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35

36 **Abstract**

37 **Context:** Maternal obesity and gestational diabetes mellitus (GDM) can both contribute to
38 adverse neonatal outcomes. The extent to which this may be mediated by differences in
39 placental metabolism and nutrient transport remains to be determined.

40 **Objective:** To examine whether raised maternal BMI and/or GDM contributed to a resetting of
41 the expression of genes within the placenta that are involved in energy sensing, oxidative stress,
42 inflammation and metabolic pathways.

43 **Methods:** Pregnant women from Spain were recruited as part of the PREOBE survey at the first
44 antenatal visit (12-20 weeks of gestation) and stratified according to pre-pregnancy BMI and the
45 incidence of GDM. At delivery, placenta and cord blood were sampled and newborn
46 anthropometry measured.

47 **Results:** Obese women with GDM had higher estimated fetal weight at 34 gestational weeks,
48 greater risk of preterm deliveries and Caesarean section. Birth weight was unaffected by BMI or
49 GDM, however, women who were obese with normal glucose tolerance had increased placental
50 weight and higher plasma glucose and leptin at term. Gene expression for markers of placental
51 energy sensing and oxidative stress, were primarily affected by maternal obesity as *mTOR* was
52 reduced whereas *SIRT-1* and *UCP2* were both upregulated. In placenta from obese women with
53 GDM gene expression for *AMPK* was also reduced whereas the downstream regulator of
54 mTOR, *p70S6KB1* was raised.

55 **Conclusions:** Placental gene expression is sensitive to both maternal obesity and GDM which
56 both impact on energy sensing, and could modulate the effect of either raised maternal BMI or
57 GDM on birth weight.

58

59 **Introduction**

60 Obesity is of great importance to individual and global health [1]. Its prevalence amongst
61 women of reproductive age is increasing [2] so that, in Spain for example, up to 17% of
62 pregnant women are obese [3]. The increased prevalence of obesity in pregnant women has
63 occurred concurrently with an increase in gestational diabetes mellitus (GDM) [4] which now
64 affects up to 14% of all pregnancies in the US, and around 2–6% of pregnancies in Europe [5,
65 6]. Raised maternal body mass index (BMI) and GDM are both associated with adverse
66 metabolic adaptations in the mother. These include increased risks of miscarriage and stillbirth,
67 preeclampsia [7] and both intrauterine growth restriction and macrosomia [8], conditions with
68 the potential to compromise fetal and newborn survival and health [9-11].

69
70 Consumption of an unhealthy diet in pregnancy has been linked to increased gestational weight
71 gain (GWG) [12], raised BMI [13] and GDM [11] that are associated with fetal overgrowth [14].
72 Placental nutrient supply is one mechanism linking maternal nutritional status and fetal growth
73 and is dependent on utero-placental blood flow, hormone production and nutrient transfer
74 capacity, which is itself dependent on the type, number and activity of a range of nutrient
75 transporters [15]. Increased glucose and lipid transport in GDM [16, 17] are also accompanied
76 by placental defects arising from compromised trophoblast invasion and blood vessel formation
77 [18]. Although the association between high pre-pregnancy BMI and fetal overgrowth is well
78 established for type 1 diabetes [19], the effect of maternal BMI on placental function in women
79 without GDM, its relationship to GWG [20] and its relationship to current diet remains
80 unknown [21, 22].

81
82 Obesity is associated with perturbed maternal metabolism, raised plasma hormones, including
83 leptin, insulin and IGF1 and the accumulation of inflammatory markers (e.g. interleukin-6) [21].

84 Insulin signalling is crucial for the regulation of intracellular and blood glucose concentrations.
85 Alterations in the number of insulin binding sites, reflecting placental IR expression, have been
86 demonstrated in obesity [23] and diabetes mellitus [24]. Fetal glucose and amino acids and
87 placental insulin/IGF1 signalling act as upstream regulators of the mammalian target of
88 rapamycin (mTOR), which is central to energy sensing and can be reset by maternal obesity and
89 GDM [25] through phosphorylation mechanisms. These responses are mediated through
90 changes in NFkB signalling, thereby resetting pro-inflammatory and pro-oxidative pathways
91 [26] acting through toll-like receptor (TLR4) [27]. Furthermore, mTOR inactivation occurs
92 through the AMP-activated protein kinase (AMPK) pathway [28], whilst uncoupling protein
93 (UCP)2 limits oxidative damage within the placenta by decreasing reactive oxygen species
94 (ROS) production [29]. Free fatty acids also decrease peroxisome proliferator-activated receptor
95 gamma (PPAR) γ expression [30] whilst activating myeloid pro-inflammatory cells, although
96 whether these placental responses can be modulated by BMI and/or GDM are not established.

97

98 In the present study, we aimed to determine whether maternal BMI and/or GDM influenced
99 placental homeostasis and energy balance and thus impact on birth outcomes. The establishment
100 of direct links between maternal nutritional status, the placenta and weight at birth will give
101 insight on mechanistic pathways thereby enabling targeted interventions designed to prevent
102 adverse outcomes under these conditions.

103

104 **Materials and Methods**

105 *Participants*

106 The subjects participated in a longitudinal study on the influence of body composition by
107 maternal genetics and nutrition (PREOBE study: P06-CTS-02341) undertaken between 2007
108 and 2010 and registered with www.ClinicalTrials.gov, (NCT01634464) [31, 32]. It was
109 conducted according to the guidelines in the Declaration of Helsinki and all experimental
110 procedures approved by the Ethics Committees for Granada University, San Cecilio University
111 Hospital and the University of Nottingham. Witnessed, written informed consent was obtained
112 from all subjects before their study inclusion and participants were assured of anonymity.
113 Anthropometric assessments of were undertaken following the standards established by the
114 Spanish Society of Gynaecology and Obstetrics, the Fetal Foundation and the Spanish
115 Association of Paediatrics.

116

117 In the overall PREOBE study (Figure 1), 474 pregnant women aged 18-45, with singleton
118 pregnancies, were assessed for eligibility between 12-20 weeks gestation at two different
119 primary health care settings (Clinical University Hospital “San Cecilio” and the “Mother-Infant”
120 University Hospital) in Granada, Spain. Amongst these, 124 declined to participate. Criteria for
121 exclusion (n=19) were participation in another study simultaneously, receiving drug treatments,
122 being underweight (BMI<18.5 kg/m²), having type 1 diabetes or pre-existing disease. Therefore,
123 331 women were included in the project and classified according to their BMI (based on self-
124 reported pre-pregnancy weight provided on enrolment) as normal weight (pre-pregnancy
125 BMI \geq 18.5 but <25 kg/m²; n=132), overweight (pre-pregnancy BMI \geq 25 but < 30 kg/m²; n=56)
126 and obese (pre-pregnancy BMI \geq 30 kg/m²; n=64). In addition, 79 women were diagnosed with
127 GDM following measurement of raised fasting plasma glucose concentrations, 25 women after a
128 75g oral glucose tolerance test (OGTT) between 16-18 weeks gestation [11], if they either had a

129 family history of GDM, or had previously had GDM, or were obese, whilst 54 women after an
130 additional 100 g OGTT between 24-28 weeks gestation.

131

132 The number of women in each BMI group for whom collection of biological samples was
133 achieved at the time of delivery are shown in Figure 1. Amongst these a subpopulation of 135
134 subjects, underwent molecular analysis in Nottingham (i.e. ~half of those sampled within each
135 group - 59 normal weight, 29 overweight, 22 obese, 25 GDM). The 25 mothers with GDM were
136 subsequently classified according to their BMI as normal weight GDM (pre-pregnancy
137 $BMI \geq 18.5$ but $< 25 \text{ kg/m}^2$; $n=14$) and obese GDM (pre-pregnancy $BMI \geq 30 \text{ kg/m}^2$; $n=11$).
138 Participants diagnosed with GDM then had increased medical supervision and received
139 nutritional advice for meal plans designed to control normoglycaemia, with none receiving
140 insulin.

141

142 During pregnancy, each mother attended additional PREOBE study medical visits at 24 (BMI
143 group) or 34 weeks of gestation (BMI and GDM groups), . Gestational age was calculated as
144 from the last menstrual period and through ultrasound scan considering a gestational age below
145 37 weeks as preterm delivery. Anthropometric characteristics of the fetus were estimated by
146 using ultrasound scan at 34 gestational weeks. **When there was a disagreement between the last
147 menstrual period and ultrasound, the measurements taken by ultrasound were used to calculate
148 the gestational age [33].**

149

150 Maternal weight gain (GWG) during pregnancy was defined as weight change to the last
151 recorded weight in the 34th gestational week and compared to the 2009 IOM guidelines [34].
152 Large (LGA) and small (SGA) for gestational age infants were defined according to the
153 Lubchenco growth curves [35] with standard adjustment for gestational age at birth i.e. birth

154 weights >90th population centile were defined as LGA infants and those <10th population centile
155 as SGA.

156

157 *Maternal nutrient intake*

158 This was collected during late gestation (34-40 weeks) using standardised 7 day dietary records
159 given during their second visit. Each participant was given verbal and written instructions on
160 how to record food and drinks consumed with a booklet of common food items and mixed
161 dishes to facilitate estimation of portion sizes. Near delivery, food records were reviewed
162 individually by a nutritionist for completeness and accuracy of food description and portion size.
163 Nutritional data were analysed for nutrient intake by using a nutritional software program
164 (CESNID 1.0: Barcelona University, Spain) based on validated Spanish food tables [36]. These
165 results were compared with a food frequency questionnaire taken at 24 weeks gestation and both
166 sets of records were reviewed with the mother around the time of delivery by a professional
167 nutritionist with respect to their accuracy, thereby avoiding the potential inaccuracies associated
168 with these types of records [37].

169

170 *Collection and analysis of blood samples*

171 Maternal venous blood was collected at 24, 34 weeks of gestation and during labour. Umbilical
172 venous blood samples were collected within 30 minutes after placental delivery from a double-
173 clamped section of umbilical cord. EDTA and serum collection tubes were used (Vacutainer®
174 Refs: 368857 & 367953) for haematological assessment and biochemical analyses respectively.
175 Blood samples for serum preparation were left at 4°C for 15 minutes to allow blood clotting,
176 centrifuged at 3,500 rpm for 10 minutes, and the serum fraction transferred into a sterile tubes.
177 Samples were stored at 4°C for same day analyses or at -80°C for further analysis.
178 Haematological parameters were analysed using a haematology analyser (Sysmec XE-2100,
179 Roche Diagnostic) and flow cytometer (Advia 120-160858, Bayer HealthCare, Tarrytown, NY).

180 Plasma glucose and triglycerides were measured enzymatically (Modular Analytics EVO,
181 Roche, Neuilly sur Seine Cedex, France), whilst serum leptin concentrations measured by
182 ELISA (Biosource Kap 2281, Denmark).

183

184 *Collection of placenta samples*

185 Placenta were collected and weighed immediately after delivery. Disc samples containing both
186 maternal and fetal tissue were obtained from identical portions of the placental plate to avoid
187 any as regional variations. Visual inspection of the placenta for necrosis or any other
188 abnormality was undertaken by experienced clinicians. **This included the measurement of**
189 **placental size, weight and morphology and if there was any abnormality such as multilobules, or**
190 **placenta spuria, annular, membranous, infarction, chorangiosis or vasculopathies, a sample was**
191 **either obtained from a healthy region.** Then after removal of the decidua a representative
192 0.5×0.5×0.5cm (200mg) sample was excised from the middle of the radius (distance between
193 the insertion of the umbilical cord and the periphery) of each placenta, rinsed twice with saline
194 solution (NaCl 0.9%) and immediately placed into sterile 1.5ml microtubes containing RNAlater
195 solution (Qiagen Ltd., Crawley, UK). All samples were stored under RNase free conditions
196 using liquid nitrogen before storage at -80°C for later analysis in Nottingham.

197

198 **Laboratory analysis**

199 *Gene expression*

200 Total RNA was extracted from 100mg of maternal placenta tissue using 200µl of chloroform per
201 1mL of TRI reagent solution (Sigma Chemical Co. Poole, UK) and RNeasy extraction kit
202 (Qiagen Ltd., Crawley, UK). Two µg RNA was used to generate 20µl cDNA using High
203 Capacity RNA-to-cDNA kit (Applied Biosystems, CA 94404, USA). Negative control RT
204 samples lacking Enzyme Mix (-RT) were included for each sample.

205

206 Real-time PCR using 15µl of reactions consisting of 4.5µl diluted 1:10 cDNA, 3.0µl (final
207 concentration of 250 nM) gene specific primers (Table 1), and 7.5µl of SYBR Green mastermix
208 (Thermo Scientific, ABgene Ltd. Epsom, UK) were performed. Duplicate samples were run for
209 40 cycles with negative controls in 96-well plates using the Techne Quantica Thermocycler
210 (Techne Inc., Barloword Scientific, Stone, UK). Ten-fold serial dilutions of cDNA for each gene
211 were used to generate standard curve analysis and only experiments with $R^2 > 0.985$ were
212 included. CT measurements, calculated by $2^{-\Delta Ct}$ method [38], were used for mRNA expression.
213 Human 18S ribosomal RNA was used as a housekeeping gene for data normalisation.

214

215 *Placental triglyceride and thiobarbituric active reactive substance (TBARS) content*

216 Total lipid extraction used an adapted Folch method and the triglyceride concentration,
217 determined spectrophotometrically (Radox Laboratories Ltd, Crumlin, UK). TBARS was
218 determined as described by Mistry et al [39].

219

220 **Statistical analysis**

221 These were performed using IBM SPSS v20.0 statistical software for Windows (IBM Corp.
222 Armonk, NY, USA). To assess the data for normality, a Kolmogorov–Smirnov test was
223 performed, where a p value > 0.05 indicated normally distribution. Thereafter, appropriate
224 parametric, or non-parametric, tests were used to analyse the effects of maternal overweight and
225 obesity as follows: 1) anthropometrical and physiological comparisons between comparable
226 groups of mothers, placentas and newborns were made using a Students t-test between relevant
227 groups; 2) comparisons of gene expression were determined by using Mann-Whitney test.
228 Categorical data were analysed using Chi-square test of independence. The study was not
229 designed to look at the effect of fetal gender on the placenta. Continuous data presented are
230 expressed as mean average with their standard errors (SEM), with p value < 0.05 deemed to
231 represent statistical significance.

232

233 **Results**

234 *Maternal characteristics, pregnancy outcome, placental composition and metabolic status*

235 Obese women with GDM were older, more likely to be unemployed and have lower educational
236 attainment. Women with obesity gained less weight up to 34 weeks gestation compared to those
237 of normal weight and glucose tolerance (Table 1). In particular obese women with GDM gained
238 significantly less weight than the 2009 IOM guidelines for their BMI group (Chi square test,
239 $p=0.04$) and reflected their lower total energy and carbohydrate intake (Table 2). They also had
240 a lower lipid intake primarily as a consequence of decreased saturated fatty acid consumption.
241 The importance of IOM classified GWG [34] on birth weight was reflected in the trend for
242 obese women to deliver bigger infants when gaining more weight than recommended (Table 1).

243

244 A majority of women gave birth normally at term, with obese women with GDM having a
245 greater risk of preterm delivery and Caesarean section (Table 1). Although estimated fetal
246 weight at 34 gestational weeks was higher when GDM was accompanied by obesity, size and
247 weight at birth were not different between these groups (Table 3). The increased fetal weight at
248 late gestation is likely to reflect the higher preterm and caesarean section delivery rate for obese
249 women with GDM (Table 1). However, although maternal obesity alone did not affect size at
250 birth, women who were obese with normal glucose tolerance had increased placental weight and
251 LGA infants.

252

253 Close to delivery, maternal blood glucose was elevated in women with GDM irrespective of
254 BMI (Table 4). Triglyceride concentrations and monocyte counts were similar between groups
255 but monocyte count was higher in the cord blood of obese women with normal glucose
256 tolerance. Serum leptin concentrations at delivery were elevated in obese compared to normal
257 weight mothers and their offspring. Placental triglyceride content was raised in obese women
258 with GDM with no difference in TBARS.

259

260 *Maternal body weight, GDM and placental markers of energy homeostasis, cell growth and*
261 *endocrine sensitivity*

262 Maternal obesity was accompanied with reduced placental gene expression for *mTOR* (Table 5),
263 whilst upstream (i.e. *Akt*) and downstream (i.e. *p70S6KB1*) signalling molecules for *mTOR* were
264 unaffected. Placental mRNA abundance for *p70S6KB1* was increased when obesity was
265 accompanied by GDM. In addition, GDM was associated with reduced placental gene
266 expression for *AMPK* irrespective of BMI. Increased placental leptin gene expression in normal
267 weight women with GDM, was reversed when GDM was accompanied by obesity. There were
268 no differences in *LEPR* gene expression between groups. Markers of oxidative stress i.e. *SIRT1*
269 and *UCP2* were up-regulated in overweight and obese women, not by GDM. Placental gene
270 expression for *glucocorticoid receptor (GR α)* increased with maternal GDM but was not
271 affected by obesity, and no differences were apparent for inflammatory markers *PPAR γ* and
272 *TLR4*, or indices of insulin action i.e. *IGF1R* or *IRS1*. There was no evidence of any effect of
273 gestational age, mode of delivery or insulin administration on any of these outcomes.

274 **Discussion**

275 Our major finding is the differential effects of perturbations in energy homeostasis on placental
276 expression of genes regulating placental size, function and endocrine sensitivity with raised BMI
277 and GDM. Maternal obesity, but not GDM, contributed to greater placental weight whereas
278 placental adaptation was demonstrated in markers of energy sensing for both groups. Reduced
279 placental *AMPK* mRNA expression with GDM but not with obesity alone, and suppression of
280 gene expression for *mTOR* with obesity are indicative of complementary control mechanisms.
281 Furthermore, the *mTOR* downstream regulator, *p70S6KB1* was increased by obesity even
282 without GDM. Consequently, as maternal glucose was raised at term, and with GDM, these
283 responses could be mediated by changes in glucose homeostasis [28, 40].

284

285 Surprisingly, placental gene expression for *IRS1* and *IGFR1* were not affected by obesity or
286 GDM, findings that differ with those described by Jansson et al. [41] in a cohort of Swedish
287 women, in which placental activation of *mTOR* was accompanied by enhanced insulin/IGF1
288 signalling with raised BMI. However, there are important demographic differences between
289 studies, as the obese Swedish women had a higher mean BMI and substantially greater GWG
290 than our Spanish women. Therefore, the discrepancy between studies may reflect placental
291 threshold effects in response to excess energy intake [42, 43]. In the overweight and obese
292 PREOBE women studied here, reduced placental *mTOR* gene expression was accompanied with
293 raised *SIRT1* and *UCP2*, suggesting enhanced antioxidant capacity [44]. These findings indicate
294 an adaptive placental response to increased BMI, in line with the physiological role of
295 mitochondria in regulating cellular ATP and AMP concentrations [45]. This could occur through
296 changes in the activity of AMPK, Akt, and mTOR with the former sensing energy depletion
297 [46], and the latter stimulated by raised energy supply [43]. Mitochondria also regulate ROS
298 production and oxidative stress by uncoupling energy supply, with both *AMPK* and *mTOR*
299 modulating oxidative stress through changes in UCP2 [47] and NFkB action [26, 48], thereby

300 promoting pro-inflammatory and pro-oxidative pathways within trophoblast cells. In contrast,
301 mitochondrial replication is dependent on *SIRT1* activity that also determines cell survival and
302 senescence by inhibiting *mTOR* activity [49]. Our findings are, therefore, indicative of a
303 protective or physiological adaptation by the placenta against oxidative stress [49, 50] with
304 raised maternal BMI. This is further supported by the stability of placental TBARS content, a
305 marker of oxidative stress [44], between groups suggesting that the fetus is protected from
306 excess ROS. These responses were accompanied by similar expression of placental genes
307 involved in inflammatory responses, i.e. *PPAR γ* [30] and *TLR4* [27], suggesting inflammation
308 was not directly promoted with raised BMI [30].

309

310 Although there were no differences in maternal triglyceride concentrations, obesity with GDM
311 lead to placental triglyceride accumulation, that [51] has been shown to be correlated with fetal
312 adiposity [52] reflected in the increase in LGA infants with maternal obesity. Increased
313 placental triglyceride storage with GDM was accompanied by up-regulation of placental *GR α*
314 that has been shown in an ovine model on nutritional manipulation of placental growth to follow
315 changes in placental mass with gestation [53].

316

317 As expected, maternal obesity was associated with higher plasma leptin irrespective of GDM
318 although whether this leads to a direct inhibitory effect on food intake [54] as reported by these
319 women or reflects maternal metabolism complicated by leptin resistance [55] is uncertain.

320 Although the placenta is a source of plasma leptin [56], which can be stimulated by obesity and
321 GDM [57, 58], we did not observe differences in leptin gene expression, suggesting that
322 adipocytes, rather than the placenta, are the main origin of differences in plasma leptin [59]. An
323 alternative explanation is that there are changes in leptin turnover or that leptin regulated its own
324 expression within the placenta through a mechanism involving the suppression of AMPK [60].
325 Effects on placental leptin expression through the action of glucocorticoids has also been

326 described [61], and is compatible with our observations of an increase in placental GR α
327 suggesting a local inflammatory response within the placenta of obese gestational diabetic
328 women [62, 63].
329
330 Plasma leptin concentrations were raised in cord blood of infants born to obese and obese GDM
331 mothers. This could reflect increased transplacental substrate supply from raised maternal
332 plasma glucose in these women acting through fetal insulin to then promote fetal fat deposition
333 [11, 64]. An enhanced glucose-insulin pathway can promote offspring adiposity [11], whilst the
334 adipokine leptin stimulates cell proliferation by inducing the IRS1/MAPK pathway in a glucose-
335 dependent manner [65]. Furthermore, whilst fetal hyperleptinemia can contribute to induce
336 leptin resistance by chronic activation of leptin receptors in the fetus [66], it is not known
337 whether hypothalamic leptin targets are responsive before birth or whether neonatal leptin
338 resistance leads to long-term adverse consequences. Enhanced circulating leptin in obese
339 women was associated with higher leptin and monocyte concentrations in cord blood. In
340 addition to its potential role in newborn adiposity [64, 67], growing evidence has linked leptin
341 with the maturation of the hypothalamus [68] and the fetal and neonatal immune system [69],
342 leading to impaired immune responses [59]. As part of the PREOBE follow-up further studies
343 are exploring potential long term implications of obesity and diabetes in offspring
344 neurodevelopment through functional measurements. This will enable a more direct assessment
345 of any impact on differences in leptin surge between infants born into the study and their
346 subsequent brain development. Increased pro-inflammatory cytokine expression, including
347 TNF α and IL6, and/or enhanced circulating monocyte chemo-attractant protein (MCP)1
348 concentrations in obese women may account for raised monocytes concentrations in cord blood
349 of their infants [70]. Higher plasma MCP1 [71] has been implicated in monocyte recruitment
350 into adipose tissue of newborns from obese individuals [70] and ultimately produce pro-

351 inflammatory cytokines, contributing to a state of insulin resistance and low grade
352 inflammation.

353

354 As the relative risk of obese and GDM women producing a LGA infant is substantial [11, 14,
355 72], one strategy to prevent this outcome [73] is through healthier food choices [74]. In our
356 study, the first line of treatment of GDM was through nutrition and lifestyle advice in maternity
357 welfare clinics. These reinforced local secular food preferences of Spanish women of primarily
358 Hispanic European white origin (95-98%) for a Mediterranean diet rich in polyunsaturated fatty
359 acids, fruit and vegetables [75, 76] which contrast with those of Northern European and
360 American women recruited in previous studies [77, 78]. However, although there was no
361 difference in mean birth weight in our study, maternal obesity was associated with a higher
362 incidence of LGA infants despite lower self-reported energy and macronutrient intakes. The
363 latter may reflect recall bias as women with increased BMI do not always accurately report their
364 food intake [79, 80]. Alternatively nutrient supply to the fetus of obese woman may be more
365 dependent on existing maternal nutrient stores and current metabolic state [81] than daily
366 intakes. This is supported by raised plasma glucose concentrations even in those obese women
367 who were not diagnosed with GDM. Furthermore, the dietary advice given to these women
368 despite being lowering GWG, did not reduce the incidence of LGA infants, although it is
369 acknowledged that the study was not powered to directly assess such an outcome.

370

371 In conclusion, placental gene expression is sensitive to both maternal BMI and GDM which
372 impacts on both placental triglyceride content and energy sensing. These adaptations could
373 modulate maternal and fetal glucose homeostasis and thus prevent some of the potential adverse
374 consequences on fetal growth and body composition.

375

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Table 1: Socio-demographic characteristics and birth weights of all participants, with and without gestational diabetes: normal weight (N), overweight (OW), obese (O), gestational diabetic normal weight (GDMN) and gestational diabetic obese (GDMO) pregnant women.

Maternal characteristics	N (n=59)	OW (n=29)	O (n=22)	GDMN (n=14)	GDMO (n=11)
Age at delivery (years)	30.4±4.5	30.9±7.2	29.0±4.7	33.1±4.1*	34.7±4.3**
Unemployed (%)	32.2	28.6	38.1	14.3	66.7*
Higher education (University) (%)	42.4	42.9	22.7	42.8	10
Smoking during pregnancy (%)	12.1	25*	9.1	0	0
Primiparous (%)	58.6	46.4	63.6	57.1	60
Height (cm)	162.9±5.7	162.5±6.4	162.7±6.2	159.3±3.9	160.5±6.0
Pre BMI (kg/m ²)	21.8±1.8	27.8±2.2***	32.5±2.6***	22.4±1.8	35.5±4.9***
BMI at 34 weeks (kg/m ²)	26.6±2.6	31.3±2.4***	35.4±2.4***	25.9±2.6	36.4±4.1***
GWG 0-34 weeks (kg) [32]	12.6±4.3	9.9±4.6**	7.3±5.1***	9.0±5.6**	2.2±7.8***
LGWG (kg & % of women in BMI category: n=15;5;8;8;6 resp.)	7.3±1.9 (25%)	2.7±1.9*** (18%)	2.5±1.7*** (36%)	5.2±3.5* (57%)	-3.4±5.0*** (55%)
AGWG (kg & % of women in BMI category: n=23;9;5;3;2 resp.)	12.1±1.1 (39%)	8.2±1.2*** (32%)	5.9±0.9*** (23%)	11.4±1.0 (22%)	5.2±0.6*** (18%)
HGWG ((kg & % of women in BMI category: n=21;14;9;3;3 resp.)	17.0±2.9 (36%)	13.6±2.4** (50%)	12.3±3.4** (41%)	17.0±1.1 (21%)	11.3±3.2** (27%)
BW for each GWG 0-34 weeks (g) [32]					
LGWG (n=15;5;8;8;6 resp.)	3410±116	3496±241	3253±158	3307±151	3373±221
AGWG (n=23;9;5;3;2 resp.)	3160±61	2870±109	3318±274	3493±334	3090±350
HGWG (n=21;14;9;3;3 resp.)	3348±101	3475±117	3707±156.	3433±98	3716±301
No. of Caesarean delivery (%)	12.3	25.9	38.1	25	50*
Preterm delivery (< 37gw) (%)	3.4	3.4	9.1	14.3	27.3*
Male new born (%)	52.5	39.3	61.9	57.1	72.7

601 Values are means ± SD or categorical data as appropriate; n: number of women per group; gw: gestational weeks.

602 Pre: pregestational; BMI: body mass index;

603 GWG: gestational weight gain during the first 34 gestational weeks based on 2009 IOM guidelines for each category [34]: LGWG
604 - gestational weight gained (kg) classified as low: <9.8 kg for normal weight, <5.9 kg for overweight and <4.2 kg for obese
605 women; AGWG - gestational weight gain (kg) classified as adequate: 9.8-13.6 kg for normal weight, 5.9-9.8 kg for overweight
606 and 4.2-7.6 kg for obese women; HGWG - gestational weight gain (kg) classified as high: >13.6 kg for normal weight, >9.8 kg for
607 overweight and >7.6 kg for obese women.

608 Statistical differences: *p<0.05, **p<0.01 ***p<0.001 compared to normal weight group (Chi-square test or t-independent test
609 for continuous variables; chi-square test for categorical variables).

610 **Table 2: Maternal energy and nutrient intake: normal weight (N), overweight (OW), obese (O), gestational diabetic normal**
 611 **weight (GDMN) and gestational diabetic obese (GDMO) pregnant women.**

Maternal dietary intake	N (n=37)	OW (n=15)	O (n=8)	GDMN (n=11)	GDMO (n=6)
Energy (kcal)	2155±339	2114±784	1831±560*	1879±379*	1656±348**
Total carbohydrates (g)	237±54	217±63	189±69*	187±31**	173±46**
Total proteins (g)	83.9±17.5	84.5±28.4	74.8±1.2	84.4±23.0	74.8±11.9
Total lipids (g)	90.5±19.4	95.6±54.2	86.5±26.4	81.7±27.4	68.7±14.7*
SFA (g)	33.8±8.3	36.6±25.9	30.3±8.9	28.3±13.8	21.4±5.6**
MUFA (g)	36.2±9.6	38.3±19.2	35.1±14.5	36.5±12.1	31.8±10.1
PUFA (g)	12.8±4.3	12.5±7.2	13.7±4.6	10.1±2.0	9.7±2.7

612 Values are means ± SD; n: number of women per group;

613 SFA: saturated fatty acid; MUFA: monounsaturated fatty acid; PUFA: polyunsaturated fatty acid.

614 Statistical differences: *p<0.05, **p<0.01 compared to normal weight group (t-independent test for continuous variables).

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616 **Table 3: Anthropometric and clinical characteristics of infants born to mothers with and without gestational diabetes:**
 617 **normal weight mother (N), overweight mother (OW), obese mother (O), gestational diabetic normal weight mother (GDMN)**
 618 **and gestational diabetic obese (GDO) mother.**

Infant characteristics	N (n=59)	OW (n=29)	O (n=22)	GDMN (n=14)	GDMO (n=11)
Estimated fetal weight at 34 weeks of gestation (g)	2363±183	2345±183	2393±383	2467±380	2541±501*
Placental weight (g)	469±120	495±135	531±114*	498±134	476±93
Placental to birth weight ratio	0.143±0.031	0.157±0.046	0.158±0.041	0.147±0.035	0.139±0.017
Gestational age (weeks)	39.2±1.0	39.4±1.6	39.3±1.7	39.3±1.3	38.8±1.3
Newborn length (cm)	50.2±1.8	50.5±1.5	50.6±2.7	50.6±1.7	50.9±3.4
Newborn weight (g)	3292±410	3230±587	3454±549	3374±402	3415±549
SGA (n) (%)	4 (6.8)	3 (10.3)	1 (4.5)	1 (7.1)	2 (18.2)
AGA (n) (%)	52 (88.1)	24 (82.8)	16 (72.8)	12 (85.8)	7 (63.6)
LGA (n) (%)	3 (5.1)	2 (6.9)	5 (22.7)*	1 (7.1)	2 (18.2)
Ponderal index (g/cm ³ *100)	2.62±0.27	2.56±0.49	2.58±0.28	2.60±0.34	2.60±0.42

619 Anthropometric characteristics of the fetus were estimated by using ultrasound scan at 34 gestational weeks.

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621 Values are means ± S.D.; n: number of women per group;

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623 SGA: small for gestational age (birthweight population centile < 10%); AGA: average for gestational age (10% < birthweight
 624 population centile < 90%); LGA: large for gestational age (birthweight population centile > 90%).

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626 Statistical differences: *p<0.05 compared to normal weight group (t-independent test for continuous variables; chi-square test
 627 for categorical variables).

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629 **Table 4: Maternal, placental and cord blood metabolic characteristics: normal weight (N), overweight (OW), obese (O),**
 630 **gestational diabetic normal weight (GDMN) and gestational diabetic obese (GDMO) pregnant women.**

Maternal blood at term	N (n=59)	OW (n=29)	O (n=22)	GDMN (n=14)	GDMO (n=11)
Glucose (mmol/L)	4.3±1.3	4.6±1.3	5.3±2.3*	6.0±2.2***	6.1±1.9***
Triglyceride (mmol/L)	11.7±3.9	13.2±4.2	12.8±4.3	11.6±3.9	12.3±3.3
Leptin (µg/L)	16.0±13.6	24.2±23.5	33.9±21.9**	21.1±16.4	36.6±19.9**
Monocyte count (x10 ⁹ /L)	0.5±0.2	0.6±0.2	0.6±0.2	0.6±0.3	0.40±0.2
Cord blood ^ψ	N (n=33)	OW (n=18)	O (n=16)	GDN (n=10)	GDO (n=7)
Glucose (mmol/L)	3.8±1.2	3.6±1.2	3.3±1.3	3.9±0.7	4.3±1.0
Triglyceride (mmol/L)	2.7±1.1	2.5±0.8	2.5±1.4	2.5±1.3	2.5±1.2
Leptin (µg/L)	19.7±17.9	32.7±19.6	62.3±69.0*	42.6±31.5*	36.6±19.9
Monocyte count (x10 ⁹ /L)	1.0±0.4	1.1±0.4	1.4±0.6*	1.2±0.8	1.1±0.4
Placental tissue	N (n=59)	OW (n=29)	O (n=22)	GDN (n=14)	GDO (n=11)
Total placental TG (mg/g) per placental weight (g)	19.9±10.0	22.0±10.9	23.9±12.8	19.7±10.6	28.3±16.5*
Relative placental TBARS	1.0±0.3	0.9±0.3	0.8±0.2	1.0±0.3	1.0±0.3

631 Values are means ± SD; n: number of women/group; ^ψ see text for information on missing individuals.

632 TBARS: thiobarbituric acid reactive substances; TG: triglyceride.

633 Statistical differences: *p<0.05, **p<0.01 compared to normal weight group (t-independent test for continuous variables).

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635 **Table 5: Effects of maternal BMI on gene expression markers of energy sensing and balance, oxidative stress and**
636 **inflammation in placenta of normal weight (N), overweight (OW), obese (O), gestational diabetic normal weight (GDMN)**
637 **and gestational diabetic obese (GDMO) pregnant women.**

Pathway	gene	NCBI sequence	Target Gene	N (n=59)	OW (n=29)	O (n=21)	GDMN (n=14)	GDMO (n=11)
Energy sensing		NM_006251	<i>AMPK</i>	1.0±0.1	0.9±0.1	1.0±0.2	0.6±0.1*	0.4±0.1***
		NM_001014432	<i>Akt1</i>	1.0±0.1	1.0±0.2	1.2±0.2	0.8±0.1	0.9±0.1
		NM_004958	<i>mTOR</i>	1.0±0.1	0.7±0.1	0.5±0.1*	0.6±0.1	0.5±0.1
		NM_003161	<i>p70S6KB1</i>	1.0±0.1	1.1±0.2	1.6±0.4	0.7±0.2	1.4±0.2*
Energy balance		NM_000230	<i>LEP</i>	1.0±0.2	1.5±0.5	0.9±0.4	4.1±1.1*	0.8±0.4
		NM_002303	<i>LEPR</i>	1.0±0.2	0.8±0.1	1.1±0.3	0.5±0.0	0.5±0.1
Insulin action		NM_000875	<i>IGF1R</i>	1.0±0.1	1.2±0.1	1.2±0.2	1.0±0.1	0.8±0.1
		NM_005544	<i>IRS1</i>	1.0±0.1	1.1±0.1	1.4±0.2	0.9±0.1	0.8±0.1
Oxidative stress		NM_001033611.1	<i>UCP2</i>	1.0±0.2	1.4±0.2**	1.4±0.2*	1.3±0.4	0.8±0.3
		NM_001142498	<i>SIRT1</i>	1.0±0.1	1.4±0.2*	1.6±0.2**	0.8±0.2	1.5±0.3
Inflammation		NM_015869.4	<i>PPARγ</i>	1.0±0.1	0.9±0.1	0.9±0.1	1.0±0.2	0.9±0.1
		NM_001135930.1	<i>TLR4</i>	1.0±0.1	0.9±0.1	0.9±0.1	0.8±0.1	0.8±0.2
		NM_000176	<i>GRα</i>	1.0±0.1	1.2±0.1	1.0±0.1	1.2±0.1*	1.5±0.2*

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641 Data expressed relative to housekeeping gene (ribosomal 18S RNA), normalised to the control group to give the fold change.
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643 a.u.: arbitrary units; n = number of women/group.

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645 AMPK: AMP-activated protein kinase; Akt: v-akt murine thymoma viral oncogene homolog; mTOR: mammalian target of
646 rapamycin; p70S6KB1: ribosomal protein S6 kinase 70kDa polypeptide; LEP: leptin; LEPR: leptin receptor; IGF1R: insulin growth
647 factor 1 receptor; IRS1: insulin receptor substrate 1; UCP: uncoupling protein; SIRT: sirtuin; PPARγ: peroxisome proliferator-
648 activated receptor gamma; TLR: toll like receptor; GRα: glucocorticoid receptor alpha.

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650 Data are non parametric and represent mean ± S.D. Statistical differences: *p<0.05, **p<0.01, ***p<0.001 compared to normal
651 weight (Mann Whitney test).

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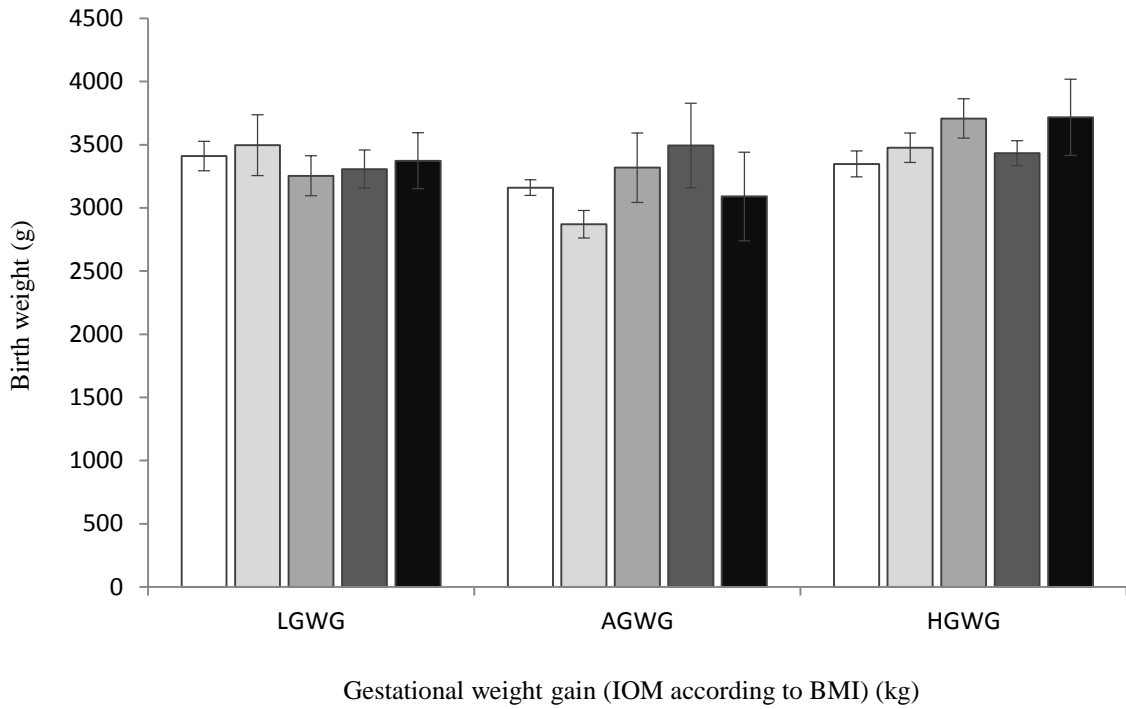


Figure 1: Offspring birth weight trend according to IOM stratification of maternal metabolic status and weight gain during the first 34 gestational weeks.

Normal weight (white bar; n=15;23;21), overweight (light grey bar; n=5;9;14) obese (grey bar; n=8;5;9), normal weight gestational diabetes (dark grey bar; n=8;3;3) and obese gestational diabetes (black bar; n=6;2;3) pregnant women. Values are means \pm SEM; weight gain during the first 34 gestational weeks is based on 2009 IOM guidelines for each BMI category^[34]: LGWG: low gestational weight gain (<9.8 kg for normal weight, <5.9 kg for overweight and <4.2 kg for obese women); AGWG: adequate gestational weight gain (9.8-13.6 kg for normal weight, 5.9-9.8 kg for overweight and 4.2-7.6 kg for obese women); HGWG: high gestational weight gain (>13.6 kg for normal weight, >9.8 kg for overweight and >7.6 kg for obese women).