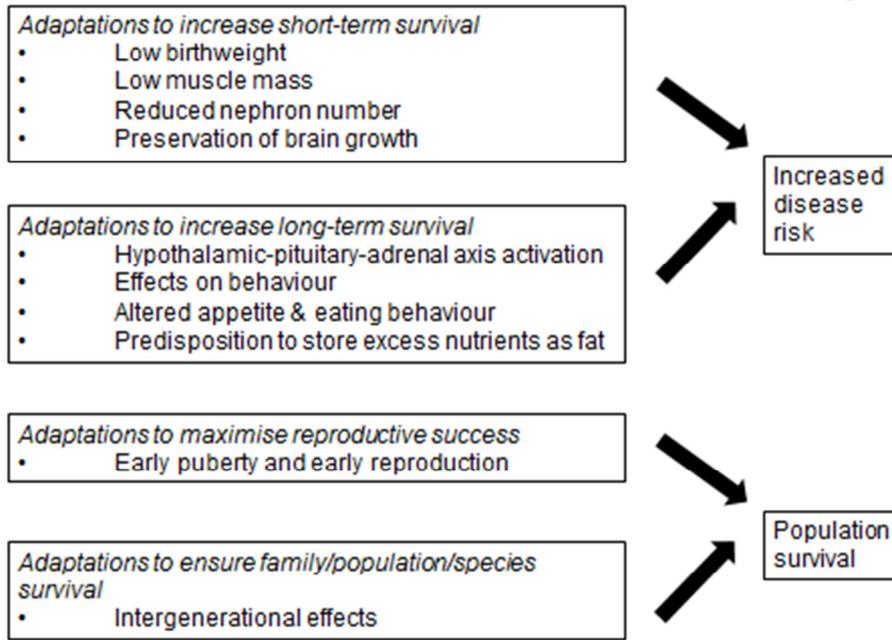


Figure 2

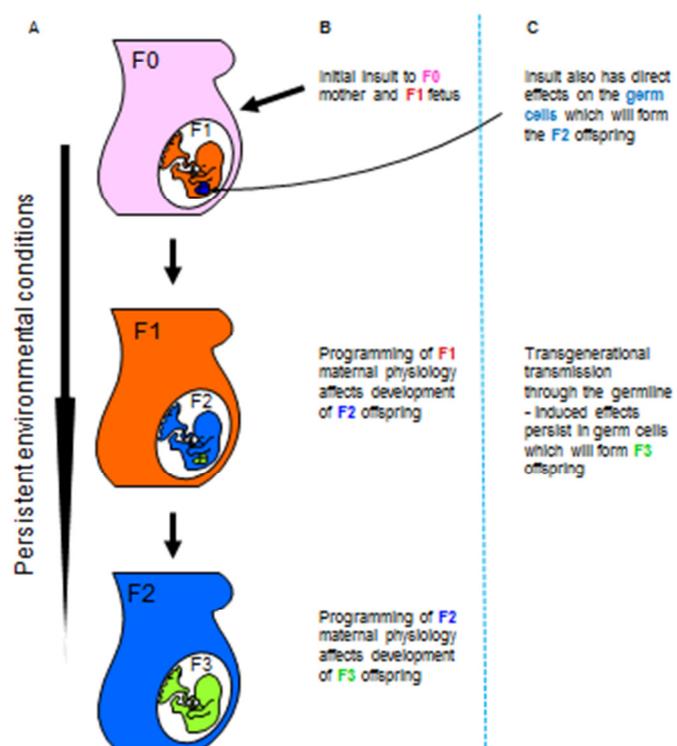


Figure 3

