Impairment of insulin receptor signal transduction in placentas of intra-uterine growth-restricted newborns and its relationship with fetal growth

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Abstract

Objective: Intra-uterine growth restriction (IUGR) is related to a higher incidence of type 2 diabetes mellitus. We previously reported reduced adiponectin and increased interleukin 6 (IL6) concentrations in IUGR placentas, which are features of insulin resistance. We aimed to investigate placental insulin receptor (IR) function and activation in human placenta and subsequently the relationships of insulin signalling peptides with placental protein content in IL6, insulin, resistin and adiponectin, and with parameters of fetal growth.

Design and methods: Whole villous tissue was collected from 18 IUGR and 24 appropriate for gestational age (AGA) placentas of comparable gestational age. Insulin signalling peptides, suppressors of cytokine signalling-2 (SOCS2), insulin, adiponectin, resistin, and IL6 concentrations were determined by using western immunoblotting or specific research kits.

Results: The amount of total IR was similar in both groups but activated IR significantly higher in IUGR. Total IR substrate-1 (IRS1) was increased in IUGR, whereas total IRS2 and activated IRS1 were similar. AKT content was reduced and activated AKT was undetectable in IUGR placentas. c-Jun N-terminal kinase content was reduced in IUGR. Total and activated ERK1/2 was similar in IUGR and AGA groups, and total SOCS2 was increased in IUGR. IL6 lysate concentrations correlated with AKT content and activated IR. Correlations were found also with adiponectin and resistin. SOCS2 correlated negatively with all growth parameters at birth.

Conclusions: IR was more activated in placentas of IUGR compared with AGA; however, signal transduction downstream of the receptor was impaired. The increase in activated IR could be in favour of a compensatory mechanism to increase insulin sensitivity. Close relationships of insulin action in placenta with fetal growth were shown.

Introduction

The failure of a fetus to attain its expected fetal growth at any gestational age is defined as intra-uterine growth retardation (IUGR). Subjects born with IUGR have been reported to have an increased incidence of developing cardiovascular disease, hypertension and type 2 diabetes mellitus in adulthood compared with the general population (1). These features are shared with obese subjects, and derive primarily from increased insulin resistance (2), which is related by association with high serum concentrations of pro-inflammatory cytokines and low adiponectin concentrations (3, 4).

Adiponectin concentrations have been found reduced in placentas of women with gestational diabetes (5). Moreover, in gestational diabetes, the insulin receptor (IR) and some of the main peptides involved in insulin signal transmission were reported to be increased (6, 7).

Insulin resistance was reported recently in human pre-eclamptic placentas also, and was related to impaired insulin signalling (8). Interestingly, both gestational diabetes and pre-eclampsia are characterised by an increased inflammatory status (7, 9).

We recently reported significantly lower adiponectin and significantly higher interleukin 6 (IL6) concentrations in placental lysates of IUGR newborns (10).

Moreover, we showed that human IUGR subjects have increased insulin-like growth factor-binding protein-1 (IGFBP1) and IGFBP2 in the placenta compared with appropriate for gestational age (AGA) newborns. The IGFB2 peptide was also significantly increased in IUGR, and it could be speculated that IGFB2 could be increased to compensate for reduced insulin bioactivity (10, 11).

However, it is unknown at present in humans, in IUGR, whether the placenta shows features of insulin resistance. Insulin is a well-recognised in utero
growth factor besides a placental growth factor (12), and IL6 and other pro-inflammatory cytokines such as TNF-α are responsible for molecular mechanisms of insulin resistance (13, 14). The insulin signalling pathway is well known to control glucose uptake, synthesis, protein synthesis, and cell proliferation, differentiation and apoptosis.

In vitro studies have shown that IL6 induces decreased phosphorylation of IR substrate-1 (IRS1) and association of subunit p85 of the PI3-kinase with IRS1, in response to physiological concentrations of insulin. Furthermore, the insulin-dependent activation of AKT is markedly reduced with subsequent inhibition of glycogen synthesis (13), and chronic infusion of IL6 in murine liver induces insulin resistance through different mechanisms (15).

In vitro studies have shown that IL6 also stimulates ERK and c-Jun N-terminal kinase (JNK) pathways (16). ERK1/2 is part of the MAPK cascade, which is involved in controlling cell growth and differentiation. JNK1/2 MAPKs are also part of this signalling pathway. JNK1/2 activation appears to be involved in serine phosphorylation of IRS1 and subsequently insulin resistance on glucose uptake (17).

IL6 is also known to induce a family of negative regulators of cytokine signal transduction, the suppressors of cytokine signalling (SOCS) which have been shown to directly interact with the IR or IRS1 inhibiting insulin signal transduction (18, 19). SOCS2 is also an inhibitor of the GH receptor, and GH is also part of the interplay in the regulation of insulin sensitivity (20).

This study aimed to assess whether IR activation and signal transduction showed changes in the placenta of IUGR compared with AGA newborns of comparable gestational age, and to establish relationships of the insulin signalling peptides with placental protein content in IL6, insulin, resistin and adiponectin.

Relationships of single peptides with placental weight and parameters of fetal growth were also investigated.

Methods
Subjects

Eighteen IUGR and 24 AGA births of comparable gestational age (35.3 ± 0.5 vs 36.5 ± 0.4 weeks respectively NS) were followed during pregnancy. All pregnancies were dated correctly by ultrasound during the first trimester of gestation. AGA birth newborns were investigated. All pregnancies, both IUGR and controls, were delivered by elective caesarean section (CS).

The causes of preterm birth (< 38 weeks of gestation) in the AGA newborns were intra-hepatic colestasis in a woman who had undergone a previous CS, premature rupture of membranes with breech presentation and delivery within 4 h with no signs of infection. Elective CS in these subjects was performed because of refusal of vaginal delivery for psychological reasons, or because of a previous CS.

At birth, the following information was collected: age of the mother, weight at birth of both parents, body mass index (BMI) of the mother before pregnancy, previous gynaecological history, medical history during pregnancy, fetal biophysical data (exact duration of pregnancy and growth trend), clinical data at delivery (indication for CS, neonatal data as sex, weight, length, head circumference, APGAR score, acid–base equilibrium and perinatal data), weight and macroscopic aspect of the placenta. The main clinical features of both IUGR and AGA newborns are shown in Table 1.

| Table 1 Clinical data of newborns born from pregnancies complicated with IUGR and AGA at birth. Data are mean ± S.E.M. |
|-------------------------------------------------|-----------------|-----------------|
|                                   | IUGR            | AGA             |
| Males/females                      | 12/6            | 15/9            |
| Gestational age (years)            | 35.3 ± 0.5      | 36.5 ± 0.4      |
| Mothers’ chronological age (years) | 31.8 ± 1.3      | 33.2 ± 0.8      |
| Mothers’ BMI at the beginning of pregnancy (kg/m²) | 22.6 ± 0.9 | 22.4 ± 0.6 |
| Placental weight (kg)              | 440 ± 55        | 543 ± 41        |
| Birth weight (kg)                  | 1.4 ± 0.1*      | 2.85 ± 0.1      |
| Birth length (cm)                  | 39.7 ± 1.4*     | 49.1 ± 0.6      |
| Head circumference (cm)            | 29.4 ± 0.7*     | 34.3 ± 0.6      |

*P < 0.05 vs AGA.
and calcification, and decidua, marginal and chorion were avoided. The fragments were immediately rinsed in ice-cold sterile saline solution and placed in RNA Later (Calbiochem-Novabiochem Corporation, affiliate of Merck HgaA, Darmstadt, Germany), kept on ice and placed within 15 min at −20 °C and subsequently at −80 °C until tissue lysis.

**Placenta lysis, total protein determination and immunoprecipitation** The samples of placenta were sliced and homogenised in a lysis buffer at 0 °C (modified RIPA buffer: 50 mM Tris–HCl, 1% NP-40, 0.25% Na deoxycholate, 150 mM NaCl, 1 mM EDTA, 1 mM phenylmethylsulphonyl fluoride, 1 µg/ml each of aprotinin, leupeptin and pepstatin, 1 mM NaF). The total protein content was measured using the microassay Bio-Rad® protocol (Bio-Rad Lab), and the standard reference curve was obtained using bovine albumin. Immunoprecipitation was performed using a protein A/G-based method (Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA). Briefly, equal amounts of placenta lysates (1 mg total protein) were subjected to immunoprecipitation overnight at 4 °C using 1 µg specific antibodies, anti-IRS1 and anti-IRS2 (Upstate Biotechnology, Lake Placid, NY, USA). The resulting immunocomplexes were adsorbed to protein A/G Sepharose beads overnight at 4 °C as recommended by the manufacturer, and then eluted with electrophoresis sample buffer (Invitrogen).

**SDS-PAGE and western blotting** Equal amounts of immunoprecipitate or protein from the whole lysates were subjected to electrophoresis using precast gels according to molecular weight (NOVEL, Invitrogen). The resolved proteins were electrophotochemically transferred to nitrocellulose membranes (Hybond-C Extra, Amersham Biosciences Ltd) and hybridised with the following antibodies: anti-IRS1 (Upstate Biotechnology, Inc., Millipore Corporate, Billerica, MA, USA; dilution 0.5 µg/ml), anti-phospho IRS1 (Y896) (Upstate Biotechnology, Inc., Millipore Corporate; dilution 1:1000), anti-phospho IRS1 (S307) (Upstate Biotechnology, Inc., Millipore Corporate; dilution 1:1500), anti-IRS2 (Upstate Biotechnology, Inc., Millipore Corporate; dilution 1:1000), anti-IRS2 (Upstate Biotechnology, Inc., Millipore Corporate; dilution 1:200). The proteins were then analysed by ECL (Amersham Pharmacia Biotech) on autoradiographic film (Kodak BioMax). The differences between IUGR subjects and controls were analysed by scanning densitometry using the programme UN-SCAN-IT gel, version 5.1 (Silk Scientific Corporation, Orem, UT, USA). Band intensity was expressed in arbitrary units (optic densitometry units, ODU).

**Assays on placenta lysates** The total amount of IR (β subunit) was measured using a specific research ELISA kit (BioSource International, Camarillo, CA, USA). The intra-assay coefficient of variation (CV) was 5.4%, and the inter-assay was 8.2%. The sensitivity of the method was 0.5 ng/ml. Activation of the IR (pY1162/1163) was assessed using a specific ELISA kit by BioSource International. The intra-assay CV was 4.9%, and the inter-assay CV was 6.6%. The sensitivity of the method was 0.8 U/ml. AKT (total) was measured using a specific research ELISA kit (BioSource International). The intra-assay CV was 7.7%, and the inter-assay CV was 9.3%. The sensitivity of the method was <0.1 ng/ml. Activated AKT (pS473) was measured using a specific research ELISA kit (BioSource International). The intra-assay CV was 6.9%, and the inter-assay CV was 8.3%. The sensitivity of the method was 0.8 U/ml.

JNK (total) was measured using a specific research ELISA kit (BioSource International). The intra-assay CV was 8.8%, and the inter-assay CV was 9.2%. The sensitivity of the method was <0.15 ng/ml. Activated JNK (pTpY 183/185) was measured using a specific research ELISA kit (BioSource International). The intra-assay CV was 8.6%, and the inter-assay CV was 9.8%. The sensitivity of the method was <0.8 U/ml. IL6 was measured using an ultrasensitive ELISA method (QuantiKine HS, R&D Systems, Minneapolis, MN, USA) as previously described (10, 11). Insulin was assayed using a specific IRMA assay (DiaSorin S.p.A., Saluggia, Italy) as previously described (10). Resistin was assayed using ultrasensitive ELISA methods (QuantiKine HS, R&D Systems), and adiponectin was assayed using a specific ELISA research kit (QuantiKine HS, R&D Systems) as previously described (11).

Insulin, resistin, IL6 and adiponectin were assayed prior to this study (10).

All concentrations of placental lysates were normalised for milligrams of total protein. We previously showed, in addition, that total protein content was similar in IUGR and AGA placentas (11).

**Ethical approval**

Informed consent was obtained from the mothers. The study was approved by the ethics committee of the University of Parma Medical School.
Statistical analysis

The statistical analysis was performed using SPSS 17.1 for Windows (SPSS, Inc., Chicago, IL, USA) and Statistica 8.0 (StatSoft, Inc., Tulsa, OK, USA). The normal distribution of the data was determined using the Kolmogorov–Smirnov/Lilliefors test. These data were analysed using an unpaired t-test. IL6 and individual peptides downstream from the IR were not normally distributed and were analysed using the Wilcoxon rank sum test. The correlation analysis was performed using Pearson’s or Spearman’s coefficients, as appropriate, followed by Bonferroni’s correction. Only significant correlations are reported in the text.

Data are expressed as mean ± S.E.M. unless otherwise stated.

Results

Differences between placentas from IUGR and AGA newborns

Total and activated IR in lysates IR content was similar in IUGR and AGA placentas (4.00 ± 1.56 vs 6.72 ± 1.64 ng/mg respectively), whereas the amount of activated IR was greater in IUGR compared with AGA placentas (4.94 ± 1.84 vs 1.54 ± 0.33 U/mg respectively, P < 0.05; Fig. 1A and B); however, the activated/total IR ratio was higher but not significantly different in the IUGR (1.23 ± 1.4 vs 0.22 ± 0.05 U/ng in AGA).

Mediators of IR signal transduction IRS1 content was significantly increased in IUGR compared with AGA placentas (147.80 ± 13.43 vs 75.11 ± 13.57 ODU in AGA, P < 0.05; Fig. 2A), whereas the amount of IRS2 was similar in IUGR and AGA (68.19 ± 10.56 vs 75.11 ± 13.57 ODU respectively). Tyrosine phosphorylated IRS1 was increased but not significantly different in IUGR compared with AGA (223.62 ± 14.35 vs 175.99 ± 12.82 ODU respectively). The activated/total IRS1 ratio was similar in IUGR and AGA placentas (1.51 ± 0.17 vs 2.34 ± 0.47 ODU respectively).

Serine phosphorylated IRS1 was undetectable in all AGA, whereas the corresponding 170 kDa band was visible in some IUGR placental lysates (n = 5/18, data not shown).

Akt content, in IUGR, was significantly reduced (20.82 ± 3.96 vs 30.37 ± 2.40 pg/mg in AGA, P < 0.05; Fig. 2B), whereas activated Akt was substantially undetectable (measurable only in two samples/18) compared with AGA (2.73 ± 0.46 U/mg).

ERK1 and 2 content and activated ERK1 (94.51 ± 15.46 vs 93.59 ± 8.21 ODU in IUGR and AGA respectively) and 2 (122.66 ± 39.50 vs 105.12 ± 8.68 ODU in IUGR and AGA respectively) by tyrosine and threonine phosphorylation were not different in IUGR compared with AGA placentas.
The activated/total ERK1 ratio was similar in IUGR and AGA placentas (1.61 ± 0.10 vs 0.67 ± 0.07 ODU respectively; NS) as also the activated/total ERK2 ratio (1.08 ± 0.2 vs 1.25 ± 0.3 ODU respectively; NS).

JNK1/2 content was significantly lower in IUGR (0.30 ± 0.07 vs 0.49 ± 0.06 ng/mg, P < 0.05; Fig. 2C). Activated tyrosine phosphorylated JNK1/2 was lower but was not significantly different in IUGR compared with AGA placentas (0.02 ± 0.007 vs 0.72 ± 0.22 U/mg respectively, P < 0.05) with a significant decreased activated/total JNK ratio in IUGR (0.067 ± 0.004 vs 0.16 ± 0.03 U/ng in AGA, P < 0.05).

The 21 kDa band, corresponding to SOCS2, was significantly increased in IUGR compared with AGA placentas (75.06 ± 8.72 vs 49.69 ± 6.78 ODU respectively, Fig. 2D).

**Correlation analysis**

**Correlations with IL6 in placenta** The activated IR content was correlated with IL6 concentrations in lysates (r = 0.81; P = 0.001; Fig. 3).

AKT content was negatively correlated with IL6 (r = −0.35; P = 0.05).

**Correlations between peptides in placenta** Total and activated IR contents were correlated as expected (r = 0.45; P = 0.02). The IR content was correlated with insulin (r = 0.62; P = 0.03), resistin (r = 0.45; P = 0.018) and adiponectin concentrations in placental lysates (r = −0.38; P = 0.044). The activated IR content was correlated with IRS1 content (r = 0.73; P = 0.011), activated JNK (r = 0.38; P = 0.050) and resistin (r = 0.45; P = 0.018) in lysates.

**Figure 3** Normalised IL6 placental content was positively correlated with activated insulin receptor placental content (p-IR(pY1162/1163)). AGA newborns are represented as open circles, and IUGR newborns as open squares.

**Figure 4 SOCS2, expressed in optic densitometry arbitrary units (ODU), was negatively correlated with birth weight (A), birth length (B) and head circumference (C) of all newborns. AGA newborns are represented as open circles, and IUGR newborns as open squares.**

Activated IRS1 was correlated with IRS1 content (r = 0.64; P = 0.048), insulin (r = 0.61; P = 0.044) and adiponectin concentrations in lysates (r = −0.66; P = 0.005), and with activated JNK (r = 0.56; P = 0.018).

IRS2 content was correlated with adiponectin concentration (r = 0.54; P = 0.015).

JNK1/2 content was correlated with adiponectin concentration (r = 0.41; P = 0.024).

**Correlations of placental peptides with placental weight and growth parameters at birth** Placental weight was correlated with IRS2 (r = 0.37; P = 0.043) and AKT placental contents (r = 0.48; P = 0.003).

Birth weight was correlated with the amount of activated IR (r = −0.37; P = 0.042), JNK1/2 content (r = 0.39; P = 0.034) and SOCS2 (r = −0.50; P = 0.012; Fig. 4A).

Birth length was correlated with the activated IR content (r = −0.56; P = 0.004) and SOCS2 (r = −0.58; P = 0.008; Fig. 4B).

Head circumference was correlated with the activated IR content (r = −0.64; P = 0.005), SOCS2 (r = −0.61; P = 0.046; Fig. 4C) and the activated/total AKT ratio (r = 0.88; P = 0.008).

The correlations between the auxological parameters at birth and the above-described peptides, except SOCS2, are summarised in Table 2.

**Discussion**

This study explored for the first time IR signal transduction in placentas of IUGR compared with AGA pregnancies. The findings showed increased activated IR content in IUGR compared with AGA
placentas but reduced signal transduction with reduced AKT activation, possibly affecting mainly glucose and protein synthesis.

Similar ERK1/2 activity, but reduced JNK1/2 content and activation, was demonstrated suggesting that cell growth, proliferation and differentiation could also be impaired in IUGR placentas. Correlation analysis showed relationships between changes in IR signal transduction and IL6 concentrations in placental tissue. Finally, the amount of activated IR, AKT, JNK1/2 and SOCS2 showed clear relationships with fetal growth parameters at birth.

The human placenta is known to contain a large number of IRSs, and a metabolic action of these receptors via fetal control has been previously hypothesised (12). Gestational age was comparable in the two groups investigated to avoid changes in IR and insulin content, related with the duration of pregnancy (12).

The similar activated IRS1 content and the detectable serine phosphorylated IRS1 in IUGR is a first step to suggest that insulin signalling might be impaired just downstream of the activated IR. This process is a common finding in insulin-resistant states and in type 2 diabetes mellitus (23, 24). IRS1 protein is considered to be crucial for regulation of glucose uptake in insulin-sensitive tissues, and our findings in IUGR placentas could suggest reduced glucose uptake.

Total and activated AKT were significantly decreased in IUGR placentas, and data suggested a very low activated/total AKT ratio in IUGR compared with AGA placentas. This finding is in agreement with findings in trophoblast cell lines and in IUGR pregnancies due to pre-eclampsia, a condition, however, excluded from our studies (25). Moreover, these same authors showed, using AKT1 null mice, a central role for AKT in IUGR in association with endoplasmic reticulum stress (25). AKT is also part of the signalling pathway of the type 1 IGF receptor. At variance with our findings, AKT content was described to be reduced, whereas activated AKT unchanged in IUGR placentas (26). These slight differences could be owed to the smaller number of placentas these authors investigated, and to the fact that in that study, preterm IUGR pregnancies were compared with AGA placentas at term. The negative relationship between AKT and IL6 confirms in vivo, in humans, previous in vitro findings where insulin-dependent activation of AKT was markedly inhibited by IL6 treatment (13).

ERK1/2 content was similar in IUGR and AGA and failed to be activated in IUGR at variance with data showing ERK1/2 activation after IL6 administration to specific brain regions of rats (27), and studies in human umbilical vein endothelial cells (28). We also failed to identify any relationship with IL6 concentration in placenta, possibly suggesting tissue- and species-specific effects of IL6. We also speculate that the absent activation could be due to crosstalk among receptors, and the increased SOCS2 could account for down-regulation of signalling through the GH and IGF type 1 receptors.

We expected JNK to be activated and/or increased by the high IL6 concentrations in IUGR placentas, and to be possibly implicated in IRS1 serine phosphorylation, thus inhibiting insulin action (16, 17); however, JNK content and activated JNK were reduced in the IUGR placentas. This suggests an impairment of the MAPK pathway being significantly reduced in IUGR. The absence of a relationship with IL6 is at variance with previous in vitro data (16). The relationship we detected between JNK and adiponectin could suggest a link with insulin action or sensitivity in the placenta. These latter findings are altogether similar to those reported by Laviola et al., (26) although less marked.

The significant increase in SOCS2 in IUGR placentas could be consistent with both reduced insulin action and reduced GH action in IUGR placentas (29). Moreover, SOCS2 can bind the type 1 IGF1 receptor both in vivo and in vitro, suggesting that it can down-regulate the bioactivity of this receptor also (20, 30).

We did not find a direct relationship of SOCS2 with IL6 concentration in placental lysates, and must therefore hypothesise SOCS2 activation by a different mechanism; however, SOCS2 could certainly play an important role in both insulin action through a direct action on the IR or IRS1 (20, 29, 30), and indirectly by inhibiting the GH receptor (31). GH action is in fact recognised as an important factor regulating insulin sensitivity (32). Placental regulation is unknown and certainly, it is largely dependent on specific placental features (12).

SOCS2 knockout mice have gigantism (20). The increased concentrations in IUGR placentas and negative relationships with all growth parameters at birth are in line with previous experimental evidence, and demonstrate in humans the importance of this mechanism for fetal growth.

A role for insulin on fetal and placental growth is well established (10, 12). Insulin concentration in lysates was directly related with the total amount of IR and activated IRS1 but not with other post-receptor mediators suggesting interferences of other signalling peptides, possibly signalling through different receptors, modifying the final effect of insulin. Our findings are in agreement with those in placentas from pregnancies complicated with gestational diabetes (33).

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<th>Variables</th>
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<tr>
<td>Birth weight (kg)</td>
<td>IR content</td>
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<tr>
<td>Birth length (cm)</td>
<td>JNK1/2 content</td>
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<td>Head circumference (cm)</td>
<td>Activated IR content</td>
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Table 2: Correlation coefficients ($r$) and $P$ values of auxological parameters at birth with insulin signalling peptides, except SOCS2, of both IUGR and AGA newborns.
The relationship of resistin with total and activated IR content, and the correlations of adiponectin with IR content, activated IRS1, IRS2 and JNK suggest that both peptides play a role in placenta in regulating insulin action, and are in line with published reports (34–36). Findings are both in agreement with previous data (5), and at variance with recent findings in placenta of women with gestational diabetes (37).

Our data showed a direct relationship of placentatal weight with IRS2 and AKT confirming that insulin is also an important placentatal growth factor (12). The relationships we described between activated IR, JNK, SOCS2 and total/activated AKT ratio and growth parameters at birth confirm that insulin signalling through the IR, and in particular via AKT activation, was crucial for fetal growth during pregnancy.

In IUGR placentas, the overall data showed reduced insulin action where the increase in activated IR could be in favour of a compensatory mechanism to increase insulin sensitivity in a condition where insulin signal transduction is impaired downstream the receptor. Finally, these findings confirmed the importance of insulin signalling and SOCS2 for normal fetal growth. It remains to be elucidated how and for how long these placentatal changes influence post-natal modifications related with insulin sensitivity.

Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement

M E Street designed and led the study, analysed the data, drafted, developed and finalised the manuscript; I Viani, M A Ziveri and A Smerieri carried out the assays, discussed the findings, prepared the content, and helped to develop and edit the manuscript; C Volta and S Bernasconi discussed the findings, helped to develop and edit the manuscript.

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