Early origins of heart disease: Low birth weight and determinants of cardiomyocyte endowment

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Summary

1. World-wide epidemiological and experimental animal studies demonstrate that adversity in fetal life, resulting in intrauterine growth restriction (IUGR), programs the offspring for a greater susceptibility to ischemic heart disease and heart failure in adult life.

2. After cardiogenesis, cardiomyocyte endowment is determined by a range of hormones and signalling pathways that regulate cardiomyocyte proliferation, apoptosis and the timing of multinucleation/terminal differentiation.

3. The small fetus may have reduced cardiomyocyte endowment due to the impact of suboptimal intrauterine environment on the signalling pathways that regulate cardiomyocyte proliferation, apoptosis and the timing of terminal differentiation.

Introduction

World-wide epidemiological and experimental animal studies demonstrate that adversity in fetal life, resulting in intrauterine growth restriction (IUGR), programs the offspring for a greater susceptibility to ischemic heart disease and heart failure in adult life.¹⁻⁹ It is currently not clear how exposure to reduced substrate supply in utero can alter heart health some fifty years later. However, the human heart undergoes considerable maturation in utero, such that the majority of cardiomyocytes, present shortly after birth, beat for a lifetime.¹⁰⁻¹³ This has led to a significant body of research focusing on the regulation of cardiomyocyte maturation, endowment and growth in utero, particularly in the last half of pregnancy; and how specific insults at critical periods of development can alter the profile of cardiomyocytes present before and after birth. This review will focus on new insights into the regulation and consequences of IUGR on cardiomyocyte endowment.

Cardiomyocyte development

After cardiogenesis, the fetal heart initially grows as a consequence of mononucleated cardiomyocyte proliferation. In the last trimester and shortly after birth, these mononucleated cardiomyocytes cease proliferating, due to the absence of karyokinesis and/or cytokinesis.¹⁴ The final endowment of cardiomyocytes in the newborn heart is the result of a highly orchestrated balance between the creation of cardiomyocytes from cardiac progenitor cells in early gestation, subsequent cardiomyocyte proliferation across gestation, apoptosis and the critical timing of terminal differentiation. Typically, perturbations during pregnancy result in IUGR in the second half of pregnancy,¹⁵ therefore this review will include the regulation of cardiomyocyte endowment after cardiogenesis.

Cardiomyocyte proliferation

During development, cardiomyocyte proliferation is regulated by paracrine factors secreted from the epicardium,¹⁶ endocardium¹⁶⁻¹⁷ and fibroblasts,¹⁸ in addition to endocrine growth hormones.11 In response to mitotic stimuli, D-type cyclins and their catalytic partners CDK4 and CDK6 accumulate in the nucleus, which then phosphorylate and deactivate retinoblastoma protein (Rb), enabling cell cycle progression from the first gap phase (G_1) to initiate DNA synthesis (S phase).¹⁹ Cyclin D/CDKs facilitate the cell cycle by sequestering CDK inhibitors, p21^{Cip1} and p27^{Kip1}, allowing S phase initiation and progression through the activity of cyclin E/CDK2 and cyclin A/CDK2, respectively²⁰ (Figure 1). Cardiomyocyte proliferation is initiated through mainstream canonical mitogenic signalling pathways, such as the phosphoinositide 3-kinase (PI3K)/Akt and Ras/extracellular signal-related kinase (ERK) pathways.^{14,16,21-22} Specifically, the PI3K/Akt pathway promotes proliferation by phosphorylating and deactivating glycogen synthase kinase-3 β (GSK-3 β). GSK-3 β acts as a negative regulator of proliferation by phosphorylating cyclin D1,²³ causing its nuclear export and proteasomal degradation, thereby preventing progression from G1 to S phase. Activation of Akt also reduces the expression of p21^{Cip1} and p27^{Kip1} through FOXO transcription factors,²⁴ which in turn promote both CDK2 activity and progression to S phase. The Ras/ERK pathway increases expression of cyclin D1 both directly²⁵ and indirectly through down-regulation of



Figure 1. Proliferation of fetal mononucleated cardiomyocytes is regulated by multiple signalling pathways that stimulate or inhibit cyclins and cytokinesis. Promotion of cell cycle progression is indicated by black lines and inhibition is indicated by red lines. FGF-2: fibroblast growth factor 2, FGFR: fibroblast growth factor receptor, NRG-1: neuregulin-1, ErbB2/4: heterodimer of ErbB2 and ErbB4, IGF-1: insulin-like growth factor-1, IGF-2: insulin-like growth factor-2, IGF-1R: insulin-like growth factor-1 receptor, CAs: catecholamines, β -AR: β adrenergic receptor, Ang-II: angiontensin II, AT-R: angiotensin receptor, ANP: atrial natriuretic peptide, PI3K: phosphoinositide-3 kinase, Akt: protein kinase B, ERK, extracellular signal-related kinase, GSK-3 β : glycogen synthase kinase-3 β , T₃: thyroid hormone, CDK: cyclin dependent kinase. G₁: first gap phase, S: DNA synthesis phase, G₂: second gap phase, M: mitosis, G₀: gap zero phase (resting/quiescent).

anti-proliferative genes such as Tob1 and JunD.²⁶⁻²⁹ Recent data suggest that ERK and Akt activation leads to phosphorylation and inhibition of p27^{Kip1}.³⁰ ERK is also required for the translocation of CDK2 to the nucleus³¹ and its subsequent phosphorylation,³² which results in cyclin E association and cell cycle progression from G₁ to S phase. There is potential cross talk between the PI3K/Akt and Ras/ERK mitogen signalling pathways.²⁷

Studies involving mice, rats and chickens have contributed greatly to the understanding of cardiomyocyte proliferation. *In vitro* administration of extracellular mitogens such as fibroblast growth factor (FGF) -2³³ and neuregulin-1 (NRG-1)³⁴ stimulate proliferation of cardiomyocytes from embryonic chickens and fetal rats, respectively. The mitotic actions of both FGF-2 and NRG-1 are dependent on the activation of the PI3K/Akt pathway.³⁵⁻³⁶ Additionally, *in vitro* exposure to insulin-like growth factor (IGF) -1³⁷ and IGF-2³⁸ results in greater DNA synthesis in fetal cardiomyocytes from mice and rats, respectively. Furthermore, conditional knockout of their receptors, insulin receptor (*INSR*) and IGF1 receptor (*IGF1R*), in the myocardium of embryonic mice, results in decreased ventricular cardiomyocyte proliferation in the first half of gestation.³⁹ Activation of IGF1R leads to activation of both the PI3K/Akt and Ras/ERK signalling pathways,⁴⁰ however, less is known about the specific mitotic signalling pathways downstream of the INSR in cardiomyocyte proliferation, such that blocking β-adrenergic receptor (β-AR) activation *in vivo*, in neonatal

rats, decreases cardiomyocyte mitosis and deactivates p70 ribosomal protein S6 kinase, which is downstream of PI3K/Akt. 41

In addition to the activation of mitotic signalling pathways, proliferation can be regulated by cytokinetic mechanisms. Embryonic mice with cardiac specific deletion of Survivin and neonatal rats with Survivin small interfering RNA (siRNA) knockdown have fewer cardiomyocytes, due to a reduction in proliferation.⁴² Survivin is a key regulator of mitosis and cytokinesis because it is a component of the chromosomal passenger complex, essential for appropriate chromosomal separation and cytokinesis (reviewed by Lens *et al.*, 2006⁴³). It is currently unknown if IUGR influences the regulation of components of the chromosomal passenger complex.

Recent studies by Heallen et al. demonstrated that the canonical Wnt signalling pathway, which is an essential regulator of pre-cardiac mesoderm cell proliferation and differentiation into cardiomyocytes, promotes cardiomyocyte proliferation.44 Through the use of conditional knockouts of Salv to selectively inhibit Hippo signalling in mice, Haellen et al. demonstrated that Hippo signalling is essential for the appropriate control of cardiomyocyte proliferation and heart size; due to its inhibition of transcription factors that are promoted by the Wnt/ β -catenin signalling pathway. Ablating Hippo signalling led to embryos with larger hearts containing more cardiomyocytes. Interestingly, this cardiomegaly was specifically due to exaggerated cardiomyocyte proliferation and was not associated with altered fibroblast, smooth muscle cell or cardiac progenitor cell proliferation; and was observed in the left and right ventricle, despite these cardiomyocytes originating from different heart fields.44

Pathological conditions in fetal life that result in fetal growth restriction occur predominantly in late gestation¹⁵ and often lead to changes not only in the endocrine environment, but also in both preload and afterload. Therefore, the effects of these changes on cardiomyocyte proliferation have been studied in sheep, where cardiomyocyte maturation begins in late gestation as in humans, unlike postnatal life such as rats, mice and chickens (Table 1). The endocrine effects have been studied in sheep models of IUGR (reviewed by Morrison, 2008⁴⁵) and show that there is a decrease in plasma IGF-1,⁴⁶ glucose⁴⁷ concentrations and an increase in plasma cortisol,⁴⁷ noradrenaline⁴⁸ and adrenaline⁴⁸ concentrations in late gestation. Different models of IUGR in sheep cause no change⁴⁹ or an increase⁵⁰⁻⁵² in mean arterial pressure.

In late gestation, increased cardiac systolic load⁵³⁻⁵⁴ and a range of hormonal and growth factors including IGF-1⁴⁰ and angiotensin II (Ang-II)⁵⁵ have been shown to stimulate proliferation of mononucleated cardiomyocytes (reviewed by Thornburg *et al.*, 2011¹¹). Ang-II acts through the Ras/ERK pathway in cardiomyocytes,⁵⁵ but evidence from mouse embryonic stem cells suggests it may also activate the PI3K/Akt pathway.⁵⁶

Studies in sheep fetuses provide conflicting results regarding the regulation of cardiomyocyte proliferation by cortisol. Increased cortisol concentrations in late gestation leads to maturation of fetal organs prior to birth, but a comprehensive study in sheep fetuses identified cortisol as a potent cardiomyocyte mitogen.⁵⁷ In contrast, a similar intrafetal infusion of cortisol has been reported to decrease DNA content in the left ventricle⁵⁸ and adrenalectomized sheep fetuses exhibit greater cardiomyocyte proliferation, thus suggesting cortisol inhibits progression through the cell cycle.⁵⁹ The signalling pathway that links cortisol to proliferation of cardiomyocytes, however, remains unclear.

In the late gestation sheep fetus, cardiomyocyte proliferation is inhibited in the presence of reduced cardiac systolic load,⁶⁰ thyroid hormone $(T_3)^{61}$ and atrial natriuretic peptide (ANP)⁶² (reviewed by Thornburg *et al.*, 2011¹¹). Specifically, T_3 increases the protein abundance of the CDK inhibitor p21^{Cip1}, while simultaneously decreasing the protein abundance of cyclin D1.⁶¹ ANP does not inhibit the basal rate of proliferation, however, Ang-II stimulated proliferation is inhibited due to reduced Akt and ERK activity⁶² (Figure 1).

Apoptosis of cardiomyocytes

Apoptosis is critical for appropriate cardiovascular development (reviewed by Poelmann & Gittenberger-de Groot, 2005⁶³ and Porrello et al., 2008⁶⁴) and is tightly regulated and controlled by two main pathways; the intrinsic and extrinsic pathways, which regulate apoptosis through mitochondrial activity and death receptors, respectively.65 Upon release of cytochrome C from the mitochondria or ligation of the death receptors (DRs), 'initiator' caspases such as procaspase 9 and procaspase 8, are cleaved to their active form (Figure 2). Activation of 'initiator' caspases results in the cleavage and activation of 'effector/executioner' caspases (e.g. 3, 6 and 7), which cause the biochemical and morphological changes associated with apoptotic cell death.⁶⁶⁻⁶⁷ The apoptotic pathway is regulated at various levels by members of the Bcl-2 family of proteins, consisting of both pro-apoptotic (Bad, Bax, Bak, tBid, Bim and Bmt) and anti-apoptotic (Bcl-2 and Bcl-X₁) elements.⁶⁵ Anti-apoptotic proteins of the Bcl-2 family act at the mitochondrial level to block the release of cytochrome C, whereas pro-apoptotic proteins inhibit the action of the anti-apoptotic proteins.⁶⁵ The cleavage and activation of anti-apoptotic Bid by caspase 8 allows for the extrinsic pathway to initiate the intrinsic (mitochondrial dependent) apoptosis pathway.⁶⁸

In the rat heart, there is very little cardiomyocyte apoptosis either prenatally $(1.4-2\%)^{69-70}$ or at 21 days postnatal life.⁷¹ However, in the first day of postnatal life, there is considerable apoptosis,⁷¹ especially in the right ventricle, which undergoes extensive remodelling due to the abrupt changes in blood flow patterns and circulatory resistance shortly after birth (reviewed by Smolich, 1995⁷²). It is important to note that in the rat heart, this period of cardiac remodelling and apoptosis corresponds to a period where the majority of cardiomyocytes are mononucleated and still capable of proliferating⁷³ (Table 1). In the sheep, this remodelling and apoptosis occurs at a time when the majority of cardiomyocytes are terminally differentiated.⁷⁴



Figure 2. Apoptosis is critical for cardiac development. It can be mediated by the mitochondrial dependent intrinsic pathway and the death receptor mediated extrinsic pathway. DRs: death receptors, FADD: Fas-Associated protein with Death Domain.

Apoptosis in the late gestation sheep heart is also minimal (<0.05% of cardiomyocytes).⁷⁵ At birth, the majority of human cardiomyocytes are mononucleated⁷⁶ (Table 1), however, they lose the ability to proliferate shortly after birth. It is currently unclear if the remodelling of the right ventricle occurs when human cardiomyocytes are still capable of proliferating.

Multinucleation and terminal differentiation of cardiomyocytes

In late gestation and shortly after birth, mononucleated cardiomyocytes cease proliferating due to the absence of karyokinesis and/or cytokinesis. If a cardiomyocyte undergoes karyokinesis in the absence of cytokinesis, it becomes a multinucleated (predominantly binucleated) cardiomyocyte and is terminally differentiated ^{10, 77} (Table 1). Unlike humans, mice and rats are born with hearts comprised of mononucleated cardiomyocytes that do not undergo binucleation or cease proliferating until after birth.73, 78 Consequently, injuries to the neonatal mouse heart before 7 days of age can result in cardiomyocyte regeneration due to the proliferation of existing mononucleated cardiomyocytes.⁷⁹ To date the signals that prevent mononucleated cardiomyocytes from undergoing karyokinesis to become polyploid, or cytokinesis to become multinucleated, are not well understood. Studies in rats demonstrate that during binucleation there is a simultaneous downregulation of cyclins and CDKs associated with G_1/S and G_2/M transition and an upregulation of cyclins and CDKs associated with G_1 phase.⁸⁰ Recent studies by Di Stefano *et al.* demonstrate that simultaneous knockdown of CDK inhibitors p21^{Cip1}, p27^{Kip1,} and p57^{Kip2} by siRNAs in cultured neonatal and adult rat cardiomyocytes can result in entry to S-phase and a proportion of cardiomyocytes completing karyokinesis.⁸¹ Transgenic mouse studies have demonstrated that overexpression of cyclin D1 and cyclin G1 induce multinucleation,⁸²⁻⁸³ but it has been suggested by Naqvi *et al.*⁸⁴ that terminal differentiation may not be simply due to altered expression of genes that regulate the cell cycle.

Recent studies by Porrello *et al.* demonstrate that cell cycle withdrawal and multinucleation may be regulated by microRNAs (miRs).⁸⁵ By comparing the expression of miRs in cardiomyocytes from mice before (1 day of age) and after (10 days of age) the transition of the cardiomyocyte pool from proliferative mononucleated cardiomyocytes, Porrello *et al.* identified the miR-15 family member, miR-195, as the most upregulated miR during this period. Furthermore, premature overexpression

| Species | Length of gestation/ incubation (days) | Timing of binucleation (days) | | Mononucleated cardiomyocytes (%) | | Binucleated cardiomyocytes (%) | | Cardiomyocytes with >2 nuclei (n) (%) |
|---------|---|--|-------------------------------|---|--|---|--|--|
| | | Before birth/ hatching | After birth/ hatching | At birth/ hatching | Adult | At birth | Adult | Adult |
| Human | 280 | Begins by ~224 ⁷⁷ | unknown | ~90 ⁷⁶ | 74 ± 8 ⁹³ | 8.8 ± 5.3^{76} | 25.5 ± 8^{93} 57 ⁷⁶ 54-63 ⁹⁴ | ~0.5 (3n and 4n) ⁹³ |
| Rat | 21 | n/a | 4-12 ⁷³ | > 95 ⁷³ | 10-14 ⁹⁵ | 2.9 ± 1.8^{73} | 85-89 ⁹⁵ | 1-5% (3n and 4n) ⁹⁵ |
| Mouse | 19 | n/a | 3-10 ⁹⁶ | ~98 ⁹⁶ | < 8.5 ⁹⁶ | 1.5 ± 0.3^{96} | 91.5 ⁹⁶ | unknown |
| Chicken | 21 | n/a | Begins by 15 ⁹⁷ | 100 ⁹⁷ | 43.6 ± 4.6 at 42 days ⁹⁷ | 097 | 44.2 ± 2.8 at 42 days ⁹⁷ | ~11.5 at 42 days ⁹⁷ |
| Sheep | 145-150 | Begins 100 ⁷⁵ -110 ⁷⁴ | Ends ~4 ⁷⁴ | Between 16.8 \pm 2.6 and 21.1 \pm 2.2 ⁷⁴ | 8 ± 1.5 at 4-6 weeks ⁷⁴ | Between 78.9 \pm 2.2 and 83.2 \pm 2.6 ⁷⁴ | 92.0 \pm 1.5 at 4-6 weeks ⁷⁴ | unknown |
| Pigs | 114 | unknown | unknown | unknown | ~5 at 6 mo ⁹⁸ | unknown | ~12 at 6 mo ⁹⁸ | ~83 (3n-16n) at 6 mo ⁹⁸ |

Table 1. Timing of multinucleation and proportion of mononucleated and multinucleated cardiomyocytes in adult life in a range of species.

of miR-195 in embroyonic hearts resulted in smaller hearts that had a reduced number of cells in the cell cycle and a greater percentage of multinucleated cardiomyocytes at postnatal day 1, suggesting premature cell cycle arrest. Furthermore, postnatal knockdown of the miR-15 family resulted in a greater number of mitotic cardiomyocytes at 12 days of age, however, this did not involve an increase in the number of cardiomyocytes undergoing cytokinesis and suggests that the miR-15 family is not alone in preventing cardiomyocyte proliferation in postnatal hearts.

Despite the absence of proliferation in the adult heart under basal conditions,86 studies of human hearts after myocardial infarction suggest that a small proportion of cardiomyocytes are capable of cytokinesis,87 albeit insufficient to maintain/repair heart function. Similarly, in vitro stimulation of adult rat cardiomyocytes with NRG-1 causes DNA synthesis followed by completion of cytokinesis in approximately 0.6% of previously quiescent mononucleated cardiomyocytes.36 Engel et al.88 have identified the signalling molecule, p38 mitogen-activated protein (MAP) kinase (p38), as an inhibitor of adult cardiomyocyte cytokinesis. Inhibiting p38 in vitro results an approximate 3.8–fold increase in in adult cardiomyocytes undergoing cytokinesis after stimulation with FGF-1 compared to FGF-1 stimulation alone.⁸⁸ p38 has also been implicated as a potential regulator of cytokenetic genes such as components of the chromosomal passenger complex (Aurora B, INCENP and Survivin) and actin assembly genes such as Anillin.⁸⁸⁻⁸⁹ In addition, Engel et al.89 have identified Anillin recruitment and localization during anaphase and late cytokinesis as being essential for the completion of cytokinesis and its absence as a cause of binucleation.

It is not clear why mature mammalian

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cardiomyocytes have a limited capacity to proliferate, whilst cardiomyocytes from species such as newts and zebrafish retain the ability to replicate DNA and divide.⁹⁰⁻⁹¹ The physiological benefit of multinucleated cardiomyocytes is uncertain, but it has been proposed to be an adaptive response in cells with a high metabolic demand, such as skeletal muscle cells, where the capacity to generate twice the ribonucleic acid (RNA) for protein synthesis might be advantageous.14 Studies in sheep, where binucleation begins prenatally, as it does in humans, demonstrate that the maximum Ca2+-activated force and adult cardiac troponin I and C protein expression increase with the decrease in the percentage of mononucleated cardiomyocytes in late gestation.⁹² The timing of binucleation can be accelerated or delayed by alterations to the fetal environment, as discussed below.

Cardiomyocyte turnover and polyploidization in postnatal life

that terminal differentiation It appears of cardiomyocytes is complete soon after birth in humans and therefore it has been proposed that the human heart lacks the capacity to generate more cardiomyocytes postnatally. Although it is still accepted that adult cardiomyocytes, unless experimentally stimulated, do not undergo cytokinesis,¹⁴ studies carried out in several laboratories have identified the presence of human cardiac stem cells (hCSCs) that have the ability to generate new cardiomyocytes.⁹⁹⁻¹⁰¹ Currently, the contribution of hCSCs to the replacement and turnover of cardiomyocytes after birth is contentious. Some studies have suggested that ~50% of cardiomyocytes are replaced in a normal life span,¹²⁻¹³ whereas others suggest that the entire cardiomyocyte pool is replaced 11 to 15 times in the life of



Figure 3. The effect of PaO₂ on fetal growth. Both fetal body weight (A; $r^2 = 0.553$, P < 0.001, y = 0.21x - 0.05) and heart weight (B; $r^2 = 0.501$, P < 0.001, y = 1.45x + 1.20) are correlated with mean gestational PaO₂ in the late gestation sheep fetus (137-145 days; term ~150 days). Open circles are control fetuses, filled circles are fetuses exposed to placental insufficiency.

men and women respectively, with the maximum age of cardiomyocytes being 23 years.¹⁰² Studies in mice demonstrate that the annual renewal rate of cardiomyocytes is approximately 1.1%,^{96,103} which is similar to an estimation made from carbon dating studies in humans.¹²⁻¹³ Regardless of the actual rate of cardiomyocyte turnover, hCSCs are not capable of replacing enough of the cardiomyocytes lost due to aging and injury (*e.g.* due to myocardial infarction) to maintain cardiac function and prevent heart failure. Therefore, the impact of restrictions to fetal substrate supply on cardiomyocyte endowment at birth has implications for the individuals' vulnerability to cardiovascular disease in adult life.

Despite the inability of adult cardiomyocytes to undergo cell division, they do retain the ability to undergo DNA replication. In the human heart, 90% of cardiomyocyte nuclei are diploid (2c) shortly after birth, whereas the majority of adult cardiomyocyte nuclei are tetraploid (4c).¹⁰⁴ Ploidy is positively related to heart weight¹⁰⁵⁻¹⁰⁶ and increases further in response to injury.¹⁰⁶⁻¹⁰⁷ Studies in sheep, where the majority of binucleation and terminal differentiation occurs prenatally, identify that premature lambs have a greater percentage of tetraploid mononucleated cardiomyocytes compared with term controls at two months of age.¹⁰⁸ Studies in rats, where gastroenteritis was induced during the period of binucleation and terminal differentiation (4-12d postnatal), demonstrated that polyploidization can also be increased.¹⁰⁹ The regulation and advantage/disadvantage of polyploidization is not well understood. Cardiomyocyte polyploidization is suggested to be protective against hypoxia induced apoptosis, but maladaptive for aerobic metabolism.¹⁰⁹ Consequently, alterations to cardiomyocyte ploidy during development may have implications for adult heart health.

The effect of IUGR in species where cardiomyocyte maturation occurs after birth

Maternal protein restriction during pregnancy results in reduced birth weight, heart weight and number of cardiomyocytes at birth in rats.¹¹⁰ Lim and colleagues extended maternal protein restriction during pregnancy to the lactation period, because cardiomyocyte binucleation in rats occurs postnatally,¹¹¹ but found no difference in the total number of cardiomyocytes in the offspring compared with controls at four weeks of age.¹¹¹ These studies are important because they suggest the presence of a critical window during cardiomyocyte maturation when cardiomyocyte endowment can be rescued. In rats, they also suggest that matching the prenatal and postnatal environment until cardiomyocyte maturation is complete, may be beneficial, but the opposite is true for nephron number.¹¹² Moreover, these studies demonstrate that heart weight and cardiomyocyte number are positively related.110-111

Maternal hypoxia during late gestation results in reduced birth weight in rats and offspring who have a greater susceptibility to ischemia reperfusion (I/R) injury in adulthood.¹¹³⁻¹¹⁵ Furthermore, I/R injury in these adult offspring results in a larger infarct and diminished postischemic recovery of left ventricular function when compared to controls,^{113,115} which is accompanied by an increase in caspase 3 activity and cardiomyocyte apoptosis.¹¹³ This vulnerability may be due, in part, to decreased capillary density¹¹⁶ and cardiac remodelling that includes increased collagen I and III and fibrillar thickness and density.¹¹⁵ Interestingly, prior to I/R injury, these rats have the same body weight, heart weight and left ventricle weight compared to controls, but their cardiomyocytes have a larger cross sectional area.¹¹⁴ These data suggest that rats,

whose mothers were exposed to hypoxia during pregnancy, have fewer cardiomyocytes and a heart weight that has been maintained by greater hypertrophy of the remaining cardiomyocytes. This premise is supported by studies of the hypertrophic heart rat (HHR) model, which develops cardiac hypertrophy in the absence of increased blood pressure, by 2 months of age.¹¹⁷ At 2 days of age, these rats have smaller hearts containing smaller and fewer cardiomyocytes, which have become prematurely binucleated and exited the cell cycle.¹¹⁸ Unlike maternal protein restriction studies, IUGR caused by maternal hypoxia can disrupt the positive relationship between heart weight and cardiomyocyte number in the postnatal heart.¹¹⁴

Fetuses of dams exposed to hypoxia have increased cardiomyocyte apoptosis and premature binucleation and exit from the cell cycle in the heart in late gestation.⁶⁹ One would presume therefore that maternal hypoxia would result in decreased numbers of cardiomyocytes in offspring at birth. In the rat, based on the observed 'catch up' of cardiomyocyte endowment in offspring exposed to maternal protein restriction,¹¹¹ it is not known if a proposed deficit in cardiomyocytes at birth will be corrected after birth, before cardiomyocytes have completed terminal differentiation.⁷³ Similarly, it is not known if the proposed deficit in cardiomyocyte number in the heart of postnatal offspring exposed to maternal hypoxia is a direct consequence of fewer cardiomyocytes at birth or if it is due to greater apoptosis in the heart of the IUGR offspring compared to control offspring post weaning. Furthermore, it has yet to be answered if IUGR, in an animal model where binucleation occurs prenatally, or in fact in humans, results in a deficit in the number of cardiomyocytes before or after birth.

The effect of IUGR in species where cardiomyocyte maturation occurs before birth

To date, there have been no studies published identifying the effect of IUGR on cardiomyocyte endowment in a species where binucleation of cardiomyocytes begins before birth. A study in sheep, a species where cardiomyocyte maturation occurs prenatally, demonstrates that naturally occurring variation in birth weight changes cardiomyocyte endowment. Specifically, birth weight and body and heart weight at nine weeks of age were positively correlated with the number of left ventricular cardiomyocytes.¹¹⁹ These data support evidence from maternal protein restriction studies in rats, which show heart weight is positively related to total cardiomyocyte number.¹¹⁰⁻¹¹¹

The use of a sheep model of chronic fetal hypoxemia, caused by fetal anemia, has identified alterations in fetal heart growth and poor cardiovascular outcomes in the adult sheep.¹²⁰⁻¹²¹ Sheep models of placental insufficiency result in chronic fetal hypoxemia, hypoglycemia, hypercortisolemia, low birth weight and reduced heart weight in late gestation, which are endocrine and growth changes that also occur in human pregnancies of IUGR.¹²²⁻¹²⁴ Studies in our laboratory have shown that in a

model of chronic placental restriction, fetal body weight and heart weight are correlated with fetal arterial PO_2 (PaO₂) across late gestation, such that the greater the degree of hypoxemia, the greater the fetal growth restriction and reduction in heart growth (Figure 3). There is, however, no change in heart weight relative to body weight in IUGR compared to control fetuses.¹²²

Studies using two different sheep models of IUGR, induced by placental insufficiency, have investigated cardiomyocyte development. One involves surgical removal of endometrial caruncles in the non-pregnant ewe to reduce placental size (placental restriction, PR),¹²² while the other involves embolization of the uterine artery of the pregnant horn to reduce uteroplacental blood flow in late gestation (UPE).¹²³⁻¹²⁴ In both models, placental insufficiency caused a delay in the transition of mononucleated cardiomyocytes binucleated to cardiomyocytes (Figure 4). This delay in maturation is in direct conflict with the results from maternal hypoxia studies in rats, which demonstrated an acceleration of binucleation,69 reflecting the importance of cardiomyocyte maturation timing differences between species.



Figure 4. Regardless of the model of IUGR employed, IUGR in sheep results in an increased percentage of mononucleated cardiomyocytes across late gestation.¹²²⁻¹²⁴ Control, blue circles; uteroplacental embolization (UPE),¹²³⁻¹²⁴ red triangles; placental restriction (PR),¹²² red diamonds.

A reduction in fetal substrate supply changes cardiomyocyte growth patterns, but these changes are dependent on the timing, duration and severity of the placental insufficiency. For example, placental insufficiency by UPE for up to 20 days is associated with a decrease in the percentage of mononucleated cardiomyocytes undergoing proliferation,¹²³ however this is not observed in PR, where placental insufficiency has occurred over at least the last half of gestation.¹²² Interestingly, the reverse is true for cardiomyocyte size, where PR results in a decrease in the absolute size of cardiomyocytes,¹²² with no change in absolute cardiomyocyte size observed after UPE.¹²³ Despite

a reduction in absolute cardiomyocyte size in PR fetuses, the size of each cardiomyocyte is larger in relation to heart weight when compared to controls. Studies in sheep demonstrate that the size of cardiomyocytes relative to heart weight decreases with gestation in the normally grown fetus, but that this is delayed in the IUGR fetus suggesting that there may be fewer cardiomyocytes in the heart of the IUGR fetus (Figure 5). The use of both sheep models of placental insufficiency highlights how differences in the degree and timing of fetal insults can result in different cardiomyocyte phenotypes.



Figure 5. The effect of placental restriction (PR) on cardiomyocyte size. In the normally grown sheep fetus (open bars) the length of binucleated cardiomyocytes relative to heart weight decreases with increasing gestational age, however, this does not occur in the PR fetus (solid bars), such that there is an increase in the relative length of binucleated cardiomyocytes in the PR fetus compared to the normally grown fetus in late gestation.^{92,122} Different superscripts (e.g. a, b, c) denote a significant difference between gestational ages in the normally grown fetus. * denotes a significant difference between normally grown fetuses and PR fetuses at 139-146 days gestation (P < 0.05).

Postnatal cardiac consequences of IUGR

The adaptation of the fetal heart to a period of reduced substrate supply and decreased body growth has critical consequences for heart health in later life because at birth, the human heart contains most of the cardiomyocytes it will have for life.¹⁰⁻¹³ Consequently, in cases where the endowment of cardiomyocytes is reduced, the remaining cells will be required to increase in size in order to increase their capacity for contractile force generation, with a consequent increased risk of coronary heart disease.11,64,125 A reduction in cardiomyocyte endowment is not the only consequence of IUGR that may change the profile of cardiomyocytes present in the postnatal heart. Increasing evidence suggests the regulation of cardiomyocyte metabolism,¹²⁶⁻¹²⁷ contractility,128 protection/ survival^{113-114,129-130} and hypertrophy^{114,122,131} may each be affected by a reduced substrate supply *in utero*.

Concluding remarks

The data discussed in this review suggest that IUGR induced by reduced substrate supply in different species leads to alterations in cardiomyocyte development and may lead to reduced cardiomyocyte endowment. It is not known, however, if the observed changes to cardiomyocyte development are all induced by a common mechanism evoked by suboptimal substrate supply, or whether deficiencies in specific substrates, such as protein or oxygen, induce specific consequences to cardiomyocytes. This is important for the understanding of the mechanisms that regulate cardiomyocyte endowment. It is also not clear, if a window of time exists when cardiomyocyte endowment can be rescued. Studies in rats suggest that the early postnatal period may represent such a window, but it is not clear if this is due to the remodelling of the heart that occurs during postnatal changes in the circulatory system or due to the postnatal timing of cardiomyocyte terminal differentiation in rats. It is clear that further studies are required to address these critical issues and to determine whether or not intervention strategies are likely to be beneficial in restoring cardiomyocyte endowment.

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