

1 **Placental phenotype and the insulin-like growth factors: resource allocation to fetal growth**

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3 Amanda N. Sferruzzi-Perri<sup>a,1</sup>, Ionel Sandovici<sup>b</sup>, Miguel Constancia<sup>b</sup>, Abigail L. Fowden<sup>a</sup>

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5 <sup>a</sup>Centre for Trophoblast Research, Department of Physiology, Development and Neuroscience,  
6 Downing Street, University of Cambridge, Cambridge, UK CB2 3EG; <sup>b</sup>Metabolic Research  
7 Laboratories, MRC Metabolic Diseases Unit, Department of Obstetrics and Gynaecology and NIHR  
8 Cambridge Biomedical Research Centre, Robinson Way, Cambridge, UK CB2 0SW

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10 <sup>1</sup>Corresponding author:

11 Amanda Sferruzzi-Perri

12 Centre for Trophoblast Research,

13 Department of Physiology, Development and Neuroscience,

14 University of Cambridge,

15 Cambridge, UK CB2 3EG

16 Telephone: +44 (0) 1223333807

17 Email: ans48@cam.ac.uk

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22 **ABSTRACT**

23 The placenta is the main determinant of fetal growth and development *in utero*. It supplies all the  
24 nutrients and oxygen required for fetal growth and secretes hormones that facilitate maternal  
25 allocation of nutrients to the fetus. Furthermore, the placenta responds to nutritional and metabolic  
26 signals in the mother by altering its structural and functional phenotype which can lead to changes in  
27 maternal resource allocation to the fetus. The molecular mechanisms by which the placenta senses  
28 and responds to environmental cues are poorly understood. This review discusses the role of the  
29 insulin-like growth factors (IGFs) in controlling placental resource allocation to fetal growth,  
30 particularly in response to adverse gestational environments. In particular, it assesses the impact of  
31 the IGFs and their signalling machinery on placental morphogenesis, substrate transport and  
32 hormone secretion, primarily in the laboratory species, although it draws on data from human and  
33 other species where relevant. It also considers the role of the IGFs as environmental signals in linking  
34 resource availability, to fetal growth through changes in the morphological and functional  
35 phenotype of the placenta. As altered fetal growth is associated with increased perinatal morbidity  
36 and mortality and a greater risk of developing adult-onset diseases in later life, understanding the  
37 role of IGFs during pregnancy in regulating placental resource allocation to fetal growth is important  
38 for identifying the mechanisms underlying the developmental programming of offspring phenotype  
39 by suboptimal intrauterine growth.

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41 **KEY POINTS SUMMARY**

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- 43 • Size at birth is critical in determining life expectancy and is dependent primarily on the placental  
44 supply of maternal nutrients and oxygen.
- 45 • The insulin-like growth factors (IGFs) are important in controlling placental resource allocation to  
46 fetal growth during development via their impacts on placental morphogenesis, substrate  
47 transport and hormone secretion.
- 48 • Placental IGFs (particularly IGF2) alter in response to environmental challenges known to affect  
49 placental phenotype and fetal growth.
- 50 • IGFs have an important role in optimising fetal growth with respect to resource availability  
51 during pregnancy via actions on placental phenotype.

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55 **AUTHOR SUMMARY**

56 Amanda Sferruzzi-Perri received her Bachelor of Science degree with Honours and PhD degree from  
57 the University of Adelaide, Australia (in 2001 and 2007, respectively). In 2008, she received a CJ  
58 Martin Overseas Biomedical Fellowship from the NH&MRC to undertake research at the University  
59 of Cambridge, UK. Through the award of a Next Generation Fellowship from the Centre for  
60 Trophoblast Research in 2011 and a Dorothy Hodgkin Research Fellowship from the Royal Society in  
61 2014, Amanda has been using a variety of strategies to decipher the role of insulin-like growth  
62 factors and their signalling pathway, PI3K in maternal-placental-fetal interactions governing  
63 pregnancy success.

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66 **INTRODUCTION**

67 Intrauterine growth is a key determinant of lifespan. Babies born growth restricted or large for  
68 gestational age are at greater risk of perinatal morbidity and mortality than those of normal birth  
69 weight. Moreover, the “memories” of an altered environment and growth *in utero* can stretch  
70 beyond the perinatal period to influence health much later in life. Epidemiological studies in humans  
71 have shown that babies grown abnormally due to poor maternal nutrition are at heightened risk of  
72 developing conditions like type 2 diabetes, heart disease and obesity as adults, and of dying younger  
73 as a consequence (Gluckman *et al.*, 2005; Jansson & Powell, 2006). Similarly, manipulating  
74 intrauterine growth experimentally by varying maternal food intake, dietary composition, oxygen  
75 availability, endocrine status or utero-placental blood flow has been shown to program  
76 cardiovascular, metabolic and endocrine function of the adult offspring in a wide range of  
77 mammalian species (Gluckman *et al.*, 2005; McMillen & Robinson, 2005; Fowden *et al.*, 2006).

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79 As the interface between the mother and fetus, the placenta is one of the main determinants of  
80 intrauterine growth. It supplies all the nutrients and oxygen required for fetal growth as well as  
81 secreting hormones that influence maternal metabolism in favour of the fetal needs. Its  
82 morphological and functional characteristics, therefore, have an important role in determining the  
83 allocation of maternal resources to fetal growth. These characteristics include cell composition,  
84 surface area, barrier thickness, blood flow, vascularity, nutrient utilisation and the abundance and  
85 activity of the various transporter molecules (Fowden *et al.*, 2009; Sandovici *et al.*, 2012). Recent  
86 studies have shown that the placenta can respond to maternal nutritional and metabolic signals by  
87 altering these characteristics which, in turn, leads to changes in the placental capacity to supply  
88 resources to the fetus (Fowden *et al.*, 2009; Sandovici *et al.*, 2012). Thus, the placenta is a key

89 mediator in linking maternal environmental conditions to development of the fetus (Burton *et al.*,  
90 2016; Sferruzzi-Perri & Camm, 2016). However, the molecular mechanisms by which the placenta  
91 senses and responds to environmental cues during pregnancy are poorly understood. This review  
92 discusses the role of the insulin-like growth factors (IGFs) in controlling placental resource allocation  
93 to intrauterine growth, particularly in relation to maternal environmental conditions during  
94 pregnancy. It focuses primarily on small laboratory animals, like mice, rats and guinea pigs that are  
95 most commonly used for these studies but also draws on data from other species, including humans,  
96 where available.

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### 99 **THE INSULIN-LIKE GROWTH FACTORS**

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101 The insulin-like growth factors (IGFs), IGF1 and IGF2, are 7.5kDa single-chained polypeptides that  
102 promote growth, both before and after birth. They affect the metabolism, mitogenesis, survival and  
103 differentiation of a wide variety of cell types by binding to IGF receptors (IGF1R and IGF2R), insulin  
104 receptor (INSR) and a hybrid IGF1R-INSR receptor with varying affinity (Sferruzzi-Perri *et al.*, 2008;  
105 Fernandez & Torres-Aleman, 2012; Harris & Westwood, 2012). Their actions are influenced by at  
106 least six different IGF binding proteins (IGFBP-1 to IGFBP-6) and numerous IGF-related binding  
107 proteins, which alter access of the IGFs to their receptors and have been reviewed in detail  
108 elsewhere (Bach *et al.*, 2005; Bach, 2015; Clemmons, 2016). The main signalling receptor for the IGFs  
109 is IGF1R, which activates the phosphoinositide-3 kinase/protein kinase A (PI3K/AKT) and mitogen-  
110 activated protein kinase (MAPK) signalling pathways. IGF2 also binds to the IGF2R, which can lead to  
111 either IGF2 degradation or activation of the G-protein-coupled signalling pathway (Okamoto *et al.*,  
112 1990).

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114 The *Igf2* gene is subject to parental imprinting and only the paternal allele is expressed. It can be  
115 expressed by different promoters, of which P0 (*Igf2P0*) is specific to the placenta in mice (Moore *et al.*  
116 *et al.*, 1997). In mice, though largely not in humans, the *Igf2r* gene is also imprinted but in a reciprocal  
117 fashion to *Igf2* with expression from the maternal allele (Monk *et al.*, 2006). The IGFs (particularly  
118 IGF2), their receptors and signalling pathways are expressed by the placenta in many species and  
119 change in their abundance both developmentally and in response to environmental cues (Sferruzzi-  
120 Perri *et al.*, 2010). In many species, circulating IGF concentrations are higher during pregnancy than  
121 in the non-pregnant animal and also change in the mother and fetus with proximity to delivery  
122 (Fowden, 2003; Sferruzzi-Perri *et al.*, 2010). IGF2 is more abundant than IGF1 in both the maternal

123 and fetal circulations in all species studied to date, with the exception of mice (Fowden, 2003;  
124 Sferruzzi-Perri *et al.*, 2010). IGF2 is also more highly expressed than IGF1 by the placenta in all  
125 species studied to date (Sferruzzi-Perri *et al.*, 2010).

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## 128 **THE EFFECTS OF THE INSULIN-LIKE GROWTH FACTORS ON PLACENTAL PHENOTYPE**

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130 The effects of the IGFs on the placenta have been studied directly in two main ways. First, they have  
131 been given exogenously either to placental cultures *in vitro* or to pregnant animals *in vivo* to study  
132 placental growth, transport and endocrine function. Secondly, the *Igf* genes, their receptors and key  
133 molecules in their downstream pathways have been under- or over-expressed in genetically  
134 modified mice to determine the morphological and functional consequences for the placenta at  
135 different stages of pregnancy. While the functions of the placenta are common across species, its  
136 structure varies in terms of shape, organisation of trophoblast lineages, extent of invasion into the  
137 maternal uterus, and degree of interdigitation at the feto-materno interface (reviewed in depth  
138 elsewhere (Carter, 2007; Wooding & Burton, 2008; Roberts *et al.*, 2016)). For instance, the human  
139 and non-human primate placenta is composed of a series of highly branched structures, called villi.  
140 These contain a mesenchymal core that has fetal capillaries which are closely associated with an  
141 overlying syncytiotrophoblast layer. The syncytiotrophoblast is directly bathed in maternal blood and  
142 functions in both transport and hormone secretion. Cytotrophoblast cells, can fuse to form the  
143 syncytiotrophoblast or migrate from the villous tree into the decidua where they invade and  
144 remodel uterine spiral arteries to promote blood flow to the placenta. The syncytiotrophoblast is  
145 also bathed in maternal blood in the mouse, rat and guinea pig placenta. However, the mouse  
146 placenta is arranged into two morphologically and functionally distinct regions; the labyrinth zone  
147 (Lz) that is responsible primarily for transport and the junctional zone (Jz; also known as basal or  
148 interlobium region) which functions in uterine remodelling/invasion and hormone secretion. In  
149 ruminant species like the sheep and cow the placenta is comprised of individual placentomes which  
150 form at specialised sites called caruncles, in the uterine wall. The overlying trophoblast layer can be  
151 a syncytium (in sheep) or remains uni-cellular (columnar epithelium; in cows) and there no invasion  
152 of the maternal blood vessels by trophoblast cells. However, in sheep some trophoblast cells migrate  
153 and fuse with caruncle epithelial cells and play an endocrine role.

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## 157 **Exogenous administration of IGFs**

158 ***In vitro experiments***

159 IGF1 and IGF2 prevent apoptosis and enhance proliferation and migration/invasion of human  
160 placental villous explants, primary trophoblast cultures and trophoblast cell lines from the first  
161 trimester and term (Table 1). IGF1 also promotes the proliferation, invasion and survival of first  
162 trimester human placental fibroblasts (Miller *et al.*, 2005) and the differentiation of term trophoblast  
163 cells into syncytiotrophoblast (Bhaumick *et al.*, 1992; Milio *et al.*, 1994; Cohran *et al.*, 1996).  
164 Similarly, IGF1 stimulates proliferation and migration of murine ectoplacental cone trophoblast in  
165 culture (Kanai-Azuma *et al.*, 1993) and early pregnancy porcine trophoblast cells (Jeong *et al.*, 2014).  
166 Furthermore, IGF2 promotes differentiation of murine ectoplacental cone trophoblast and migration  
167 of ovine trophoblast cells *in vitro* (Kim *et al.*, 2008). Using receptor and pathway inhibitors and IGF  
168 analogues with selectivity for particular receptors, some of the molecular mechanisms mediating the  
169 actions of IGFs on the human placenta have begun to be identified *in vitro*. IGFs appear to mediate  
170 their proliferative and anti-apoptotic effects on trophoblast through activating IGF1R and triggering  
171 the MAPK and PI3K/AKT signalling pathways, respectively (Forbes *et al.*, 2008). IGFs also induce  
172 trophoblast migration and invasion through IGF1R, and possibly INSR with subsequent activation of  
173 MAPK and PI3K/AKT signalling pathways (Diaz *et al.*, 2007; Shields *et al.*, 2007; Forbes *et al.*, 2008;  
174 Mayama *et al.*, 2013). However, IGF2 may also signal via IGF2R and G<sub>i</sub> proteins, MAPK and Rho  
175 GTPase pathways to trigger trophoblast migration and invasion (McKinnon *et al.*, 2001; Shields *et al.*,  
176 2007; Harris *et al.*, 2011). Thus IGFs promote the growth of different cell lineages in the placenta via  
177 multiple mechanisms (Figure 1A).

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179 In addition to stimulating placental growth, both IGFs stimulate glucose and System A amino acid  
180 uptake and IGF1 increases System L activity but reduces lipoprotein lipase activity in human  
181 trophoblast *in vitro* (Table 1). However, these changes in nutrient uptake do not always track with  
182 the expression of the transporter genes or proteins, suggesting that the IGFs may also affect post-  
183 transcriptional/translational mechanisms (Fang *et al.*, 2006; Jones *et al.*, 2013; Jones *et al.*, 2014).  
184 Indeed, IGF1 was recently shown to stimulate glucose transporter capacity by increasing the  
185 translocation of GLUT1/SLC2A1 to the trophoblast plasma membrane (Baumann *et al.*, 2014). In  
186 culture, IGF1 prevents the release of the vaso-constrictors, prostaglandin E and F, and thromboxane,  
187 by the term human placenta and reduces the agonist-mediated vasoconstriction of human  
188 myometrial arteries (Siler-Khodr *et al.*, 1995; Corcoran *et al.*, 2012). *In vivo*, these effects could  
189 increase utero-placental blood flow and substrate transfer in late gestation. Both IGF1 and IGF2 also  
190 enhance trophoblast endocrine capacity in culture. IGFs increase the secretion of hormones  
191 including progesterone, human chorionic gonadotrophin and placental lactogen *in vitro* although

192 others, like placental growth hormone may not be affected (Maruo *et al.*, 1995; Zeck *et al.*, 2008;  
193 Rak-Mardyla & Gregoraszczyk, 2010). In addition, IGF2 simulates the differentiation of hormone-  
194 producing murine and ovine trophoblast *in vitro* (Kanai-Azuma *et al.*, 1993; Kim *et al.*, 2008). Thus,  
195 IGFs have the capacity to promote growth, hormone secretion and substrate transport capacity of  
196 the placenta.

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### 199 ***In vivo experiments***

200 Treatment of guinea pig dams with either IGF1 or IGF2 in early-mid pregnancy increases fetal weight  
201 near term [Table 1; (Sferruzzi-Perri *et al.*, 2006)]. With exogenous IGF1, placental Lz area and *Igf2*  
202 gene expression is reduced during the treatment, even though fetal weight is increased already in  
203 mid pregnancy (Sohlstrom *et al.*, 2001; Sferruzzi-Perri *et al.*, 2007b; Standen *et al.*, 2015). Whilst  
204 there is no sustained effect of either IGF on placental weight, IGF2 increases the volume and surface  
205 area of the transport Lz, near term [Table 1, (Sferruzzi-Perri *et al.*, 2006)]. Development of the  
206 placental exchange region was further enhanced when the IGF2R-selective synthetic analogue,  
207 Leu<sup>27</sup>-IGF2 was administered maternally (Sferruzzi-Perri *et al.*, 2008). In mice, maternal Leu<sup>27</sup>-IGF2  
208 treatment from day 13 of pregnancy halves the number of fetuses naturally growth-restricted within  
209 the litter near term (Charnock *et al.*, 2016). Taken together, these findings suggest that maternal  
210 IGF2 in early gestation may act, in part, via the IGF2R to enhance functional development of the  
211 placenta with beneficial impacts on fetal growth. However, caution is warranted as part of the  
212 effects of Leu<sup>27</sup>-IGF2 could be due to the displacement of endogenous IGF2 and its subsequent  
213 interaction with IGF1R and INSR in the placenta.

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215 Exogenous IGFs also modify the functional capacity of the placenta to supply resources for fetal  
216 growth. In the late pregnant ewe, increasing IGF1 in the fetal circulation increases amino acid and  
217 glucose uptake by the placenta but may reduce materno-fetal transfer of these substrates, lactate  
218 production and the number of placentomes (Table 1). Increasing IGF1 in the maternal circulation  
219 also alters placental metabolic function in the pregnant ewe near term; glucose transfer capacity  
220 and lactate production are enhanced by an acute infusion of IGF1 (Liu *et al.*, 1994). In guinea pigs,  
221 placental delivery of glucose and/or neutral amino acids to the fetus is increased in late gestation by  
222 chronic maternal IGF treatment in early-mid pregnancy (Table 1). This enhanced placental transfer in  
223 late gestation is partly due to increased expression of nutrient transporters (System A amino acid;  
224 *SNAT2/Slc38a2*) by IGF1 in mid pregnancy and improved development of the exchange region by  
225 IGF2 in late pregnancy (Sferruzzi-Perri *et al.*, 2006; Sferruzzi-Perri *et al.*, 2007b). In mice, the

226 variability in System A amino acid transport capacity and conceptus weight within the litter is  
227 abolished by maternal Leu<sup>27</sup>-IGF2 (Charnock *et al.*, 2016) and data suggest that IGFs may have most  
228 benefit for improving growth of the smallest pups. Indeed, maternal Leu<sup>27</sup>-IGF2 improves the weight  
229 of fetuses that are growth restricted due to a lack of the endothelial nitric oxide gene and reduces  
230 the number of pups below the fifth centile of the wild-type population in late gestation (Charnock *et al.*,  
231 2016). In addition to improving placental transport function, exogenous IGFs also affect  
232 endocrine capacity *in vivo* (Figure 1B). Maternal IGF2 treatment simulates the development of the  
233 endocrine I<sub>2</sub> of the rat placenta (Van Mieghem *et al.*, 2009) and exogenous IGF1 and IGF2 increase  
234 placental pro-renin activation in guinea pigs (Standen *et al.*, 2015). Thus, IGFs may also increase fetal  
235 resource supply through changing placental endocrine function and thus maternal adaptations to  
236 pregnancy, however further studies are warranted.

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238 To circumvent possible confounding effects of systemic IGF treatment on the mother, approaches  
239 are being developed to target IGFs to the placenta. In mice, adenoviral-mediated site-specific  
240 intraplacental transfer of the *Igf1* gene on day 14 of pregnancy, increases the area of the placenta,  
241 the size of the L<sub>2</sub> and of the maternal and fetal facing areas three days later, although there is no  
242 change in conceptus weight [Table 1; (Katz *et al.*, 2009)]. In response to liposome-mediated targeting  
243 of IGF2 to the mouse placenta, placental growth is also increased although fetal weight is not  
244 affected (King *et al.*, 2016). In rabbits, the weight of natural runt fetuses in the litter is increased two  
245 days following placental *Igf1* transgene delivery without a change in placental weight however how  
246 it impacts on structure and function of the placenta remains unknown (Keswani *et al.*, 2015). These  
247 data suggest that targeting of IGF delivery to the placenta may prove an effective method of  
248 improving placental function and, thus, fetal growth, particularly when fetoplacental growth is  
249 impaired.

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### 252 **Genetic manipulation of the IGF system**

253 In mice, knockout of the *Igf2* gene in the entire conceptus or predominantly within the fetal or  
254 trophoblast cell lineages leads to placental and fetal growth restriction, with the greatest reduction  
255 in growth seen with ubiquitous *Igf2* loss (Table 2). Similarly, a heterozygous deficiency in the PI3K-  
256 p110 $\alpha$  (*Pik3ca*; homozygous deficiency is lethal) or complete ablation of the AKT1 (*Pkba*) or MAPK1  
257 (*Erk2*) genes, causes fetoplacental growth restriction (Cho *et al.*, 2001; Hatano *et al.*, 2003; Yang *et al.*,  
258 2003; Yung *et al.*, 2008; Kent *et al.*, 2012; Sferruzzi-Perri *et al.*, 2016). In contrast, over-expressing  
259 the *Igf2* gene through activating the normally silent maternal gene copy in the *H19* null, increasing



260 IGF2 availability via *Igf2r* ablation, or deletion of the PI3K signalling inhibitor (*Pten*), results in over-  
261 growth of the fetus and placenta (Leighton *et al.*, 1995; Ludwig *et al.*, 1996; Louvi *et al.*, 1997;  
262 Ripoche *et al.*, 1997; Church *et al.*, 2012). Deletion of the *Igf1*, *Igf1r* or *Insr* genes in mice also leads  
263 to fetal growth restriction, but placental weight is unaffected (DeChiara *et al.*, 1990; Baker *et al.*,  
264 1993; Louvi *et al.*, 1997). This suggests that the growth-promoting effect of IGF2 in the mouse  
265 placenta occurs independently of IGF1R and INSR, possibly through an unknown, distinct placental-  
266 specific receptor (XRp) (Louvi *et al.*, 1997). However, evidence from *H19* null mutants suggests that  
267 IGF1R could contribute to the control of placental growth in mice as the first exon of the *H19* gene  
268 encodes *miR-675* which targets *Igf1r* for reduced expression (Keniry *et al.*, 2012). Overgrowth of the  
269 *H19* null placenta (Leighton *et al.*, 1995; Esquiliano *et al.*, 2009; Angiolini *et al.*, 2011; Church *et al.*,  
270 2012), is thus thought to be due to biallelic *Igf2* via imprinting mechanisms, as well as, enhanced  
271 *Igf1r* expression through loss of *miR-675* (Keniry *et al.*, 2012). Taken together, these data highlight  
272 the importance and complexity of the IGF system in controlling conceptus growth in mice.

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274 Genetic manipulations of *Igf2*, *Igf2r* and the downstream signalling pathways also affect the  
275 morphology of the placenta (Table 2). For instance, loss of *Igf2* (complete and *Igf2P0* null), *Pik3ca*,  
276 *Pkba* or *Erk2* gene expression causes defective Lz formation. In particular, Lz volume/thickness,  
277 exchange surface area and vascularisation are all reduced and the interhaemal barrier to diffusion of  
278 gases like oxygen is greater in the placenta of all these mutants (Table 2). In contrast, in the *H19* null,  
279 the Lz surface area is increased in line with the placentomegally observed (Angiolini *et al.*, 2011). In  
280 addition, IGF2 affects the formation of endocrine cells in the placenta. In particular, loss or gain of  
281 *Igf2* or the PI3K-AKT signalling pathway causes a disproportionate decrease or expansion of the  
282 glycogen cells in the Jz, whereas *Igf1r* or *Insr* nulls show no changes in Jz glycogen cell abundance  
283 (Table 2). Collectively, the available data suggest that IGF2 acts via both the PI3K/AKT and MAPK  
284 pathways to attain normal placental weight and Lz structure, and through PI3K/AKT signalling to  
285 drive placental glycogen cell formation in mice (Figure 2A).

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287 Placental function also changes when the IGF system is genetically modified in mice (Table 2). The  
288 passive permeability of the placenta to hydrophilic nutrients/solutes is reduced in the complete *Igf2*  
289 null, placental-specific *Igf2P0* null and *H19* null (Constancia *et al.*, 2002; Sibley *et al.*, 2004; Coan *et*  
290 *al.*, 2008b; Angiolini *et al.*, 2011). The complete *Igf2* null placenta transports less neutral amino acid  
291 (methyl amino-isobutyric acid, MeAIB) via the System A transporters in association with reduced  
292 *SNAT2/Slc38a2* expression (Constancia *et al.*, 2005). There is also reduced abundance of System X<sub>AG</sub><sup>-</sup>  
293 and System Y<sup>+</sup> transporters, responsible for placental transfer of cationic and anionic amino acids, in

294 the complete *Igf2* and the *Igf1r* null (Matthews *et al.*, 1999). In contrast, the *Igf2P0* null placenta  
295 transports more neutral amino acids via System A, as well as, more glucose and calcium in late  
296 gestation (Table 2). Up-regulation of placental transport capacity is associated with increased  
297 expression of *SNAT4/Slc38a4* and *GLUT3/Slc2a3* by the *Igf2P0* deficient placenta. In contrast to *Igf2*,  
298 there is little or no information on the capacity of the *Igf1* or *Insr* null placenta to supply nutrients to  
299 the fetus. In the complete *Igf2* null, placental and fetal growth restriction occurs concurrently and  
300 becomes evident in mid-gestation [Table 2; (Baker *et al.*, 1993; Constancia *et al.*, 2005)]. In the  
301 *Igf2P0* null, placental weight is reduced at a similar time in gestation, but fetal growth only becomes  
302 restricted much closer to term and to a lesser extent than in the complete *Igf2* null (Baker *et al.*,  
303 1993; Constancia *et al.*, 2002; Constancia *et al.*, 2005). Liposome-mediated targeting of IGF2 to the  
304 placenta has recently been shown to increase the weight of *Igf2P0* null mouse fetuses near term  
305 [Table 1; (King *et al.*, 2016)]. Collectively, these findings suggest that the *Igf2P0* null placenta  
306 compensates for its defective development and compromised permeability by adaptively up-  
307 regulating its nutrient transport systems and thereby, minimises the degree of fetal growth  
308 restriction, relative to the complete *Igf2* null. The *Pik3ca* heterozygote deficient placenta also  
309 transfers glucose and amino acids via System A transporters with increased efficiency in  
310 compensation for its impaired development, which is associated a less severe reduction in fetal  
311 weight close to term than earlier in gestation (Sferruzzi-Perri *et al.*, 2016). Moreover, the naturally  
312 small placenta that supports more fetal mass per gram shows increased expression of *Igf2P0*  
313 coupled with a preservation of Lz growth and with increased placental System A transport capacity  
314 and *SNAT2/Slc38a2* abundance compared to the large placenta in the litter (Coan *et al.*, 2008a). In  
315 contrast, the over-grown *H19* null placenta shows diminished neutral amino acid and glucose  
316 transport which is thought to limit fetal over-growth and avoid an excessive drain of maternal  
317 resources into the fetus (Angiolini *et al.*, 2011). Thus, IGF2 in the placenta is important for fine-  
318 tuning nutrient supply to the fetus (Figure 2B).

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320 In addition to effects on placental transport, the *Igf2* gene may also affect the endocrine function of  
321 the placenta with consequences for maternal physiology during pregnancy. Evidence for this stems  
322 from associations between altered placental Jz formation in *H19* and *Igf2P0* null mutants and raised  
323 circulating glucose, insulin and/or corticosterone in phenotypically wild-type dams (Petry *et al.*,  
324 2010; Sferruzzi-Perri *et al.*, 2011). Thus, IGF2 has an important role in nutrient allocation to the  
325 fetus. By regulating placental phenotype, it balances the fetal genetic drive for growth with the  
326 maternal ability to supply the required resources, thereby optimising both offspring and maternal  
327 fitness.

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## **IGFS AS ENVIRONMENTAL SIGNALS IN REGULATING PLACENTAL RESOURCE ALLOCATION TO FETAL GROWTH**

IGFs may also play an important role in changing placental resource allocation to the fetus in environmentally-challenged pregnancies. As *Igf1* expression is relatively low in the placenta, studies have largely focussed on placental expression of *Igf2* and activation of its signalling pathways (Table 3). However, since the signalling pathways are responsive to both IGFs, the placenta can also respond to changes in circulating IGF1 and IGF2 induced by nutritional or other environmental cues.

### **Maternal nutrition**

#### ***Undernutrition***

In mice, guinea pigs and baboons, undernutrition restricts placental growth in association with a decrease in the expression of *Igf2* and/or signalling via the PI3K/AKT and MAPK pathways (Table 3). There are also reductions in placental vascularisation, exchange surface area, Jz volume and glycogen cell abundance and/or a greater barrier to diffusion with maternal undernutrition in mice and guinea pigs; morphological parameters that were altered similarly by a genetic deficiency in *Igf2*, *Pik3ca*, *Pkb* and *Erk2* (Tables 2 and 3). Together, these studies suggest that decreases in IGF2 expression and signalling within the placenta could underlie the growth and morphological defects observed with maternal undernutrition in these species. In larger animals, the expression of *Igf2* and its signalling machinery reduces, is unchanged or even increases in response to undernutrition (Table 3). For instance, signalling via MAPK and PI3K/AKT in the placenta is up-regulated in nutrient-restricted ewes and hiefers (Zhu *et al.*, 2007a; Zhu *et al.*, 2007b; Ma *et al.*, 2011). In these models, changes in signalling relate to a normalisation of placental weight or an increase in placental cotyledon vascularity. They also correlate with a maintenance or restoration of fetal weight in later gestation, despite an exposure to undernutrition. In ewes of a moderate condition, which have the smallest placentas supporting more mass of fetus per gram, placental expression of *Igf2* is greatest (Osgerby *et al.*, 2003). These studies therefore suggest that in larger species, there is morphological adaptation of the placental to an adverse maternal nutritional state through increasing *Igf2* and growth signalling locally.

361 In the undernourished sheep placenta with increased PI3K/AKT and MAPK signalling, the expression  
362 of glucose and fatty acid transporters is also increased (Ma *et al.*, 2011). However, in  
363 undernourished baboons, diminished *Igf2* expression and signalling in the placenta accompanies  
364 reductions in Systems A and L amino acid transporter capacity and glucose transporter gene  
365 expression (Pantham *et al.*, 2015; Pantham *et al.*, 2016). Taken together, these studies suggest that  
366 IGF2 and the PI3K/AKT and MAPK signalling pathways could also mediate changes in placental  
367 transport function during undernutrition. In mice, despite a 20% reduction in maternal food intake  
368 and placental growth restriction earlier in gestation, fetal weight is normal until just prior to term  
369 (Coan *et al.*, 2010). This maintenance of fetal growth relates to an initial preservation of Lz  
370 development in earlier gestation and an adaptive up-regulation of System A amino acid transporter  
371 capacity and *SNAT2/Slc38a2* expression near term, by the growth restricted undernourished  
372 placenta. However, in mice lacking the placental-specific *Igf2* isoform (*Igf2P0*) these adaptations to  
373 maternal undernutrition fail to occur. The development of the placental exchange region is  
374 compromised earlier in gestation, there is no up-regulation of amino acid transport or  
375 *SNAT2/Slc38a2* expression and reduced *SNAT4/Slc38a4* abundance near term in *Igf2P0* null  
376 placentas compared to wildtype in undernourished mice (Sferruzzi-Perri *et al.*, 2011). As a result,  
377 fetal growth is restricted earlier in gestation and more adversely affected near term by  
378 undernutrition, in *Igf2P0* nulls. The *Igf2P0* transcript is, therefore, a major determinant of the  
379 environmental modification of placental phenotype with undernutrition in mice. The expression of  
380 genes involved in glucose, neutral amino acid and fatty acid transport, as well as, the IGF signalling  
381 pathways in the human placenta are modified by the diet and physical activity of the mother during  
382 pregnancy (Brett *et al.*, 2015). Thus, the IGF system may also be important for modifying resource  
383 capacity of the human placenta in response to changes in the maternal environment.

384

385

### 386 ***Low-protein diets***

387 During rodent pregnancy, consumption of an iso-caloric low protein diet has inconsistent impacts  
388 on both placental weight and placental *Igf2* expression [Table 3 and (Sferruzzi-Perri & Camm, 2016)].  
389 However, the nature of the specific effect appears to depend on the degree of protein deprivation,  
390 stage of pregnancy studied and sex of the conceptus (Jansson *et al.*, 2006; Coan *et al.*, 2011; Nusken  
391 *et al.*, 2011; Gao *et al.*, 2012a). Despite the contrasting results, placental *Igf2* expression seems to  
392 track positively with the weight of the placenta in mice and rats (Coan *et al.*, 2011; Nusken *et al.*,  
393 2011; Gao *et al.*, 2012a). For instance, in pregnant mice, low protein diets cause placentomegaly and  
394 the degree of placental weight increase relates to the level of *Igf2* up-regulation at first appearance

395 of growth enhancement (Coan *et al.*, 2011). The variation in placental growth and *Igf2* expression  
396 observed in different models of protein deficiency could be caused by the content and source of  
397 carbohydrate used to maintain calorie intake. Nevertheless, taken together, these findings suggest  
398 that at least part of the changes in placental growth seen with protein deprivation could be  
399 mediated through local changes in *Igf2*.

400

401 There are also changes in placental transport capacity with gestational protein malnutrition. For  
402 instance, in response to a diet with 8% protein, the mouse placenta adaptively transports more  
403 glucose to the fetus on day 16 of pregnancy (Coan *et al.*, 2011). This up-regulation occurs when  
404 placental *Igf2* expression is also increased and when fetal growth is maintained despite maternal  
405 protein deprivation (Coan *et al.*, 2011). A few days later however, glucose transport is unchanged,  
406 System A amino acid transporter abundance is reduced and *Igf2* expression no longer increased in  
407 the placenta by a low protein diet, and fetal growth restriction ensues (Coan *et al.*, 2011). These data  
408 suggest that placental *Igf2* may be important for adapting nutrient supply to the fetus in response to  
409 maternal protein malnutrition in mice. However, there is evidence that pathways downstream of  
410 *Igf2* may also be important. For instance, the mechanistic target of rapamycin (mTORC1) mediates  
411 the mitogenic and metabolic actions of IGFs (Jansson *et al.*, 2012b). In rats, protein deprivation  
412 reduces mTORC1 signalling, Systems A and L amino acid transport and *SNAT2/Slc38a2*, *LAT1/Slc7a5*  
413 and *LAT2/Slc7a8* expression by the placenta, prior to the appearance of placental and fetal growth  
414 restriction (Jansson *et al.*, 2006; Rosario *et al.*, 2011). These findings suggest that down-regulation of  
415 signalling pathways like mTORC1 and amino acid transporters in the placenta could link maternal  
416 protein restriction to decreases in fetal growth. The availability of protein and specific amino acids  
417 during pre-implantation rodent development is linked to alterations in the expression of genes  
418 within the *H19-Igf2* locus, mTORC1 signalling and trophoblast cell formation and differentiation with  
419 consequences for feto-placental phenotype in late gestation (Kwong *et al.*, 2006; Van Winkle *et al.*,  
420 2006; Eckert *et al.*, 2012; Watkins *et al.*, 2015). Thus, changes in *Igf2* expression and its signalling  
421 pathways could be responsive to the availability of nutrients from the earliest stages of  
422 development.

423

424

#### 425 ***Diets with excess sugar and/or fat***

426 The expression of *Igf2* and its signalling pathways in the placenta are inconsistently altered by diets  
427 with excess sugar and/or fat (Table 3). Weight of the conceptus may also be reduced, increased or  
428 unchanged, depending on the level of fat in the diet, the amount of simple sugars consumed and the

429 timing of the dietary manipulation [Table 3 and reviewed in (Sferruzzi-Perri & Camm, 2016)]. Part of  
430 these variations in *Igf2* expression and conceptus growth could be due to the differences in protein  
431 and micronutrient intake, as species like mice and rats control their calorie intake tightly (Keesey &  
432 Hirvonen, 1997). In mice fed a diet containing 2.5-times the fat of the controls, placental weight is  
433 reduced in early pregnancy in association with decreases in the expression of *Igf2* and signalling  
434 machinery, including *Mtor* (Sasson *et al.*, 2015). These placental changes accompanied reductions in  
435 the expression of System A amino acid transporter, *SNAT1/Slc38a1*, glucose transporter  
436 *GLUT1/Slc2a1* and/or fatty acid translocase, *CD36* depending on the length of high fat feeding and  
437 whether the diet was eaten before pregnancy (Sasson *et al.*, 2015). In over-nourished ewes,  
438 placental weight is reduced in mid-gestation in association with decreased activity of the IGF  
439 signalling pathway (including activation of IRS1 and mTORC1) and changes in vessel size and density  
440 in the placenta (Zhu *et al.*, 2009; Ma *et al.*, 2010). However, fetal weight is increased along with fatty  
441 acid transporters and translocases in the placenta, suggesting that alternative signalling pathways  
442 may be activated to adapt placental nutrient supply to the fetus in ewes with excess food intake  
443 (Zhu *et al.*, 2010; Tuersunjiang *et al.*, 2013).

444

445 In other studies, increases in the placental IGF system are coupled with improved placental resource  
446 allocation to the fetus in dams fed obesogenic diets (King *et al.*, 2013; Sferruzzi-Perri *et al.*, 2013;  
447 Diaz *et al.*, 2015; Rosario *et al.*, 2015; Rosario *et al.*, 2016). For instance, in mice, consumption of a  
448 high sugar and fat diet from day 1 of pregnancy initially causes conceptus growth restriction and  
449 morphological defects in the placental Lz. However, fetal weight normalises by term, despite the  
450 persistence of placental growth and morphological defects through adaptive up-regulation of  
451 glucose and neutral amino acid transport to the fetus by the placenta (Sferruzzi-Perri *et al.*, 2013).  
452 Up-regulation of transport capacity relates to increased expression of *GLUT3/Slc2a3* and  
453 *SNAT2/Slc38a2*, as well as, elevated expression of the placental-specific *Igf2* isoform and PI3K/AKT  
454 signalling in the placenta in dams fed a diet with excess sugar and fat. Obesogenic diets fed from  
455 before pregnancy also increase placental nutrient transporter capacity (glucose, Systems A and L  
456 amino acid and fatty acids) in line with greater *Igf2* or PI3K/AKT and mTORC1 signalling, however  
457 responses varied with the precise composition of the diet and possibly, fetal sex (King *et al.*, 2013;  
458 Aye *et al.*, 2015; Diaz *et al.*, 2015; Rosario *et al.*, 2015; Rosario *et al.*, 2016). The expression of IGF  
459 signalling machinery (receptors, AKT, mTORC1) and nutrient transporters is also altered in the  
460 placenta from obese women, however, the specific nature of these changes appears to depend on  
461 the level of maternal body fat mass, gestational weight gain and whether macrosomia is observed  
462 (Jansson *et al.*, 2012a; Brett *et al.*, 2016; Martino *et al.*, 2016). Taken together, these findings suggest

463 that obesity and obesogenic diets alter placental phenotype in association with changes in placental  
464 *Igf2* system and fetal growth.

465

466

#### 467 **Maternal hypoxia**

468 In mice, hypoxia typically reduces fetal growth in a severity-dependent manner without a change in  
469 placental weight [Table 3 and reviewed in (Sferruzzi-Perri & Camm, 2016)]. However, if the hypoxic  
470 challenge commences early in pregnancy, placentomegaly is observed in associated with greater  
471 maternal blood spaces and activation of the PI3K/AKT and mTORC1 signalling pathways in the  
472 placenta (Matheson *et al.*, 2015). Even though placental weight may not be altered when maternal  
473 hypoxia commenced later in pregnancy, placental expression of the IGF system and capacity to  
474 supply resources to the fetus is altered (Table 3). In particular, placental expression of IGF receptors,  
475 INSR and PI3K isoforms is decreased in response to five days of 13%-10% maternal hypoxia in late  
476 mouse gestation, and in 10% hypoxia this effect is due to reductions in maternal food intake (Cuffe  
477 *et al.*, 2014; Higgins *et al.*, 2015). However, expression of *Igf2*, *Igf2P0* and activated AKT increases  
478 with 13% hypoxia, but is unchanged or even decreased in response to 5 days of 12-10% hypoxia near  
479 term (Cuffe *et al.*, 2014; Higgins *et al.*, 2015). In the 13% hypoxic mouse placenta showing increases  
480 in IGF2 expression and signalling, there are beneficial changes in Lz structure including improved  
481 vascularisation, maternal blood spaces and a thinner diffusion barrier to exchange; changes that  
482 would optimise oxygen delivery to the fetus near term (Higgins *et al.*, 2015; Matheson *et al.*, 2015).  
483 There is also greater placental glucose uptake and transport and maintained delivery of neutral  
484 amino acids to the fetus when 13% hypoxia occurs in the last third of pregnancy (Higgins *et al.*,  
485 2015). In contrast, in the 12-10% hypoxic placenta with unchanged or decreased expression of the  
486 IGF2 system, the morphology of the placental Lz is compromised, with reductions in maternal blood  
487 spaces and surface area and a greater barrier to diffusion; changes that would further limit fetal  
488 oxygen supply in hypoxic dams (Cuffe *et al.*, 2014; Higgins *et al.*, 2015). Moreover, placental glucose  
489 uptake and transport capacity is not up-regulated or even reduced (less *GLUT1/Slc2a1*) and delivery  
490 of neutral amino acids diminished, in dams exposed to 12-10% hypoxia, depending on whether food  
491 intake is reduced and the sex of the fetus (Cuffe *et al.*, 2014; Higgins *et al.*, 2015). In culture, 1%  
492 hypoxia reduces the outgrowth of mouse ectoplacental cone trophoblast in association with  
493 diminished *Igf2* expression (Pringle *et al.*, 2007). Hypoxia (1% oxygen) also diminishes the expression  
494 of PI3K/AKT and mTORC1 signalling in human trophoblast cell lines (Yung *et al.*, 2012a) and  
495 modulates IGF1 and IGF2 signalling in early pregnancy placental mesenchymal stem cells (Youssef *et*  
496 *al.*, 2014; Youssef & Han, 2016). Placental expression of the PI3K/AKT and mTORC1 signalling

497 pathways and *GLUT1/Slc2a1* expression are decreased in women at 3100m above sea level who  
498 deliver growth-restricted babies (Zamudio *et al.*, 2006; Yung *et al.*, 2012a). In addition, inducing  
499 endoplasmic stress in the mouse placental Jz genetically is associated with defects in PI3K/AKT and  
500 mTORC1 signalling, altered IGF2 glycosylation and bioactivity, and with feto-placental growth  
501 restriction (Yung *et al.*, 2012b). Taken together, these findings suggest that activating IGF2 and/or  
502 PI3K/AKT signalling in the placenta may be critical for adapting placental resource allocation to the  
503 fetus during hypoxia in late pregnancy. They also suggest that the placenta may integrate signals of  
504 oxygen and nutrient availability through the IGF2 system to adapt its phenotype and optimize  
505 maternal resource supply to fetal growth. Indeed, the mouse *Igf2* gene harbours a hypoxia-  
506 responsive element in its promoter (Feldser *et al.*, 1999), as well as CHORE motifs, which bind the  
507 glucose-responsive transcription factor, MLX (Hunt *et al.*, 2015). Therefore, the availability of oxygen  
508 and nutrients *in utero* could have direct effects on placental *Igf2*. Nutritional and hypoxic challenges  
509 alter the concentration of hormones like the glucocorticoid stress hormone and insulin, in the  
510 maternal circulation (Sferruzzi-Perri *et al.*, 2011; Cuffe *et al.*, 2014). Thus, changes in placental  
511 phenotype may reflect alterations in the metabolic and endocrine state of the mother.

512

### 513 **Maternal endocrine challenges**

514 Endocrine challenges can affect maternal metabolism and utilisation of nutrients and thus the  
515 partitioning of resource to the conceptus in pregnancy (Vaughan *et al.*, 2011). Administering  
516 corticosterone or the synthetic glucocorticoid, dexamethasone to rodents for 3-7 days reduces fetal  
517 and placental weights during gestation [Table 3 and (Vaughan *et al.*, 2011)]. In mice, corticosterone  
518 decreases AKT and mTORC1 activation in association with reductions in feto-placental weight, Lz  
519 vascularisation and glucose and System A amino acid transporter capacity, however the specific  
520 nature of these effects depend on when in pregnancy the over-exposure occurs (Table 3).  
521 Administering the synthetic glucocorticoid, dexamethasone reduces the expression of MAPK and  
522 weight of placenta in female, but not male conceptuses and there is no change in glucose and  
523 *SNAT/Slc38a* amino acid transporters irrespective of fetal sex (Cuffe *et al.*, 2011). In mice, placental  
524 *Igf2* expression is unaffected by maternal administration of corticosterone and dexamethasone even  
525 though the conceptus may be growth-restricted (Cuffe *et al.*, 2011; Vaughan *et al.*, 2012; Vaughan *et al.*,  
526 2015). Whereas restrain stress increases placental *Igf2* but does not alter offspring weight in  
527 mice (Pankevich *et al.*, 2009). In rats, dexamethasone decreases placental *Igf2* and the level of  
528 activated AKT, particularly in the endocrine Jz (Ain *et al.*, 2005). In dexamethasone-treated rats,  
529 there are reductions in the expression of prolactin-related family genes by the Jz in late gestation,  
530 which may influence the maternal adaptations to pregnancy and, thus, alter the fetal supply of



531 nutrients indirectly (Ain *et al.*, 2005). The expression of *Igf2* by the term human placenta is also  
532 altered in women with elevated plasma cortisol during pregnancy due to emotional distress (Mina *et*  
533 *al.*, 2015). Glucocorticoid response elements have been identified in the human *Igf1* gene promoter  
534 (He *et al.*, 2016) however, very little is known about whether glucocorticoids could have direct  
535 effects on placental *Igf2* expression. Collectively these findings suggest that reductions in placental  
536 *Igf2* system and the functional phenotype of the placenta could link elevated maternal  
537 glucocorticoids to decreases in fetal growth.

538

539 In rats, pre-existing maternal diabetes also alters the expression of the IGF system in the placenta, as  
540 well as, materno-fetal resource allocation, however, the direction of change depends on how long  
541 the dam was insulin deficient/dysglycemic. For instance, *Igf* expression and IGF1R activation are  
542 elevated in association with greater glycerol and free fatty acid transfer by the placenta and an  
543 increase in fetal weight by 13% in rats that are diabetic from neonatal life (White *et al.*, 2015).  
544 Placental lipid transport capacity is also increased in rat dams that are diabetic for 1 week prior to  
545 pregnancy (increase in placental lipoprotein lipase), however, the expression of *Igf2* and the IGF  
546 signalling machinery is decreased and the abundance of *GLUT1/Slc2a1* reduced in association with a  
547 more minor increase in fetal weight (by 5%) (Cisse *et al.*, 2013). Genetically-inducing maternal insulin  
548 insensitivity by a global heterozygous deficiency in PI3K-p110 $\alpha$  signalling capacity in the mouse dam  
549 is associated with improved placental Lz development (larger surface area and thinner barrier to  
550 diffusion), but reduced glucose transport and expression of nutrient (*GLUT1/Slc2a1*, *SNAT1/Slc38a1*,  
551 *SNAT2/Slc38a2*) and prolactin-related family genes near term (Sferruzzi-Perri *et al.*, 2016). However,  
552 the specific nature of these placental changes depended on whether the conceptus itself was  
553 heterozygous for the PI3K-p110 $\alpha$  deficiency (Sferruzzi-Perri *et al.*, 2016). Moreover, there is no  
554 effect of maternal heterozygous deficiency in PI3K-p110 $\alpha$  signalling on fetal weight in this model,  
555 irrespective of fetal genotype (Sferruzzi-Perri *et al.*, 2016). Taken together, these studies suggest  
556 that the IGF2/PI3K-p110 $\alpha$  system plays an important role in modulating fetal nutrition and growth in  
557 response to maternal insulin deficiency and/or insensitivity, by acting at the level of placental  
558 transport phenotype.

559

560

### 561 ***Other environmental challenges affecting conceptus growth***

562 The expression of *Igfs*, receptors and signalling machinery in the placenta also changes in response  
563 insults that affect the placental capacity to supply the fetus with nutrients. Such insults include  
564 reduced utero-placental blood flow, heat stress and alcohol consumption (Table 3). Reducing both

565 maternal oxygen and nutrient supply to the conceptus using uterine artery ligation in mice, rats and  
566 guinea pigs, or placental embolism in sheep, reduces placental expression of components of the IGF  
567 system in association with defects in placental Lz structure and in transporter expression and activity  
568 of the glucose and Systems A and L amino acid transporters (Table 3). The extent of these changes  
569 however, depends on timing of the insult in the pregnancy. In sheep, removal of uterine caruncles  
570 prior to pregnancy is associated with increased placental *Igf2* expression and an adaptive increase in  
571 placentome size, trophoblast and maternal capillary volume and surface area, although total  
572 placental mass and fetal weight, are reduced [Table 3; (Zhang *et al.*, 2016b)]. Acute exogenous IGF1  
573 does not alter nutrient metabolism by the embolised sheep placenta [Table 1; (Jensen *et al.*, 1999)].  
574 However, several doses of intra-amniotic IGF1 increases glucose and Systems A and L amino acid  
575 transporter expression by the embolised placenta and improves fetoplacental growth *in vivo* [Table  
576 1; (Eremia *et al.*, 2007; Wali *et al.*, 2012)]. Moreover, in mice with uterine artery ligation, targeting of  
577 IGF1 to the placenta using a nanoparticle or adenoviral-mediated approach increases the abundance  
578 of glucose and Systems A and L amino acid transporters in the placenta, placental width and fetal  
579 growth [Table 1; (Jones *et al.*, 2013; Jones *et al.*, 2014; Abd Allah *et al.*, 2015)]. These findings  
580 highlight the therapeutic potential of IGFs for improving the capacity of the placenta to supply  
581 nutrients to the fetus in compromised pregnancies.

582

583

584 In ewes, heat stress reduces placental growth and glucose transport capacity, as well as, alters the  
585 expression of IGF1 and IGF2, and AKT, mTORC1 and MAPK signalling pathways during gestation  
586 (Table 3). In rats, alcohol consumption during the peri-conceptual period leads to late gestational  
587 fetal growth restriction but no change in placental weight (Gardebjer *et al.*, 2014). However, Lz  
588 development and *Igf1*, *Igf1r* and *SNAT2/Slc38a2* expression is decreased, but Jz glycogen cell  
589 formation, *Igf2* and GLUT1/Slc2a1 may be increased in response to peri-conceptual alcohol  
590 exposure near term (Gardebjer *et al.*, 2014). This suggests there can be programmed changes in the  
591 conceptus leading to changes in the IGF system and the structural and functional phenotype of the  
592 placenta that link the maternal environment from the earliest stages of pregnancy to fetal growth  
593 near term.

594

## 595 **CONCLUSIONS AND PERSPECTIVES**

596 Thus, IGFs are important regulators of placental resource allocation to fetal growth both  
597 developmentally and in response to environmental manipulations known to program the ill health of  
598 offspring. They increase placental morphogenesis, substrate transport and hormone secretion,

599 which, in turn promotes fetal growth either directly via the supply of nutrients and oxygen or  
600 indirectly via the maternal metabolic adaptation to pregnancy and the availability of nutrients for  
601 transplacental transport. In response to environmental challenges, the IGFs (particularly IGF2) and  
602 their signalling pathways change in line with the alterations in placental structure and function, and  
603 thereby, link changes in the maternal environment to fetal substrate supply and growth during  
604 pregnancy with implications for developmental programming. The environmentally-induced changes  
605 in the IGF system and placental phenotype may be beneficial (obesogenic diets, moderate hypoxia)  
606 or detrimental (eg. severe oxygen and nutrient deprivation and glucocorticoid excess) to resource  
607 allocation to the fetus depending on the type, severity and timing of the challenge during pregnancy  
608 (Figure 3). The beneficial effects of IGF treatments on placental phenotype show promising  
609 therapeutic potential for improving fetal growth in situations in which placental growth is impaired  
610 without major maternal compromise, particularly when the treatment with IGF1 or IGF2 is targeted  
611 directly to the placenta. However, efforts to understand the regulation of endogenous placental IGF  
612 expression may also be fruitful, particularly in the case of *Igf2* which appears to be most important  
613 for mediating adaptive responses locally in mice. These findings are important in the context of  
614 human pregnancy as dysregulated expression of the IGFs and signalling components are often  
615 reported in the human placenta associated with abnormal fetal growth (Abu-Amero *et al.*, 1998;  
616 Sheikh *et al.*, 2001; Gratton *et al.*, 2002; Gurel *et al.*, 2003; Laviola *et al.*, 2005; Scioscia *et al.*, 2006;  
617 Street *et al.*, 2006; Trollmann *et al.*, 2007; Akram *et al.*, 2008; Yung *et al.*, 2008; Colomiere *et al.*,  
618 2009; Borzsonyi *et al.*, 2011; Street *et al.*, 2011; Demendi *et al.*, 2012; Jansson *et al.*, 2012a; Iniguez  
619 *et al.*, 2014; Nawathe *et al.*, 2016; Zhang *et al.*, 2016a). However, it is important to note, that several  
620 causes of environmental, maternal, and fetal origin, can lead to changes in placental phenotype and  
621 fetal growth in humans (Gaccioli & Lager, 2016). Thus studies of animal models showing alterations  
622 in the expression of IGFs and their signalling pathways provides insight but further information is  
623 required on the natural conditions of variable placental phenotype among humans.

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1488

1489

1490 **Abstract figure. The proposed actions of IGFs on placental resource allocation to drive fetal**  
1491 **growth.** Note that changes in placental IGFs and resource allocation depend on the timing and  
1492 severity of the environmental insult.

1493

1494 **Figure 1. Impact of exogenous IGFs on the placenta. A)** The effect of exogenous IGFs on placental  
1495 human trophoblast *in vitro*. Proposed signalling pathways mediating the actions of IGFs shown. **B)**  
1496 The effect of exogenous maternal IGFs on the mouse, rat and/or guinea pig placenta *in vivo*. Dashed  
1497 lines indicate a potential interaction (A) or impact (B) of IGF1. IGF = insulin-like growth factor, IGF1R  
1498 = type 1 IGF receptor, IGF2R = type 2 IGF receptor, INSR = insulin receptor, Jz = junctional zone, Lz =  
1499 labyrinthine zone, MAPK = mitogen-activated protein kinase, PI3K = phosphoinositol 3-kinase.

1500

1501 **Figure 2. The effect of genetically manipulating IGF2 expression or signalling on placental**  
1502 **phenotype in mice. A)** shows the effect of complete loss of IGF2 and **B)** shows the effect of partial  
1503 loss of IGF2, either by deleting the placental-exclusive isoform, *Igf2P0* or through a constitutive  
1504 heterozygous deficiency of PI3K-p110 $\alpha$ . Dashed line indicates a potential interaction of IGF2 with  
1505 receptor. Line with a round head indicates parameters reduced by loss of IGF2 signalling. Loss of  
1506 IGF2 signalling leads to reductions in placental development and transport function (A). Partial loss of  
1507 IGF2 signalling also leads to reductions in placental development, but is associated with adaptive up-  
1508 regulation in transport function (B). AA = amino acids, IGF = insulin-like growth factor, IGF1R = type 1  
1509 IGF receptor, Lz = labyrinthine zone, MAPK = mitogen-activated protein kinase, PI3K =  
1510 phosphoinositol 3-kinase, XRp = unknown placental-specific IGF receptor.

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1512 **Figure 3. The effect of different environmental manipulations on the placental IGF system and**  
1513 **resource allocation phenotype in the mouse. A)** shows manipulations which down-regulate IGF2

1514 signalling. **B)** shows manipulations which up-regulate IGF2 signalling. AKT = protein kinase B, IGF =  
1515 insulin-like growth factor, Lz = labyrinthine zone. \* Note *Igf2P0* is required for the placenta to up-  
1516 regulate amino acid transport to the fetus in response to maternal undernutrition.

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1532 **Table 1.** The impact of exogenous IGF1 or IGF2 on the placental phenotype and fetal outcome (where available)

IGF	System	Species	Treatment	Study	Placental size and morphology	Placental function	Fetal weight	References
IGF1	<i>In vitro</i>	Mouse	Primary ectoplacental cone trophoblast	First trimester	↑ proliferation and migration			(Kanai-Azuma <i>et al.</i> , 1993)
		Pig	Primary trophoblast cells	First trimester	↑ proliferation and migration			(Jeong <i>et al.</i> , 2014)
		Human	1 <sup>st</sup> trimester primary trophoblast	First trimester	↑ invasion via INSR and IGF1R activation of Akt			(Mayama <i>et al.</i> , 2013)
		Human	1 <sup>st</sup> trimester placental explant	First trimester	↑ proliferation and syncytial formation via IGF1R-mediated MAPK signalling, ↓ apoptosis via IGF1R-mediated PI3K signalling			(Forbes <i>et al.</i> , 2008; Forbes <i>et al.</i> , 2015)
		Human	1 <sup>st</sup> trimester placental trophoblast	First trimester	↑ proliferation, migration			(Hashimoto <i>et al.</i> , 2010)
		Human	1 <sup>st</sup> trimester placental explant	First trimester	↑ proliferation			(Forbes <i>et al.</i> , 2009; Forbes <i>et al.</i> , 2015)
		Human	1 <sup>st</sup> trimester placental explant	First trimester	↑ migration			(Lacey <i>et al.</i> , 2002)

		Human	1 <sup>st</sup> trimester placental explant	First trimester	↑ proliferation	↑hCG, hPL		(Maruo <i>et al.</i> , 1995)
		Human	1 <sup>st</sup> trimester trophoblast	First trimester		↑System A amino acid and glucose uptake		(Kniss <i>et al.</i> , 1994)
		Human	1 <sup>st</sup> trimester primary placental fibroblasts	First trimester	↑proliferation, invasion, ↓ apoptosis			(Miller <i>et al.</i> , 2005)  *Ad-IGF-I
		Human	BeWo syncytial cell line		↑ proliferation, invasion, ↓ apoptosis	↑ System A and System L amino acid transporter activity, Snat1, Snat2, Lat1, 4F2hc, GLUT1, GLUT3 and GLUT8, ↓Lat2		(Jones <i>et al.</i> , 2013; Jones <i>et al.</i> , 2014)  *Ad-hIGF-I
		Human	BeWo			↔ pGH		(Zeck <i>et al.</i> , 2008)
		Human	JEG-3 choriocarcinoma cell line		↑proliferation, ↓apoptosis	↑ P4, hCG secretion		(Rak-Mardyla & Gregoraszcuk, 2010)
		Human	JEG-3		↑invasion via induction of adhesion and migration through IGF1R-PI3K and MAPK signalling			(Diaz <i>et al.</i> , 2007)
		Human	BeWo			↑ System A amino acid transporter activity via PI3K signalling, ↔ Snat1 or		(Fang <i>et al.</i> , 2006)

						Snat2		
		Human	BeWo, term explants and term perfused human placenta			↑ glucose transport, GLUT1 membrane abundance		(Baumann <i>et al.</i> , 2014)
		human	Term human placenta	Term		↓ LPL activity in		(Magnusson-Olsson <i>et al.</i> , 2006)
		Human	Term trophoblast	Term		↑ System A amino acid uptake		(Bloxam <i>et al.</i> , 1994; Karl, 1995; Yu <i>et al.</i> , 1998)
		Human	Term trophoblast and cell lines	Term	↑ syncytialisation			(Bhaumick <i>et al.</i> , 1992; Milio <i>et al.</i> , 1994; Cohran <i>et al.</i> , 1996)
<b>IGF2</b>	<i>In vitro</i>	Mouse	Primary ectoplacental cone trophoblast	First trimester	↑ differentiation into endocrine cells			(Kanai-Azuma <i>et al.</i> , 1993)
		Sheep	Primary trophoblast	First trimester	↑ migration			(Kim <i>et al.</i> , 2008)
		Human	1 <sup>st</sup> trimester HTR8_SVneo cell line	First trimester	↑ migration via Rho GTPases			(Qiu <i>et al.</i> , 2005; Shields <i>et al.</i> , 2007)
		Human	1 <sup>st</sup> trimester HTR8_SVneo cell line	First trimester	↑ migration via signalling through IGF2R involving inhibitory G proteins and the MAPK pathway			(McKinnon <i>et al.</i> , 2001)

		Human	1 <sup>st</sup> trimester primary trophoblast	First trimester	↑ migration/invasion			(Irving & Lala, 1995; Hamilton <i>et al.</i> , 1998)
		Human	1 <sup>st</sup> trimester placental explant	First trimester	↑ trophoblast proliferation and syncytial formation via IGF1R-mediated MAPK signalling, ↓ apoptosis via IGF1R-mediated PI3K signalling			(Forbes <i>et al.</i> , 2008; Forbes <i>et al.</i> , 2009; Forbes <i>et al.</i> , 2015)
		Human	JEG-3 choriocarcinoma cell line		↑ invasion via induction of adhesion and migration through INSR-PI3K and MAPK signalling			(Diaz <i>et al.</i> , 2007)
		Human	SGHPL4 and 1 <sup>st</sup> trimester villous explants	First trimester	↑ proliferation, migration and invasion			(Pollheimer <i>et al.</i> , 2011)
		Human	1 <sup>st</sup> trimester primary placental fibroblasts	First trimester	↑ proliferation and invasion, ↓ apoptosis			(Miller <i>et al.</i> , 2005) *Ad-IGF-II
		Human	1 <sup>st</sup> trimester and term trophoblast	First trimester	↓ apoptosis, ↑ proliferation and survival against TNF-α and IFN-γ-induced apoptosis			(Hills <i>et al.</i> , 2012)
		Human	1 <sup>st</sup> trimester placental	First		↑ glucose uptake		(Kniss <i>et al.</i> , 1994)

			trophoblast	trimester				
		Human	1 <sup>st</sup> trimester placental trophoblast	First trimester		↑ glucose and System A amino acid uptake		(Kniss <i>et al.</i> , 1994)
		Human	1 <sup>st</sup> trimester placental explant	First trimester	↑ migration			(Lacey <i>et al.</i> , 2002)
		Human	In BeWo and term explants		↑ proliferation ↓ apoptosis and necrosis			(Harris <i>et al.</i> , 2011)
IGF1	<i>In vivo</i>	Mouse	D14	D17	↔ weight, ↑ placental cross-sectional area, Lz and fetal and maternal facing areas		↔ weight or viability	(Katz <i>et al.</i> , 2009)  * Ad-hIGF-I
		Mouse uterine artery ligation	D16	D20	↔ weight, ↑ placental thickness		↑27%, ↔ fetal viability	(Abd Ellah <i>et al.</i> , 2015)  * nanoparticle targeted delivery to placenta: PLAC1-hIGF-1
		Mouse uterine artery ligation	D18	D20	ND	↑4F2hc, Lat1, Lat2, GLUT8, GLUT9a/b, ↔ Snat1, Snat2,	ND	(Jones <i>et al.</i> , 2013; Jones <i>et al.</i> , 2014)

						GLUT1		* Ad-hIGF-I
	Guinea pig	D20-37	D40	↔ weight			↑ 6%, ↓ litter size	(Sohlstrom <i>et al.</i> , 2001)
	Guinea pig	D20-38	D35	↑ 17% weight, ↓ placental and Lz area, ↔ Lz, Jz, FC, MBS, Troph Vd	↑ glucose and System A amino acid transfer, Snat2 and prorenin activation, ↓ Igf2, ↔ Glut1, Igf1		↑ 15%	(Sferruzzi-Perri <i>et al.</i> , 2007b; Standen <i>et al.</i> , 2015)
	Guinea pig	D20-38	D62	↔ weight ↔ structure	↑ glucose and System A amino acid transfer		↑ 17% ↑ fetal viability	(Sferruzzi-Perri <i>et al.</i> , 2006; Sferruzzi-Perri <i>et al.</i> , 2007a)
	Guinea pig 30%UN	D20-37	D40	↑ 13% weight			↔	(Sohlstrom <i>et al.</i> , 2001)
	Rabbit  Natural runt in litter	D19	D21	↔ weight			↑ 19%	(Keswani <i>et al.</i> , 2015)  * Ad-hIGF-I

		Sheep	D128, 4hr infusion	D128	ND	↑ glucose transfer and lactate production, ↔ blood flow, urea or glucose transfer	ND	(Liu <i>et al.</i> , 1994)
		Sheep * Fetal infusion	D121-132	D132	↔ weight, ↓ placentome number	↓ glucose and System A amino acid transfer	↔	(Bloomfield <i>et al.</i> , 2002b)
		Sheep * Fetal infusion	D128, 4hr infusion	D128	ND	↓ glucose transfer, lactate uptake and umbilical flow, ↔ urea transfer or serine uptake	ND	(Harding <i>et al.</i> , 1994; Jensen <i>et al.</i> , 1999; Jensen <i>et al.</i> , 2000)
		Sheep Embolised * Fetal infusion	D128, 4hr infusion	D128	ND	↔ glucose or urea transfer, lactate uptake and umbilical flow	ND	(Jensen <i>et al.</i> , 1999)
		Sheep Spontaneous growth restriction * Fetal infusion	D128, 4hr infusion	D128	ND	↔ glucose or urea transfer, lactate uptake and umbilical flow	ND	(Jensen <i>et al.</i> , 1999)
		Sheep Embolised * Intra-	D110, D117, D124	D120-131	↔ but placentas no longer significantly different to untreated controls	↔ glucose uptake, ↑ Glut1, Glut4, Systems y+ and L transporters (Slc7a1 and	↔ weight, ↑ fetal growth rate and fetuses no longer significantly different	(Ereimia <i>et al.</i> , 2007; Wali <i>et al.</i> , 2012)

		amniotic infusion				Slc7a8) ↔ Glut3, Snat4, Slc7a5	to untreated controls	
<b>IGF2</b>	<i>In vivo</i>	Mouse	D14, D16, D18  IGF2 (1mg/kg/day) or iRGD-liposome with IGF2 (0.3mg/kg/day)	D18	↑ weight		↔	(King <i>et al.</i> , 2016)
		Mouse IGF2P0	D14, D16, D18  treatment with iRGD-liposome with IGF2 (0.3mg/kg/day)	D18	↔ weight of IGF2P0 and WT		↑ IGF2P0 but not WT	(King <i>et al.</i> , 2016)
		Rat	D16-22	D22	↔ weight, ↑ Jz		↔	(Van Mieghem <i>et al.</i> , 2009)
		Guinea pig	D20-37	D40	↑ 9% weight		↑ 7%	(Sohlstrom <i>et al.</i> , 2001)
		Guinea pig	D20-38	D35	↔ weight and structure	↔ glucose or System A amino acid transfer, Glut1, Snat2, IGF1 and IGF2, ↑ prorenin activation	↔	(Sferruzzi-Perri <i>et al.</i> , 2007b; Standen <i>et al.</i> , 2015)
		Guinea pig	D20-38	D62	↔ weight, ↑ Lz area, Vd, Vol, SA, ↓ Jz Vd, ↔ BT	↑ glucose transfer, ↔ System A amino acid transfer	↑ 11% weight and ↑ fetal viability	(Sferruzzi-Perri <i>et al.</i> , 2006; Sferruzzi-Perri <i>et al.</i> , 2007a)



		Guinea pig 30%UN	D20-37	D40	↔ weight		↔	(Sohlstrom <i>et al.</i> , 2001)
<b>Leu<sup>-</sup><sub>27</sub></b> <b>IGF-II</b>	<i>In vivo</i>	Mouse	D13-19	D19	↔ weight	↔ System A amino acid transfer, ↓ litter System A amino acid variability	↔ weight, ↓ variability in fetal weight	(Charnock <i>et al.</i> , 2016)
		Mouse eNOS-/-	D13-19	D19	↔ weight		↑	(Charnock <i>et al.</i> , 2016)
		Guinea pig	D20-38	D62	↔ weight, ↑ Lz vd, Troph, MBS Vd and Vol and SA, ↓ Jz area, Vd, Vol, FC Vd, Vol and BT	↑ glucose and System A transfer and prorenin activation	↑ 11%	(Sferruzzi-Perri <i>et al.</i> , 2008)

1533 For *in vivo* studies, exogenous IGF was administered to the mother, unless stated otherwise. Abbreviations: BT=barrier thickness, D=day, FC=fetal capillaries, GLUT=glucose transporter,  
1534 hCG=human chorionic gonadotrophin, hPL=human placental lactogen, IGF1/Igf1=insulin-like growth factor-1, IGF2/Igf2=insulin-like growth factor-2, Jz=junctional zone, LAT=cationic amino  
1535 acid transporter, Lz=labyrinthine zone, MAPK/ERK=mitogen activated kinase, MBS=maternal blood space, ND=not determined; P4=progesterone; PI3K=phosphoinositol 3-kinase,  
1536 pGH=placental growth hormone, Prl=prolactin-related hormone, SA=surface area, SNAT/Slc38a= Sodium-coupled neutral amino acid transporter, vol=volume, vd=volume density.

1537 **Search terms used:** trophoblast, placenta, fetus, insulin-like growth factor, IGF and/or transport

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1539 Table 2. The effect of genetically manipulating IGF abundance and/or signalling on fetoplacental growth in mice

Manipulation	Approach	Placental			Fetal weight	Reference
		size	morphology	function		
<b>Deficiency of IGF and downstream signalling</b>						
Global IGF1 KO	Igf1 <sup>-/-</sup>	D18/19 ↔			D18/19 ↓40%	(Baker <i>et al.</i> , 1993)
Global IGF2 KO	Paternal Igf2 <sup>-</sup>	D15 ↓47%  D18/19 ↓20-30%	D15 ↓ Jz GlyT  D18/19 ↔ Lz and Jz Vd ↓ Jz GlyT	D17 ↓ EAAT1, EAAT2 (Jz), EAAT3 (Jz), EAAT4, ↑ CAT1, ↔ 4f2hc  D18/19 ND	D15 ND  D18/19 ↓40% ↑ fetal loss	(DeChiara <i>et al.</i> , 1990; DeChiara <i>et al.</i> , 1991; Baker <i>et al.</i> , 1993; Liu <i>et al.</i> , 1993; Lopez <i>et al.</i> , 1996; Matthews <i>et al.</i> , 1999; Esquiliano <i>et al.</i> , 2009; Church <i>et al.</i> , 2012; Kent <i>et al.</i> , 2012)
Global IGF2 KO	Paternal transmission LacZDMR2 <sup>-</sup>	D16 ↓27%  D19 ↓40%	D16 ND  D19 ↓ Lz vd and volume of all Lz components, SA, FC length and diffusing capacity, ↑ Jz vd and BT	D16 ↔ System A and glucose transport, Snat1, Snat2, Snat4  D19 ↓ System A transfer and passive permeability and Snat2, ↔ glucose transport, Snat1 and Snat4	D16 ↓24%  D19 ↓52%	(Constancia <i>et al.</i> , 2005; Coan <i>et al.</i> , 2008b)
Fetal specific IGF2 KO	Inner cell mass Igf2 <sup>-</sup>	D17 ↓14%			D17 ↓27%	(Gardner <i>et al.</i> , 1999)
Placental trophoblast specific IGF2 KO	Trophectoderm Igf2 <sup>-</sup>	D17 ↓21%			D17 ↓12%	(Gardner <i>et al.</i> , 1999)
Placental Lz specific	Paternal transmission	D16 ↓20%	D16 ↔ Lz or Jz Vd	D16 ↑ System A and	D16 ↔/↓4%	(Constancia <i>et al.</i> , 2002;

IGF2 KO	Igf2P0-	D17 ↓24%  D19 ↓35%	↓ Lz Trophoblast, GlyT  D19 ↓ SA, trophoblast, FC volume, FC length, diffusing capacity, ↑ BT ↔ Lz or Jz vd and umbilical artery flow	glucose transport, Snat4, Glut3, ↓ passive permeability, calbindin, ↔ Snat1, Snat2, Glut1  D19 ↑/↔ System A transport, ↑glucose and calcium transport, ↓ passive permeability, ↔ Snat1, Snat2, Snat4, calcium transport, calbindin, PMCA1, TRPV6	D17 ↓24%  D19 ↓24%	Sibley <i>et al.</i> , 2004; Constancia <i>et al.</i> , 2005; Coan <i>et al.</i> , 2008b; Dilworth <i>et al.</i> , 2010; Kusinski <i>et al.</i> , 2011; Sferruzzi-Perri <i>et al.</i> , 2011; Dilworth <i>et al.</i> , 2013)
Global IGF1R KO	Igf1r-	D19 ↔	D18/19 ↔ Jz GlyT	D17 ↓ EAAT2 (Jz), EAAT3 (Lz and Jz), ↑ CAT1, ↔ EAAT1,EAAT4	D19 ↓55%	(DeChiara <i>et al.</i> , 1990; Louvi <i>et al.</i> , 1997; Matthews <i>et al.</i> , 1999) (Esquiliano <i>et al.</i> , 2009)
Global INSR KO	INSR-	D15 ↔ D18/19 ↔	D15 ↔ Jz GlyT D18/19 ↔Jz GlyT		D15 ND D18/19 ↓10%	(Louvi <i>et al.</i> , 1997; Esquiliano <i>et al.</i> , 2009)
PI3K p110α (Pik3ca)	Kinase dead heterozygote Pik3ca-D933A	D16 ↓9%	D16 ↓Lz vol, FC vol, FC length, MBS vol, SA, diffusing capacity, ↑ BT, ↔ Jz	D16 ↑ glucose and System A transfer per unit SA, ↔ Glut1, Glut3, Snat1, Snat2, Snat4	D16 ↓19%	(Sferruzzi-Perri <i>et al.</i> , 2016)

		D19 ↓12%	D19 ↓ Lz vol, FC vol, FC length, Troph vol, SA, diffusing capacity, ↑ BT, ↔ Jz	D19 ↑ glucose and System A transfer per unit SA, ↑ Prl3b1, ↔ Glut1, Glut3, Snat1, Snat2, Snat4	D19 ↓11%	
Global decreased AKT signalling through increased PTEN	Prl2-/-	D17 ↓22%	D17 ↓Jz, GlyT and Lz	D17 ↓ passive transport	D17 ↓17%	(Dong <i>et al.</i> , 2012)
Global decreased AKT1 signalling	Pkba-/- (exons 4-8 deleted)	D17 ↓33%  D19 ↓45%	D17 ↓ thickness, GlyT, Lz vessel density, length, area	D17 ↓ pAkt  D19 ↓ total Akt, pAkt ↑ Akt2 and Akt3	D17 ↓17%	(Yang <i>et al.</i> , 2003; Yung <i>et al.</i> , 2008)
Global decreased AKT1 signalling	Pkba-/- (exon 1 deleted)	D18 ↓30%	D18 ↔ Lz and Jz Vd	D18 ↓ pAkt, ↔ p-Akt	D18 ↓22% weight and ↑ fetal loss	(Cho <i>et al.</i> , 2001; Kent <i>et al.</i> , 2012)
Global decreased MAPK signalling	Erk2-/-	D11 ↓	D11 ↓ Lz thickness, FC development	↓ MAPK signalling	D11 ↓ weight and ↑ fetal loss	(Hatano <i>et al.</i> , 2003)
<b>Over-expression of IGF and downstream signalling</b>						
Global IGF2 over-expression	Maternal Igf2r-	D16 ↑40% D18 ↑25%			D16 ↑40% D18 ↑40%	(Ludwig <i>et al.</i> , 1996; Louvi <i>et al.</i> , 1997)
Global IGF2 over-expression*	Maternal H19Δ13-	D15 ↑37%  D16 ↑30%	D15 ↑ Jz GlyT  D16 ↑ volume of all placental components, ↑ SA, diffusing capacity, ↔BT	D15 ↑ Akt1 ↔ pAkt, p-ERK1/2 D16 ↓ glucose transfer, passive permeability and Glut3, ↔Glut1, Snat1, Snat2, Snat4	D15 ↑30%  D16 ↑12%	(Leighton <i>et al.</i> , 1995; Esquiliano <i>et al.</i> , 2009; Angiolini <i>et al.</i> , 2011; Church <i>et al.</i> , 2012)

		D18 ↑60% D19 ↑45%	D18 ↑Jz GlyT D19 ↑ volume of all placental components, SA, diffusing capacity	D19 ↓ glucose and System A transfer, passive permeability and Snat4, ↔ Glut1, Glut3, Snat1, Snat2	D18 ↑20% D19 ↑23%	
Global increased IGF2 and signalling via AKT	Double KO of maternal H19 and Pten +/-	D16 ↑65% D19 ↑80%	D16 ↑ Jz, GlyT D19 ↑ Jz, GlyT	↑ p-AKT and IGF2	D16 ↑31% D19 ↑31%	(Church <i>et al.</i> , 2012)
Global increased pAKT	Pten +/-	D16 ↑22% D19 ↑22%	D16 ↑ Jz, GlyT D19 ↑ Jz, GlyT	↑ p-AKT, ↔ IGF2	D16 ↑19% D19 ↑7%	(Church <i>et al.</i> , 2012)

1540 \*H19 null has biallelic expression of Igf2 combined with absence miR675 (encoded by H19)

1541 Abbreviations: BT=barrier thickness, D=day, FC=fetal capillaries, GLUT/Slc2a=glucose transporter, GlyT=trophoblast glycogen cells, IGF1/Igf1 =insulin-like growth factor-1, IGF2/Igf2=insulin-like growth factor-2, Jz=junctional zone, LAT=L-type amino acid transporter, Lz=labyrinthine zone, MBS=maternal blood space, ND=not determined, PI3K=phosphoinositol 3-kinase, SA=surface area, SNAT/Slc38a= Sodium-coupled neutral amino acid transporter, Vd=volume density.

1544 **Search terms used:** placenta, fetus, insulin-like growth factor, IGF, PI3K, ERK, MAPK, knock out, deficiency and/or transgenic

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1547 Table 3. The effect of maternal environmental challenge on fetal growth and placental structure, function and IGF signalling.

Maternal manipulation	Species	Timing	Placental				Fetal weight	Reference
			IGF and signalling	size	morphology	function		
<b>Nutrient restriction</b>								
20% UN	Mouse	D3-D19	D16 ↑ IGF1R ↓ Igf2P0 and PI3K signalling D19 ↓ Igf2P0 and PI3K signalling	D16 ↓6%  D19 ↓9%	D16 ↔Lz but ↓Jz and GlyT  D19 ↓Lz (MBS and FC vols and SA), ↔BT	D16 ↓ Glut1  D19 ↑ System A amino acid transport, ↑ Glut1, Snat2, ↓Snat4	D16 ↔  D19 ↓13%	(Coan <i>et al.</i> , 2010; Sferruzzi-Perri <i>et al.</i> , 2011)
10-30% UN	Guinea pig	-D28	D35/40 ↓Igf2, ↔ Igf1	D35 ↓20%  D60 ↓30%	D35 ↓ Jz volume ↔ Lz, but ↓ MBS, SA and ↑ BT  D60: ↓ Lz volume, MBS, FC, SA, ↑ BT, ↔ Jz,		D35 ↓29%  D60 ↓35%	(Roberts <i>et al.</i> , 2001; Olausson & Sohlstrom, 2003)
30% UN	Sheep	D22-D135	D135 ↔ Igf2	D135 ↓19% , altered placentome distribution			D135 ↓12%	(Osgerby <i>et al.</i> , 2002, 2004)
50% UN	Sheep	-D60-D30	D78 ↑ Insulin-IGF signalling (p-Akt and p-	D78 ↓29%	D78 ↑ vascularity		D78 ↔	(Zhu <i>et al.</i> , 2007b)

			ERK1/2)					
50% UN	Sheep	D28-D78	D78 ↑ Insulin-IGF signalling (p-ERK1/2, ↔pAkt) ↔ mTORC1 signalling D135 ↔	D78 ↓21%  D135 ↔		D78 ↑Glut3, GLUT1, Fatp4  D135 ↑ Fatp4	D78 ↓26%  D135 ↔	(Ma <i>et al.</i> , 2011)
UN gradual decrease to full food withdrawal	Sheep	D83-D90	D90 ↔ Igf2  D135 ↓Igf2	D90 ↓22%  D135 ↔		D90 ↔ Glut1, Glut3 D135 ↔ Glut1, Glut3	D90 ↔  D135 ↔	(McMullen <i>et al.</i> , 2005)
UN 50%	Cow	D30-D125	D125 ↑ Insulin-IGF signalling (p-Akt and p-ERK1/2) D250 ↔	D125 ↓27%  D250 ↓20%	D125 ↑ vascularity  D250 ↔ vascularity		D125 ↔  D250 ↔	(Zhu <i>et al.</i> , 2007a)
70% UN	Baboons	D30-D165	D90 ↓ Igf2, IGF2R, ↑ IGF1R, ↔ Igf1 or IGF1 D120 ND	D90 ↔  D120 ↔		D90 ND  D120 ↓ System A amino acid transport, ↔ system L amino acid transport, GLUT1 TAUT, SNAT1, SNAT2, SNAT4, LAT1,	D90 ↔  D120 ↔	(Li <i>et al.</i> , 2007; Pantham <i>et al.</i> , 2015) (Kavitha <i>et al.</i> , 2014)

			D165 ↓ insulin/IGF-I, MAPK (IRS-1, Akt S6K, ERK-1) and mTOR signalling	D165 ↓20%		LAT2 D165 ↓ System A and L amino acid transport, GLUT1, TAUT, SNAT2, LAT1, LAT 2	D165 ↓19%	
<b>Low protein diets</b>								
16% v 20% protein (0.80CT)	Mouse	D3-19	D16 ↔ Igf2, H19  D19 ↔ Igf2, H19	D16 ↑5%  D19 ↑5%	D16 ↓ Lz/Jz ratio  D19 ↓ Lz/Jz ratio	D16 ↑ glucose transport, Glut1, ↔ System A amino acid transport  D19 ↓ System A amino acid transport, Snat4, ↔ glucose transport,	D16 ↔  D19 ↔	(Coan <i>et al.</i> , 2011)
8% vs 20% protein (0.40CT)	Mouse	D3-19	D16 ↑ total Igf2 ↔ Igf2P0, H19	D16 ↔	D16 ↔	D16 ↑ glucose transport, Snat2, ↔ System A amino acid	D16 ↔	(Coan <i>et al.</i> , 2011)



			D19 ↔ Igf2, H19	D19 ↑4%	D19 ↔	transport D19 ↓ Snat1, Snat4, ↔ glucose and System A amino acid transport	D19 ↓9%	
9% vs 17% protein (0.53CT)	Rat	D1-22	D22 ↑Igf1 ↓Igf2 ↔Igf1r,Igf2r,Insr	ND			D22 ↓8%	(Nusken <i>et al.</i> , 2011)
6% vs 20% protein (0.30CT)	Rat	D1-21	D14 ↓ Lz Igf2, Insr in female and ↑ Lz IGF2, ↓ Igf1r in male  D16 ↓ Lz Igf2 in female and male D21 ↓ Lz IGF2 in male and female	D14 ↓25% D18 ↓12%  D21 ↔	D14 ↓Lz and Jz vol D18 ↓Lz vol, ↑ trophoblast stem cells and Lz sinuosoidal GiT, ↓ spongiotrophoblast and GiT cells, ↔Jz D21↔Lz ↓Jz		D14 ↓21.5 D18 ↓27  D21 ↓14%	(Gao <i>et al.</i> , 2012a; Gao <i>et al.</i> , 2012b, 2013)
4% vs 18% protein (0.22CT)	Rat	D2-21	D19 and D21 ↓ mTOR  D21 ↓ PI3K signalling (p-Akt-T308)	D15-19 ↔  D21 ↓12.5%	ND	D19 and D21 ↓ Systems A and L amino acid transport, LAT1, LAT2, SNAT2, ↔ glucose transport,	D15-19 ↔  D21 ↓21%	(Jansson <i>et al.</i> , 2006{Rosario, 2011 #3227; Pantham <i>et al.</i> , 2016)

						SNAT4		
<b>Obesogenic diets</b>								
2.5-times fat	Mouse	-D28-D1	D13 ↓ Igf2, Mtor, ↔ Igf1  D18 ↓ Igf2, Igf2r, ↔ Igf1	D13 ↓20%  D18 ↑15% in males, ↔ females	↔ Lz	D13 ↓ Snat1, Glut1, ↔ Cd36  D18 ↓ Cd36, ↔ Snat1, Glut1	D13 ↓28%  D18 ↓15%	(Sasson <i>et al.</i> , 2015)
2.5-times fat	Mouse	-D28-D18	D1 ↑ Igf1r, ↓ Igf2, Igf2r, Mtor, ↔ Igf1  D18 ↓ Igf2, Igf2r, ↔ Igf1	D13 ↓20% in males  D18 ↔ males or females	↔ Lz	D13 ↓ Snat1, Glut1, ↔ Cd36  D18 ↓ Cd36, ↔ Snat1, Glut1	D13 ↓25%  D18 ↓25%	(Sasson <i>et al.</i> , 2015)
2.5-times fat	Mouse	D1-D18	D13 ↑ Igf1r, ↓ Igf2, Igf2r, Mtor, ↔ Igf1  D18 ↓ Igf2, Igf2r ↔ Igf1	D13 ↓20% in males  D18 ↑15% in males	↔ Lz	D13 ↓ Snat1, Glut1, Cd36  D18 ↓ Glut1, Cd36 ↔ Snat1	D13 ↓28%  D18 ↓28%	(Sasson <i>et al.</i> , 2015)
5.3-times fat	Mouse	-D84-D19	D15 ↑ Igf2 and Igf2r male, ↔ female  D19 ↔ Igf2 and Igf2r	D15 ↔  D19 ↔		D15 ↑ Lz Snat2 in male, ↑ Lz Snat4 in female  D19 ↔	D15 ↔  D19 ↓8% in males	(King <i>et al.</i> , 2013)
6-times fat	Mouse	D1-15	D15 ↑ Igf1, ↓ Irs1 in	D15 ↑7%	D15 ↔ Lz or	D15 ↓ Slc22a1,	D15 ↔	(Gallou-Kabani <i>et al.</i> ,

			males, ↔ IGF2, IGF2P0, IGF2R, H19		vascularity	↑ Slc22a2 *sexually dimorphic response of placenta		2010; Gabory <i>et al.</i> , 2012)
2.5-times fat	Rat	D1-D21	D21 ↔ mTORC1 signalling	D21 ↔	D21 ↓ Jz		D21 ↓5%	(Mark <i>et al.</i> , 2011)
5-6-times fat	Rat	-D49-D21	D21 ↑ mTORC1 signalling, ↔ Insulin-IGF signalling (p-Akt or p-MAPK)	D21 ↔		D21 ↓ SNAT1, ↔ Systems A and L amino acid transport and LPL activity, SNAT2, SNAT4, GLUT1, GLUT3, GLUT9, FATP4, FATP6, LPL	D21 ↑7%	(Gaccioli <i>et al.</i> , 2013)
3-times fat and 5-times sugar diet	Mouse	-D42-D18	D18 ↓ mTORC1 signalling, ↔ Insulin-IGF PI3K (p-AKT, IRS1, PI3K-p85)	D18 ↔			D18 ↔	(Lager <i>et al.</i> , 2014)
4-times fat and 1.3-times sugar	Mouse	-D20-D19	D19 ↑ Insulin/IGF-PI3K (p-IRS1, p-Akt-T308) and mTORC1 signalling, ↔MAPK	D19 ↔		D19 ↑ Systems A and L amino acid transport, SNAT2, LAT1, GLUT1, GLUT3,	D19 ↑18%	(Diaz <i>et al.</i> , 2015; Rosario <i>et al.</i> , 2015; Rosario <i>et al.</i> , 2016)

						FATP6, ↔ SNAT4, LAT2, CD98, FAT/CD36, FATP2, FATP4		
3-times fat and 5-times sugar diet	Mouse	D1-D19	D16 ↑ Igf2, IgfP0, H19, Insulin/IGF-PI3K signalling (PI3K-p110α, p-Akt), ↓ INSR, ↔ mTORC1 or MAPK  D19 ↑ Insulin/IGF-PI3K signalling (PI3K-p110α, p-Akt, p-MAPK), ↔ Igf2, Igf2P0, H19, INSR or mTORC1	D16 ↓11%  D19 ↓8%	D16 ↓Lz FC ↑BT  D19 ↓Lz, MBS, BT, SA and ↓GlyT	D16 ↑ glucose and System A amino acid transport, Glut3, Snat2  D19 ↑ FATP1, ↔ glucose and System A amino acid transport	D16 ↓9%  D19 ↔	(Sferruzzi-Perri <i>et al.</i> , 2013)
50% greater food intake	Sheep	-D60-D135	D70-75 ↓ p-IRS1, p-mTORC1, p-MAPK in the arterial tissues, ↔ INSR, IGF1R  D165 ND	D70-75 ↓22%  D165 ↔	D70-75 ↑ arteriole diameters, ↓ vessel density  D165 ↔	D70-75 ↑ Fatp1, Fatp4, Cd36, Lpl  D165 ↑ GLUT3, FATP1, Fatp4, Cd36, ↔ Lpl	D70-75 ↑20-26%  D165 ↔	(Zhu <i>et al.</i> , 2009; Ma <i>et al.</i> , 2010; Zhu <i>et al.</i> , 2010; Tuersunjiang <i>et al.</i> , 2013)
<b>Hypoxia</b>								
13%	Mouse	D1-D19	D19 ↑ Insulin-IGF	D19 ↑10%	D19 ↑ Maternal	ND	D19 ↓12% weight and	(Matheson <i>et al.</i> ,

			(↑ p-Akt) and mTORC1 signalling		arterial and venous blood space		litter size	2015)
13% hypoxia	Mouse	D11-16	D16 ↓ Igf2, ↔ Igf2PO, altered p-Akt (depending on site phosphorylated)	D16 ↔	D16 ↑ Lz ↑ MBS, trophoblast vol, SA exchange	D16 ↔ System A amino acid amino acid or glucose transport, Gluts and Snats	D16 ↔	(Higgins, 2015 #5541
13% hypoxia	Mouse	D14-19	D19 ↑ Igf2, Igf2PO, altered insulin-IGF signalling (↓ INSR, IGF1R, PI3K-p85α, PI3K-p110α but ↑ p-Akt)	D19 ↔	D19 ↑ FC volume and density, ↓ BT	D19 ↑ glucose transport, Snat1, ↔ System A amino acid amino acid transport	D19 ↓5%	(Higgins <i>et al.</i> , 2015)
12% Hypoxia	Mouse	D14.5-18.5	D18.5 ↓ Igf2r and Igf2, Igf1r in females	D18.5 ↔	D18.5 ↓ Lz blood space, ↑ tissue in females	D18.5 ↓ Glut1, ↑ Snat1 in females, ↔ Glut3	D18.5 ↓6.5%	(Cuffe <i>et al.</i> , 2014)
10% hypoxia	Mouse	D14-19	D19 ↓ Insulin-IGF signalling (↓ INSR, IGF1R, PI3K-p85α, PI3K-p110α and p-Akt), ↔ Igf2, Igf2PO	D19 ↔	D19 ↓ Lz vd, MBS volume, SA exchange, ↑ Jz vd, trophoblast vol and BT	D19 ↓ System A amino acid transport, ↔ glucose transport but altered uterine artery vasoreactivity	D19 ↓21%	(Higgins <i>et al.</i> , 2015; Skeffington <i>et al.</i> , 2015)

<b>Endocrine disruption</b>								
Corticosterone 83µg/g/day	Mouse	D11-D16	D16 ↓ p-Akt, ↔ Igf2, Igf2PO, INSR, IGF1R, mTORC1 signalling	D16 ↓6%	D16 ↓ FC vol and Vd, ↑ MBS and Troph Vd, ↔ SA, BT	D16 ↓ Glut1, Glut3, Snat1, Snat2, ↔ glucose or System A amino acid transport and Snat4	D16 ↓7%	(Vaughan <i>et al.</i> , 2012; Vaughan <i>et al.</i> , 2015)
Corticosterone 81µg/g/day	Mouse	D11-D19	D19 ↓ mTORC1 signalling, ↔ Igf2, Igf2PO, INSR, IGF1R, p- Akt	D19 ↓12%	D19 ↔ FC, MBS, Troph, SA, BT	D19 ↓ glucose and System A amino acid transport, ↑ Snat1, ↔ Glut1, Glut3, Snat2, Snat4	D19 ↓19%	(Vaughan <i>et al.</i> , 2012; Vaughan <i>et al.</i> , 2015)
Dexamethasone 24µg/kg/day	Mouse	D13-D16	D16 ↓ MAPK1 D18 ↔ MAPK1  D16 and D18 ↔ Igf2	D16 ↓20% female only  D18 ↔	D16 ↓Jz area female only  D18 ND	D16 and D18 ↔ Glut1, Glut3, Snat1, Snat2, Snat4	D16 ↓20%  D18 ↔	(Cuffe <i>et al.</i> , 2011)
Dexamethasone 24µg/kg/day	Rat	D13-D20	D20 ↓pAkt in Jz	D20 ↓50%		D20 ↓ Prls in Jz, ↑ Prls in Lz	D20 ↓22%	(Ain <i>et al.</i> , 2005)
Diabetes via streptozotocin administration neonatally	Rat		D20 ↑ Igf1, Igf2, Igf2r, IGF1R kinase and autophosphorylation activity, ↔ Igf1r, Insr	D21 ↑22%	D21	D21 ↑ glycerol and FFA release	D21 ↑13%	(Hauguel-de Mouzon <i>et al.</i> , 1992; Martinez <i>et al.</i> , 2008; White <i>et al.</i> , 2015)

Diabetes via streptozotocin administration 1 week before mating	Rat	-D7-D21	D21 ↓ Insr, Irs1, Igf2, Igf2r, ↔ Irs2, Igf1r	D21 ↑ 22%	D21 ↑ Lz, ↑ lacunae	D21 ↓ Glut1, ↑ Lpl, ↔ Glut3, Snat2, Snat4, Lat1	D21 ↑ 5% or ↔	(Cisse <i>et al.</i> , 2013)
Insulin resistance via heterozygous p110 $\alpha$ deficiency	Mouse		D16 ↓ PI3K signalling  D19 ↓ PI3K signalling	D16 ↔  D19 ↑ 15%	D16 ↓ Lz Troph vol, ↓ BT  D19 ↑ Jz vol ↑ SA diffusing capacity	D16 ↓ glucose transfer, Snat1, ↔ System A amino acid transfer, Glut1, Glut3, Snat2, Snat4  D19 ↓ glucose transfer, Glut1, Snat1, Snat2, Prls, ↔ System A amino acid transfer, Glut3, Snat4	D16 ↔  D19 ↔	(Sferruzzi-Perri <i>et al.</i> , 2016)  *Depends on fetal genotype
<b>Other manipulations affecting conceptus growth</b>								
<b>Restriction of utero-placental blood flow</b>								
Uterine ligation	Mouse	D18	D20 ↓ Igf1, Igf2	D20 ↔	D20 ↓ Lz depth, vol,	D20 ↓ Slc5a9,	D20 ↓ 11%	(Habli <i>et al.</i> , 2013;

					vessel area	Slc7a10, 4F2hc, Lat1, Lat2, Snat2, GLUT1, GLUT8, ↑ Snat1, ↔ GLUT3, GLUT9		Jones <i>et al.</i> , 2013; Jones <i>et al.</i> , 2014)
Uterine ligation	Rat	D17	D20 ↓ Igf2	D20 ↓8%			D20 ↓20%	(Price <i>et al.</i> , 1992)
Uterine ligation	Rat	D18 or D19	D20 ↓ IGF1R, ↔ INSR	D20 ↔ or ↓25%	D20↑ diameter, ↔ Lz vd	D20 ↓ GLUT1, ↔ GLUT3	D20 ↓7% or 27% weight and ↓ litter size	(Das <i>et al.</i> , 1998; Reid <i>et al.</i> , 2002; Wlodek <i>et al.</i> , 2005)
Uterine ligation	Rat	D19	D22 ↓ Igf1, ↔ Igf2, Insr, Igf1r, Igf2r	ND			D22 ↔	(Nusken <i>et al.</i> , 2011)
Uterine ligation	Guinea pig	D30	D55-60 ↔ Igf1, Igf2	D55-60 ↔ or 37%		D55-60 ↓ System A amino acid transfer, ↔ glucose transfer	D55-60 ↓7% or 38%	(Jansson & Persson, 1990; Carter <i>et al.</i> , 2005)
Placental embolism	sheep	D113-120	D131 ↓ IGF1R, ↔ IGF-I	D131 ↓30%			D131 ↓21%	(Bloomfield <i>et al.</i> , 2002a; Shaikh <i>et al.</i> , 2005)
Placental embolism	sheep	D103-109	D131 ↔ Mtor	D131 ↓43%		D131 ↓ Glut1, Slc7a1, Slc7a8, ↔ Glut3, Glut4, Snat4, Slc7a5	D131 ↓20%	(Wali <i>et al.</i> , 2012)



Uterine carunclectomy	sheep	-D70	D130-134 ↑ Igf2, ↔ Igf1, Igf1r, Igf2r	D130-134 ↓ 30-40%	D130-134 altered distribution of placentome types and ↓ placentome number but ↑ individual weight of placentomes, trophoblast and maternal capillary volume and SA of placentomes	D130-134 ↓ Fatp4, ↔ Glut1, Glut3, Glut4, Slc7a1, Slc7a5, Snat1, Snat4, Fatp1, Cd36, Fabp5	D130-134 ↓26%	(Zhang <i>et al.</i> , 2016b)
<b>Hyperthermia</b>	Sheep	D39-D125	D55 ↑ IGF2  D90 ↑ IGF1 , p-mTORC1, ↓ p-Akt, ↔ MAPK  D135 ↑ pAkt, p-MAPK dys-regulated mTORC1 signalling (↑ p- mTORC1 but ↓ p-p70)	D55 ↔  D90 ↓24%  D135 ↓58%		D135 ↑ Slc7a5, Slc7a8, uterine blood flow, trans-placental oxygen diffusion, ↓ branched amino acid and glucose transport,	D55 ↔  D90 ↔  D135 ↓47%	(Thureen <i>et al.</i> , 1992; Ross <i>et al.</i> , 1996; Anderson <i>et al.</i> , 1997; Regnault <i>et al.</i> , 2003; de Vrijer <i>et al.</i> , 2004; Regnault <i>et al.</i> , 2005; de Vrijer <i>et al.</i> , 2006; Regnault <i>et al.</i> , 2007; Arroyo <i>et al.</i> , 2009; Arroyo <i>et al.</i> , 2010)

						↔ utero-placental oxygen uptake		
<b>Alcohol consumption</b>	Rat	-D4-D4	D20 ↓ Lz Igf1, and Lz Igf1r in males, ↔ Igf2  ↑ Jz Igf2, ↔ Jz Igf1, Lz or Jz Igf2r	D20 ↔	D20 ↑ length and width, ↓ Lz and ↑ Jz and ↑ GlyT in females	D20 ↓ Lz Snat2, ↔ Lz Snat1, Snat4, Glut1, Glut3 and ↓ Jz Glut1 in males, ↑ Jz Glut1 in females	D20 ↓7%	(Gardebjer <i>et al.</i> , 2014)

1548 Gestational age: mouse ~20 days, rats ~23 days, guinea pigs ~70 days, sheep ~150 days, cows ~283 days, baboons ~183 days.

1549 Abbreviations: BT=barrier thickness, D=day, FATP=fatty acid transport protein, FC=fetal capillaries, GLUT=glucose transporter, GiT- giant trophoblast cells, GlyT- trophoblast glycogen cells,

1550 IGF1/Igf1=insulin-like growth factor-1, IGF2/Igf2=insulin-like growth factor-2, Jz=junctional zone, LAT=cationic amino acid transporter, LPL=lipoprotein lipase, Lz=labyrinthine zone,

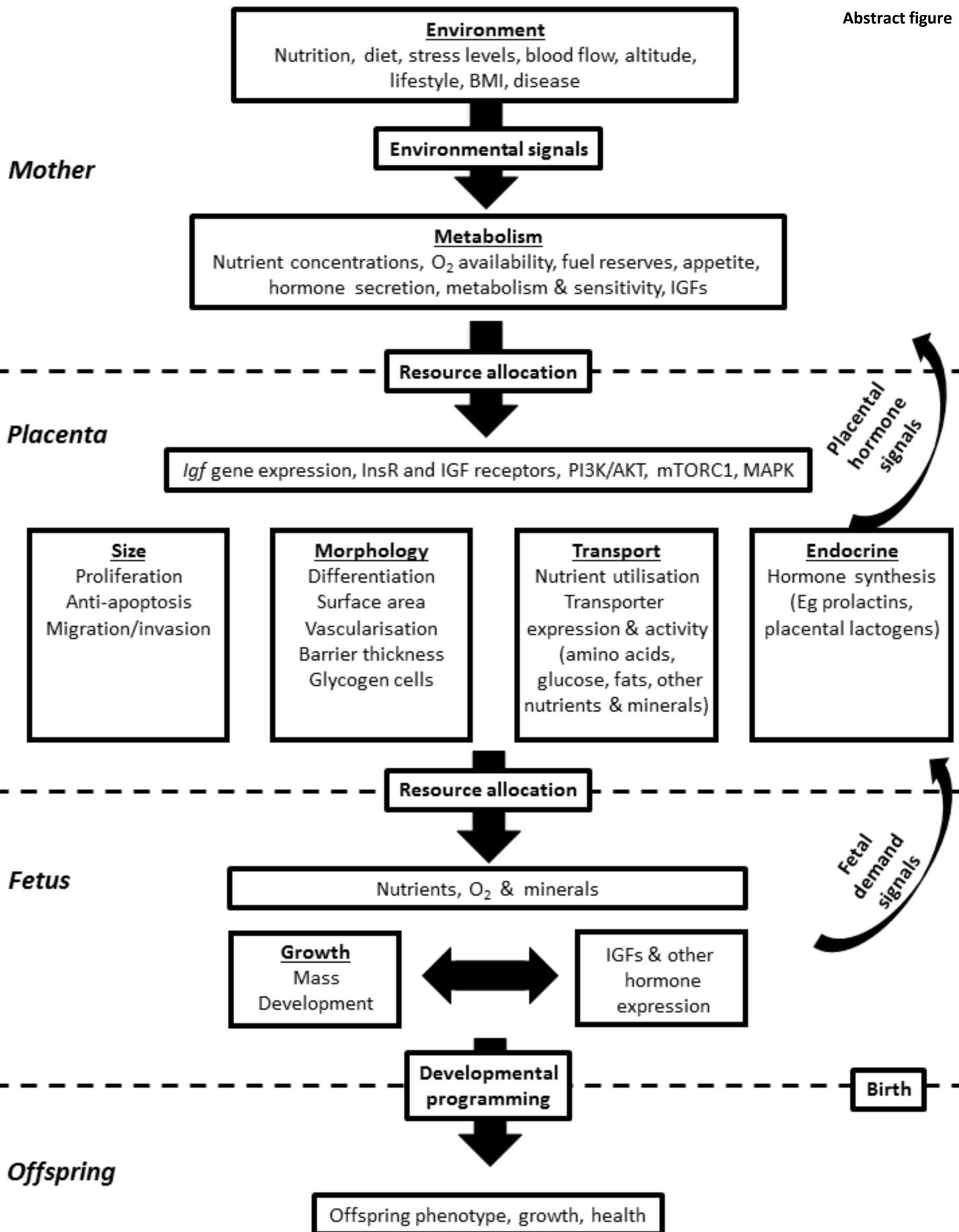
1551 MAPK/ERK=mitogen activated kinase, MBS=maternal blood space, mTOR=mechanistic target of rapamycin, p=phosphorylation, PI3K=phosphoinositol 3-kinase, Prl=prolactin-related hormone,

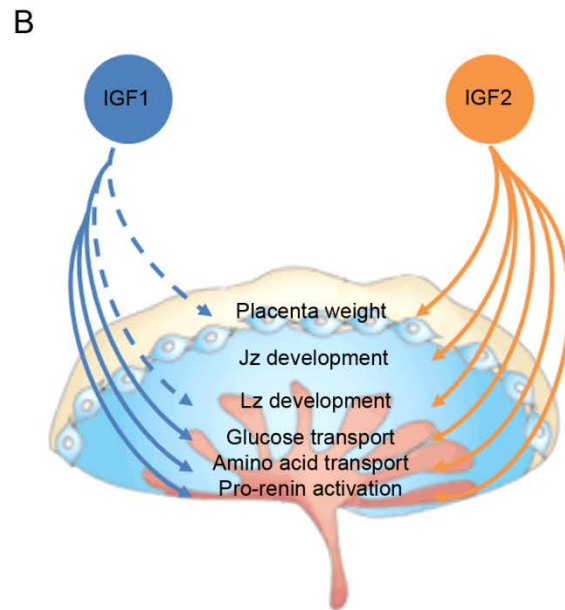
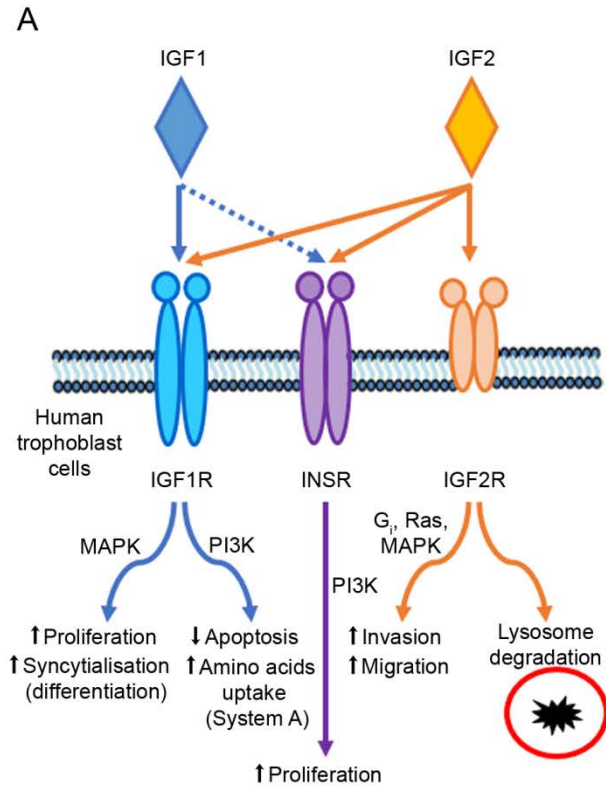
1552 SA=surface area, SNAT/Slc38a= Sodium-coupled neutral amino acid transporter, UN=undernutrition; vol=volume, vd=volume density.

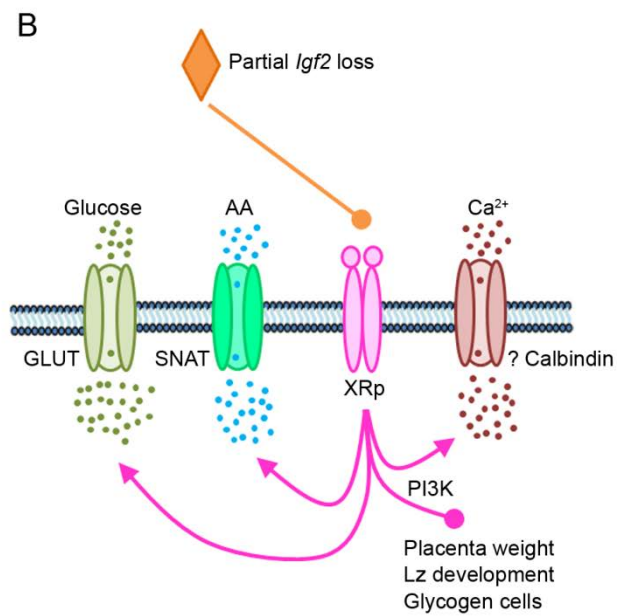
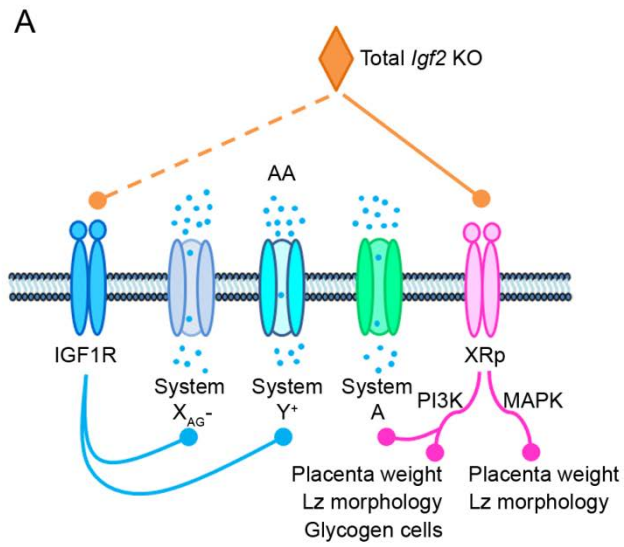
1553 **Search terms used:** placenta, fetus, insulin-like growth factor, IGF, nutrient restriction, undernutrition, low protein diet, high sugar, high fat, obesogenic, IUGR, PI, hypoxia, uterine ligation,

1554 corticosterone, dexamethasone carunclectomy, heat stress and/or diabetes.

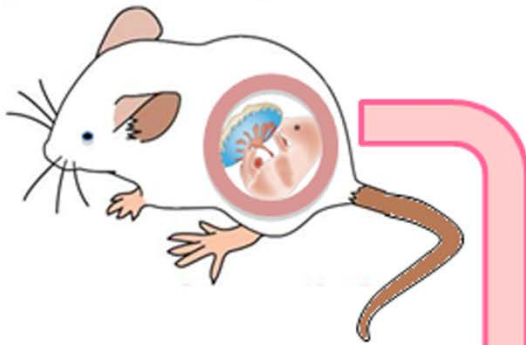
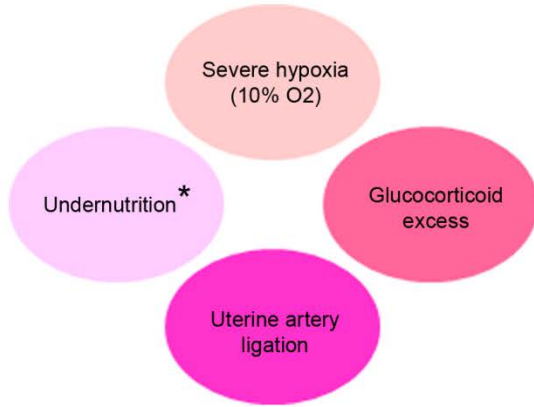
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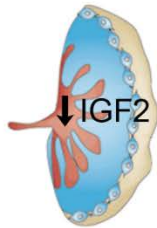




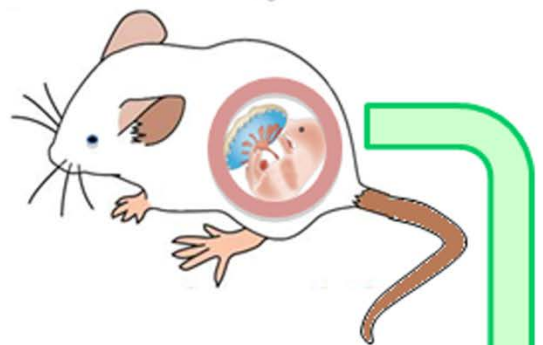
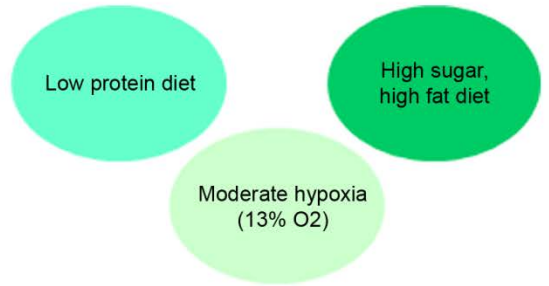
A



- ↓ AKT signalling
- ↓↔ Placenta weight
- ↓ Lz development
- ↓ Glucose transport
- ↓↑ Amino acid transport



B



- ↑ AKT signalling
- ↓↑ Placenta weight
- ↓↑ Lz development
- ↑ Glucose transport
- ↓↑ Amino acid transport

