Programming of glucose-insulin metabolism in adult sheep after maternal undernutrition


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Gardner, D. S., K. Tingey, B. W. M. Van Bon, S. E. Ozanne, V. Wilson, J. Dandrea, D. H. Keisler, T. Stephenson, and M. E. Symonds. Programming of glucose-insulin metabolism in adult sheep after maternal undernutrition. Am J Physiol Regul Integr Comp Physiol 289: R947–R954, 2005. First published June 16, 2005; doi:10.1152/ajpregu.00120.2005.—The present study examines the effects of late vs. early gestation undernutrition on adult glucose-insulin homeostasis in sheep and investigates whether the lower birth weight of twins alters glucose-insulin handling in adult life. Pregnant sheep were fed to requirement (100% intake) from day 0 of gestation to term (~147 days of gestation (dGA)), control singles (CS) n = 5; control twins (CT) n = 5 or to 50% requirement from days 0–30 dGA [nutrient restricted during early gestation (NRE); n = 5] or day 110-term [NR during late nutrition (NRL); n = 4]. At all other times, NR sheep received 100% intake. All sheep lambed naturally; offspring were weaned at 10 wk and were reared on pasture until 1 yr of age. At this time, indwelling catheters were inserted, and 2–4 days later, basal metabolic and endocrine status and responses to an intravenous glucose tolerance test (IVGTT) and feeding were assessed. Adipose and skeletal muscle were then sampled after humane euthanasia and were analyzed for expression of insulin-signaling proteins and GLUT4. Between groups, birth weight of singletons was similar and increased relative to twins. At 1 yr of age, weights were similar between groups. The areas under the curve for glucose and insulin during the IVGTT were greater in NRL vs. other groups, indicating glucose intolerance. This was associated with reduced adipose, but not muscle, GLUT4, and increased adipose tissue mass. Adult glucose-insulin homeostasis in sheep was unaffected by fetal number. In conclusion, prenatal undernutrition, specifically during late gestation, affects adult offspring intermediary metabolism, and, in particular, glucose-insulin homeostasis.

In middle age has its origins early in life, specifically during fetal and early postnatal development (5, 22). Some of the most compelling data regarding the developmental programming of obesity and related metabolic disorders have come from retrospective studies examining a proportion of the Dutch population exposed to famine during World War II—the Dutch Hunger Winter Famine. Very early on, it was realized that the period of famine exposure (3–4 mo) produced different outcomes in the adult offspring, depending on when the famine conditions were endured during pregnancy (36). For example, exposure during the first trimester of pregnancy only, gave rise to offspring with increased risk of coronary heart disease (odds ratio, 8.8%) compared with exposure during midgestation, late gestation, or nonexposed individuals [midgestation, 0.9%; late gestation, 2.5%; nonexposed, 3.0% (37)]. In contrast, famine exposure during late gestation only tended to impact on intermediary metabolism, in particular, glucose-insulin homeostasis (35). Such effects can be readily explained when one considers the chronological development of system/growth in the fetus; cardiovascular growth has a priority early in gestation, with adipose and muscle growth (highly insulin-sensitive tissues) occurring relatively late.

Animal studies have, in part, replicated these original hypotheses and shown that undernutrition confined to early gestation may impact on adult cardiovascular function (19, 27), whereas undernutrition throughout gestation and/or lactation produces both hypertensive (26) and insulin-resistant (16, 32) offspring. Surprisingly, to date, no animal study has investigated adult glucocorticoid-insulin metabolism after prenatal undernutrition confined to late gestation alone, nor to have directly compared the effect of early vs. late undernutrition on aspects of glucose-insulin intermediary metabolism in the resultant adult offspring. The current study was therefore established to directly address the hypothesis that late, rather than early, gestation undernutrition has the greatest impact on adult glucocorticoid-insulin homeostasis.

In addition, one of the many criticisms leveled at the developmental origins of adult disease, hypothesis concerns the incidence of noncommunicable disease in twins that are generally of lower weight at birth (7) but do not seem to exhibit a higher risk for either cardiovascular or metabolic disease (4, 8, 11, 33).
catheters were inserted under anesthesia (1–2% halothane in 50:50 
O2/N2O) with appropriate antibiotic [15 mg/kg im amoxicillin
(Duphamox); Fort Dodge Animal Health, Southampton, UK] and
analgescic coverage [1 mg/kg flunixin meglumine (Finadyne); Shering-
Plough, Kenilworth, UK] postoperatively. Catheters were kept patent
by daily flushing with heparinized saline (50 IU heparin/ml).

Experimental Protocols

No experiment was performed until after 2–4 days postoperative
recovery. The investigator was blinded to the dietary origin of the
sheep before any experiment was performed. In all sheep, two ex-
periments were conducted on separate days.

Experiment 1. METABOLIC RESPONSE TO FEEDING. Both hay and
concentrate were withheld from the sheep for a period of 24 h. The
following morning, two baseline blood samples (2 ml; −10 and −5
min) were taken before feeding and then subsequently at 5, 15, 30, 45,
and 60 min after feed was given at time 0, which coincides with the
rapid switch from a dependence on lipid to carbohydrate metabolism
in ruminants (38). All concentrate feed for all animals was consumed
within 10 min, and blood samples were handled as previously de-
scribed for experiment 1. The combined 45- and 60-min sample was
used for a paired comparison of fastest i.e., mean of baseline samples
(−10 and −5 min) vs. fed state (45- and 60-min sample).

Experiment 2. INTRAVENOUS GLUCOSE TOLERANCE TEST. Both hay
and concentrate were withheld from the sheep for a period of 24 h.
The following morning, two baseline blood samples (2 ml; −10 and −5
min) were collected into chilled heparinized tubes before feeding,
and then subsequently at 2, 5, 10, 15, 20, 30, 40, 60, 80, 120, 150, 180,
210, and 240 min after 0.5 g/kg glucose was administered intrave-
rously (in 0.9% NaCl) within 2 min, through the jugular catheter at
time 0. All blood samples were centrifuged immediately for 5 min at
4°C and, after decanting of the supernatant plasma, were stored at
−20°C for further analysis of plasma glucose, insulin, and nonsteri-
fied fatty acids (NEFA) concentrations. In two sheep, a similar
volume (≈45 ml) of 0.9% NaCl only was administered as a control.
Values for glucose, insulin, and NEFA remained unchanged from
baseline in these sheep (data not shown).

Biochemical Analyses

Plasma concentrations of glucose and NEFA were measured enzy-
merically as described by Symonds et al. (39), whereas plasma
triglycerides were determined with the use of a commercial coloromet-
ric kit (GPO-PAP TR212; Randox Laboratories, Crumlin, Antrim,
UK) adapted for use with 96-well plates. The intra- and interassay
coefficients of variation were 6.4 and 9.6% (n = 10 and 8, respec-
tively). Plasma cortisol and leptin were both determined by radioim-
unoassay as previously described in detail (6, 10). Insulin (50 µ l
plasma) was assayed in duplicate using a commercially available
coated tube assay (INSI-CTK IRMA; Diasorin, Wokingham, UK).
The assay uses human insulin as the reference standard and has a
minimum detection limit of 4.3 pmol/l. The intra-assay variation was
<5%. All samples for insulin were assayed at the same time with
the same standard curve.

Molecular Analyses

At the end of all experimental protocols, the 1-yr-old sheep were
humanely euthanized with a lethal dose of pentobarbital sodium
(Euthatal; 100 mg/kg). All major organ weights were recorded, and
tissue samples were flash frozen in liquid nitrogen and stored at
−80°C. A portion of perirenal fat and skeletal muscle were also
collected for Western blot analysis as previously described for the rat
(16) but with the following modifications for sheep. The cleared
protein lysates from each animal were standardized to a final concen-
tration of 0.5 mg/ml in Laemmli’s sample buffer and equal amounts
of protein for each animal (10 µg) were loaded onto 10% SDS

Materials and Methods

Animals

All procedures were approved by the local ethics committee of The
University of Nottingham and performed under United Kingdom
Animals (Scientific Procedures) Act, 1986. The general principles of
laboratory animal care were strictly followed (29). Nineteen blue-
faced Leicester cross Swaledale (Mule) sheep from the University of
Nottingham’s commercial flock were used in the study. Sheep were
mated during their natural breeding season without the use of ovarian
stimulants. After mating, sheep were individually housed and ran-
domly assigned to receive one of three diets during pregnancy: 1) 100%
metabolizable energy (ME) requirements as defined by the A

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polycrylamide gels for separation by electrophoresis. The ovine antibodies used in this study were to insulin receptor β-subunit (Irfβ; Santa Cruz sc-711, Autogen Bioclear, UK); phosphatidylinositol 3-kinase (PI3-kinase) p110 β-subunit (p110β; Santa Cruz); PI3-kinase p85 ε-subunit (p85ε; Upstate Biotech); and GLUT4 (Abcam, Cambridge, UK). Secondary antibodies and an antibody-binding (enhanced chemiluminescence) kit were both obtained from Amersham.

**Calculations**

For the intravenous glucose tolerance test (IVGTT) a glucose and insulin curve was plotted. From this plot of the rapid increases and gradual decay of glucose, and subsequently insulin, a value for area under the curve (AUC) was derived using the GraphPad Prism software (Version 3; GraphPad). Variables derived were 1) maximum change in glucose/insulin calculated as the maximum value achieved minus the average baseline value, 2) time to restore fasting baseline glucose/insulin calculated from the intercept of the slopes for the glucose/insulin curves with fasting glucose/insulin concentration, 3) glucose tolerance measured as the area above fasting plasma glucose under the plasma glucose curve (Glucose: min·mmol−1·l−1·t−1), 4) absolute insulin secretion measured as the area above fasting plasma insulin under the plasma insulin curve (Insulin: min·mmol−1·l−1·t−1), 5) relative insulin secretion (nmol·mmol−1·l−1·t−1) calculated as the absolute insulin secretion divided by glucose intolerance, and 6) glucose clearance rate (mmol/min) calculated as (t<sub>broad minus time at peak glucose) divided by the peak glucose achieved (Glucose clearance).

**Statistical Analyses**

All data are expressed as means ± SE unless otherwise stated. Effects of the nutritional group on measures of postnatal growth rates, glucose homeostasis, insulin secretion, and plasma metabolites or hormones were analyzed by univariate general linear models with group and sex as fixed effects using SPSS version 11.5 (SPSS, Chicago, IL). Relationships between pairs of variables were analyzed by Pearson’s correlation across all animals (SPSS version 11.5). Descriptive data for maternal weight, food intake, and birth weight were analyzed by one-way ANOVA (SPSS version 11.5). For a comparison between hormone concentrations in the fed and fasted state, a paired Student’s t-test was employed. For all comparisons, statistical significance was accepted when P < 0.05.

**RESULTS**

**Gestation and Postnatal Growth**

Control sheep consumed from 7.6 ± 0.3 MJ/day at week 1 of pregnancy to 15.8 ± 0.9 control singletons (CS) and 18.9 ± 0.8 control twins (CT) MJ/day at week 21 of pregnancy. NR days 1–30 (NRE) and NR days 110–140 sheep (NRL) consumed 50% of that amount for the first 30 days (3.61 ± 0.11 MJ/day) or days 110–140 (8.67 ± 0.15 MJ/day), respectively. At all other times, the undernourished groups consumed 100% of control intake. Birth weights were similar between singleton lambs in all groups (CS, 4.4 ± 0.4 kg; NRE, 4.2 ± 0.3 kg; NRL, 4.9 ± 0.3 kg) but control-fed twin lambs were significantly smaller at birth (CT, 3.4 ± 0.2 kg). There was no difference in the growth rates (current wt – birth wt/time) of lambs in each group during the first 3 mo (0–3 mo), second 3 mo (4–6 mo), and between 7–12 mo: CS, 1.66 ± 0.31, 1.27 ± 0.15, 0.92 ± 0.07 kg/wk; CT, 1.99 ± 0.16, 1.47 ± 0.09, 0.89 ± 0.11; NRE, 1.62 ± 0.18, 1.32 ± 0.08, 0.88 ± 0.09; NRL, 2.62 ± 0.19, 2.01 ± 0.08, and 1.02 ± 0.09, respectively. Rates of growth in all groups slowed over time (P < 0.001). At 1 yr of age, when the metabolic studies were undertaken, there was no difference in body weight between the groups of sheep (CS, 47.1 ± 4.0; CT, 50.2 ± 5.9; NRE, 49.8 ± 4.3; NRL, 58 ± 5 kg).

**Fed vs. Fasted Metabolite and Leptin Concentrations**

The fed and fasted plasma concentrations of metabolites and leptin are given in Table 1. Fasted plasma glucose concentration was similar between groups and values did not change significantly during feeding. In contrast, plasma NEFA significantly decreased from fasted values during feeding in CS (P = 0.01) and NRE (P = 0.002), but no effect was observed in NRL. There was no effect of fasting or feeding on plasma triglycerides or leptin. Fasted values for plasma leptin and triglycerides showed a strong positive correlation with each other (r = 0.74, P < 0.01). Gender had no effect on any metabolite or hormone values as assessed using a univariate general linear model with sex as a fixed influence. Similarly basal plasma cortisol concentration was unaffected by any of the variables assessed (e.g., CS, 31.6 ± 7.5; CT, 29.5 ± 5.5; NRE, 42.7 ± 7.0; NRL, 35.7 ± 12.7 nM).

**IVGTT**

Baseline plasma glucose (CS, 5.11 ± 0.43; CT, 5.35 ± 0.35; NRE, 5.42 ± 0.52; NRL, 5.25 ± 0.66 mM) and insulin (CS, 100 ± 18; CT, 113 ± 25; NRE, 76 ± 8.3; NRL, 148 ± 41 nM) concentrations were similar between dietary groups. Infusion of an intravenous glucose load significantly increased plasma glucose (Fig. 1) and insulin (Fig. 2) concentration in all groups of sheep with the peak values achieved being similar (CS, 21.7 ± 1.0 and 1.146 ± 196; CT, 21.7 ± 0.5 and 990 ± 102; NRE, 22.6 ± 0.5 and 803 ± 132; NRL, 24.5 ± 1.8 mM and 1,165 ± 129 nM for glucose and insulin, respectively). There was no significant effect of diet or fetal number on the maximal change in plasma glucose, or insulin, or time to restore plasma glucose to baseline; however, the time taken for plasma insulin to return to baseline after the IVGTT was significantly increased in NRL relative to CS (Table 2, P = 0.02). In addition, glucose tolerance as defined by glucose AUC was decreased

Table 1. Fed and fasted plasma concentrations of glucose, nonesterified fatty acids (NEFA), triglycerides, and leptin in control and nutrient-restricted sheep at 1 yr of age

<table>
<thead>
<tr>
<th></th>
<th>CS</th>
<th>CT</th>
<th>NRE</th>
<th>NRL</th>
</tr>
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<tbody>
<tr>
<td>n</td>
<td>5</td>
<td>5</td>
<td>5</td>
<td>4</td>
</tr>
<tr>
<td>Glucose, mM</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fasted</td>
<td>5.59±0.55</td>
<td>4.88±0.17</td>
<td>5.42±0.54</td>
<td>5.25±0.51</td>
</tr>
<tr>
<td>Fed</td>
<td>4.94±0.60</td>
<td>6.08±0.91</td>
<td>7.28±1.28</td>
<td>7.14±1.82</td>
</tr>
<tr>
<td>NEFA, mM</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fasted</td>
<td>0.63±0.13</td>
<td>0.45±0.11</td>
<td>0.95±0.12</td>
<td>0.45±0.11</td>
</tr>
<tr>
<td>Fed</td>
<td>0.12±0.01</td>
<td>0.15±0.04</td>
<td>0.13±0.04</td>
<td>0.42±0.19</td>
</tr>
<tr>
<td>Triglycerides, mM</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fasted</td>
<td>0.18±0.02</td>
<td>0.20±0.04</td>
<td>0.23±0.03</td>
<td>0.31±0.07</td>
</tr>
<tr>
<td>Fed</td>
<td>0.21±0.02</td>
<td>0.25±0.02</td>
<td>0.24±0.03</td>
<td>0.26±0.02</td>
</tr>
<tr>
<td>Leptin, ng/ml</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fasted</td>
<td>0.99±0.10</td>
<td>1.36±0.28</td>
<td>0.93±0.19</td>
<td>3.27±1.52</td>
</tr>
<tr>
<td>Fed</td>
<td>2.60±0.73</td>
<td>1.84±0.54</td>
<td>2.20±1.23</td>
<td>1.49±0.52</td>
</tr>
</tbody>
</table>

Values are means ± SE for control singlets (CS), control twins (CT), early nutrient-restricted single (NRE), and late nutrient-restricted single (NRL) sheep. Fasted concentrations reflect the mean values for 2 baseline samples, and the fed samples are the mean values for 2 samples collected between 45–60 min after feeding. Data were statistically tested using Student’s paired t-test and differences are: *P < 0.05 fasted vs. fed.

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Interestigly, the expression of p110α was significantly increased during baseline at NaCl. Blood samples were taken and late nutrient-restricted single (NRL) sheep. Glucose (0.5 g/kg in 0.9% NaCl) was administered as a bolus over 2 min. Protein expression in the muscle. There was, however, a significant increase in the protein for the p85α of PI3-kinase (CS, 20,833 AU; NRL, 24,074 AU) protein expression, p85α, and GLUT4 expression (r = -0.48, P = 0.049) predicted worse glucose tolerance (i.e., increased GluAUC of the offspring). However, when current weight was controlled for in a partial correlation analysis, the relationship with GLUT4 became nonsignificant (r = -0.32, P = 0.22), and the significance of the relationships with p110α and Irβ were weakened (r = 0.56, P = 0.06; r = 0.44, P = 0.07, respectively). The positive relationship between muscle Irβ and GluAUC remained after controlling for current weight. Fasting plasma insulin and InsAUC correlated negatively with muscle (r = -0.56, P = 0.01) and adipose GLUT4 expression (r = -0.61, P = 0.01), respectively. Allowing for current weight weakened the relationship between InsAUC and GLUT4 (r = -0.38, P = 0.001) but had no effect on the relationship between fasting plasma insulin and muscle GLUT4. There was a strong positive correlation between InsAUC and p110α expression in adipose tissue (r = 0.71, P < 0.001), which remained after controlling for current weight.

**Sheep biometry at 1 yr of age.** There were no differences between dietary groups in the weights of any organ measured. However, the proportion of adipose tissue in central (omental) and peripheral (perirenal) depots was significantly (P = 0.01) greater in NRL relative to other groups of sheep (Table 3).

**DISCUSSION**

In this study, it has been shown that undernutrition confined to late gestation, during the period of maximal fetal growth, impacts negatively on the subsequent young adults’ glucose-insulin homeostasis i.e., they show evidence of glucose intolerance (increased GluAUC and insulin resistance (increased InsAUC). Importantly, no major change in glucose tolerance and/or insulin sensitivity was observed in the young adult offspring of sheep undernourished for an equivalent period during very early gestation. The underlying mechanism for disturbed glucose-insulin homeostasis in these late gestation undernourished offspring appears to be altered tissue (specifically, adipose rather than muscle) glucose uptake, through reduced cellular expression of GLUT4. In addition, the study has clearly shown that the lower birth weight of twins has no

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**Fig. 1.** Plasma glucose after an intravenous glucose tolerance test (IVGTT) in control and nutrient-restricted sheep at 1 yr of age. Values are means ± SE for control singles (CS), control twins (CT), early nutrient-restricted single (NRE) and late nutrient-restricted single (NRL) sheep. Glucose (0.5 g/kg in 0.9% NaCl) was administered as a bolus over 2 min. Blood samples were taken during baseline at −10 and −5 min and subsequently at 2, 5, 10, 15, 20, 30, 40, 60, 80, 120, 150, 180, 210, and 240 min after the glucose bolus.

**Fig. 2.** Plasma insulin after an IVGTT in control and nutrient-restricted sheep at 1 yr of age. Values are means ± SE for CS, CT, NRE, and NRL sheep. Glucose (0.5 g/kg in 0.9% NaCl) was administered as a bolus over 2 min. Blood samples were taken during baseline at −10 and −5 min and subsequently at 2, 5, 10, 15, 20, 30, 40, 60, 80, 120, 150, 180, 210, and 240 min after the glucose bolus.
bearing on their future metabolic competence, at least in terms of their glucose-insulin handling.

Retrospective data from the Dutch Hunger Winter Famine have shown quite clearly that specific periods of famine exposure may impact on specific physiological control systems in adult life. For example, exposure during early gestation influenced the cardiovascular system, clinically reflected as an increased risk of coronary heart disease (37); whereas exposure during late gestation tended to affect intermediary metabolism, in particular, glucose-insulin homeostasis, and was clinically reflected as an increased risk of type 2 diabetes (35). Subsequent animal studies established to model these epidemiological data have recognized early programming of the metabolic syndrome and have used a similar degree of undernutrition i.e., 50% normal intake (protein and/or energy) (for reviews see Refs. 3 and 30). However, in general, the diet in these studies has been restricted either throughout gestation or throughout gestation and lactation, thereby covering the entire period of embryonic, fetal, and postnatal growth. This is important when one further considers the relatively greater nutritive demands of the small compared with large animal species used in these studies (28). Nevertheless, in larger farm animal species, programming of cardiovascular and metabolic function has been described, although the magnitude of adaptation is attenuated compared with that described in small laboratory species. These studies have focused on the impact of either birth size (33, 34) or excess glucocorticoid exposure (12) rather than the plane of maternal nutrition. Two recent studies have now directly examined the effect of a specific period of maternal undernutrition (early/early-mid gestation) on adult physiological function in a larger species and have shown programming of cardiovascular control (19, 21) in common with the observations after early famine exposure in the Dutch Hunger Winter Famine. In the current study, these observations are extended to include a further group of late-gestation undernourished animals for comparison and to an examination of metabolic function in the young adult offspring of a large animal species, two hitherto unknown endpoints.

When the current study is considered alongside a contemporaneous study (19), it is clear that late-gestation undernutrition has a more profound impact on intermediary metabolism, whereas early-gestation undernutrition affects cardiovascular control (19), two observations that replicate the Dutch Famine data. Furthermore, we have shown that despite the lower birth weight and relatively increased early growth rate of twins, their intermediary metabolism (current study) and cardiovascular control (19) are not compromised. Thus developmental programming is not related to birth weight per se, which in multiple births is largely due to maternal constraint but is apparent only when the intra/extraterine environment is specifically challenged or inappropriate at that point in time e.g., through nutritional imbalance (under/overnutrition). Twins appear to have specific adaptations to accommodate their compromised prenatal environment (13, 18), but these adaptations appear to confer no further disadvantage in later life. In contrast, brief periods of maternal undernutrition elicit changes

Table 2. Derived values from intravenous glucose tolerance test in control and nutrient-restricted sheep at 1 yr of age

<table>
<thead>
<tr>
<th></th>
<th>CS</th>
<th>CT</th>
<th>NRE</th>
<th>NRL</th>
</tr>
</thead>
<tbody>
<tr>
<td>Peak change in glucose, mM</td>
<td>16.6±1.2</td>
<td>16.4±0.7</td>
<td>17.2±0.3</td>
<td>19.3±1.4</td>
</tr>
<tr>
<td>Peak change in insulin, nM</td>
<td>1045±194</td>
<td>876±112</td>
<td>726±131</td>
<td>1134±36</td>
</tr>
<tr>
<td>Time to baseline (glucose), min</td>
<td>175±23</td>
<td>137±14</td>
<td>163±20</td>
<td>200±10</td>
</tr>
<tr>
<td>Time to baseline (insulin), min</td>
<td>137±7</td>
<td>172±18</td>
<td>149±20</td>
<td>225±9</td>
</tr>
<tr>
<td>Glucose tolerance, (min:mmol−1·l−1)</td>
<td>1111±54a</td>
<td>964±109a</td>
<td>1340±206b</td>
<td>1619±170b</td>
</tr>
<tr>
<td>Insulin secretion: (min:mmol−1·l−1)</td>
<td>69.5±9.0a</td>
<td>69.5±6.9a</td>
<td>53.0±10.6a</td>
<td>115.6±22.0b</td>
</tr>
<tr>
<td>Relative insulin secretion: (nmol:mmol−1·l−1)</td>
<td>63.2±9.2a</td>
<td>73.6±6.2b</td>
<td>39.2±4.3a</td>
<td>80.6±15.9b</td>
</tr>
<tr>
<td>Glucose clearance: (mmol−1·min−1)</td>
<td>10.6±1.3</td>
<td>8.1±1.2</td>
<td>8.4±0.6</td>
<td>10.3±0.9</td>
</tr>
</tbody>
</table>

Values are means ± SE for CS, CT, NRE, and NRL sheep. Values were derived as described in MATERIALS AND METHODS. Values arranged in individual rows with differing superscript letters are statistically different at P < 0.05 (1-way ANOVA).
in the developmental pattern and have further consequences as the individual ages.

The reduced glucose tolerance in NRL vs. other groups in the current study is very clear despite the relatively low number in this group. This glucose intolerance is related to insulin resistance and not insulin deficiency as indicated by the increase in adiposity, alteration in expression of insulin-signaling components, and reduced glucose tolerance of NRL in the current study, it has previously been shown (16) that the subcellular insulin signaling proteins downstream of the receptor may be programmed before development of increased adiposity and glucose intolerance. Thus it is conceivable that late-gestation undernutrition provokes a compensatory increase in adipose insulin receptor in response to the reduction in maternal and/or fetal glucose concentrations (50% reduction with NRL dietary regimen, see Ref. 14). Although the NRL regimen may reduce term fetal adipose mass (40), irreversibly increased adipose insulin receptor may sustain a postnatal anabolic drive toward increased adipose deposition. The underlying mechanism for reduced GLUT4 following late-gestation undernutrition is unclear; however, it has been shown that hyperinsulinemia during late gestation in the ovine fetus produces tissue-specific (i.e., in insulin sensitive tissues such as myocardium, adipose) insulin resistance, and reduced GLUT4 expression (2).

### Table 3. Sheep biometry at 1 yr of age

<table>
<thead>
<tr>
<th>Tissue Mass</th>
<th>CS</th>
<th>CT</th>
<th>NRE</th>
<th>NRL</th>
</tr>
</thead>
<tbody>
<tr>
<td>Brain weight</td>
<td>96.7±3.5</td>
<td>99.7±2.8</td>
<td>93.5±1.5</td>
<td>94.3±3.5</td>
</tr>
<tr>
<td>Heart weight</td>
<td>178±17</td>
<td>195±21</td>
<td>213±14</td>
<td>225±22</td>
</tr>
<tr>
<td>Liver weight</td>
<td>633±32</td>
<td>688±107</td>
<td>669±99</td>
<td>672±72</td>
</tr>
<tr>
<td>Kidney weight</td>
<td>111±9</td>
<td>116±6</td>
<td>119±6</td>
<td>172±35</td>
</tr>
<tr>
<td>Adrenal weight</td>
<td>2.90±0.64</td>
<td>3.17±0.40</td>
<td>3.17±0.20</td>
<td>2.66±0.66</td>
</tr>
<tr>
<td>Spleen weight</td>
<td>111±19</td>
<td>110±10</td>
<td>92±15</td>
<td>145±33</td>
</tr>
<tr>
<td>Lung weight</td>
<td>475±19</td>
<td>476±21</td>
<td>464±22</td>
<td>478±40</td>
</tr>
<tr>
<td>Pancreas weight</td>
<td>42.5±3.5</td>
<td>49.0±10.1</td>
<td>43.3±7.8</td>
<td>29.0±3.4</td>
</tr>
<tr>
<td>Perirenal adipose tissue</td>
<td>104±25a</td>
<td>189±33a</td>
<td>151±27a</td>
<td>361±65b</td>
</tr>
<tr>
<td>Omental adipose tissue</td>
<td>259±96</td>
<td>295±41</td>
<td>273±31</td>
<td>558±132</td>
</tr>
<tr>
<td>Relative fat mass</td>
<td>8.09±1.89</td>
<td>11.24±1.36</td>
<td>9.72±0.60</td>
<td>17.52±4.33a</td>
</tr>
</tbody>
</table>

Values are means ± SE in grams for all weights and tissues, except relative fat mass, which is in grams per kilogram, for (CS, n = 5), CT, (n = 5), NRE and NRL sheep. Values arranged in individual rows with differing superscript letters are significantly different at P < 0.05.

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![Image](http://ajpregu.physiology.org/doi/10.1152/ajpregu.00920.2005)
It is clear, therefore, that the period of exposure to a dietary imbalance during gestation may have different effects on the offspring depending on when the imposition occurs. The extent to which these effects may be amplified or ameliorated by the postnatal environment; i.e., the thrifty phenotype cannot be determined in the current study, because all sheep were treated similarly postnatally; but the potential is obvious (15). In addition, it is clear from studies in rats that the age-related decline in glucose tolerance in prenatally protein-restricted offspring is sex specific; i.e., it is delayed in female compared with male rats (16). When included as an independent covariate, the sex of the offspring did not influence any of the outcome measures in the current study; however, it is recognized that we do not have sufficient statistical power to draw firm conclusions regarding sex-specific effects on adult glucose metabolism after prenatal undernutrition in sheep. In addition, future studies should consider longitudinal assessments of glucose tolerance and insulin sensitivity of prenatally programmed male and female sheep to determine whether the observed decline with age (20) is exacerbated by a poor prenatal environment, specifically during functional development of fetal pancreas, i.e., mid-late gestation (17).

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