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Maternal diet quality during pregnancy is associated with biomarkers of metabolic risk among male offspring

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Abstract

Aims/hypothesis—Limited data exist on the association between maternal diet quality during pregnancy and metabolic traits in offspring during early childhood, which is a sensitive period for risk of obesity-related disorders later in life. We aimed to examine the association of maternal diet quality, as indicated by the Healthy Eating Index-2010 (HEI), in pregnancy with offspring metabolic biomarkers and body composition at age 4–7 years.

Methods—We used data from 761 mother–offspring pairs from the Healthy Start study to examine sex-specific associations of HEI >57 vs ≤57 with offspring fasting glucose, leptin, cholesterol, HDL, LDL, percentage fat mass, BMI z score and log-transformed insulin, 1/insulin, HOMA-IR, adiponectin, triacylglycerols, triacylglycerols:HDL, fat mass, and sum of skinfolds. Multivariable linear regression models accounted for maternal race/ethnicity, age, education, smoking habits during pregnancy and physical activity, and child’s age.

Results—During pregnancy, mean (SD) HEI score was 55.0 (13.3), and 43.0% had an HEI score >57. Among boys, there was an inverse association of maternal HEI with offspring glucose, insulin, HOMA-IR and adiponectin. For instance, maternal HEI >57 was associated with lower fasting glucose (−0.11; 95% CI −0.20, −0.02 mmol/l), and lower concentrations of: insulin by 15.3% (95% CI −24.6, −5.0), HOMA-IR by 16.3% (95% CI −25.7, −5.6) and adiponectin by

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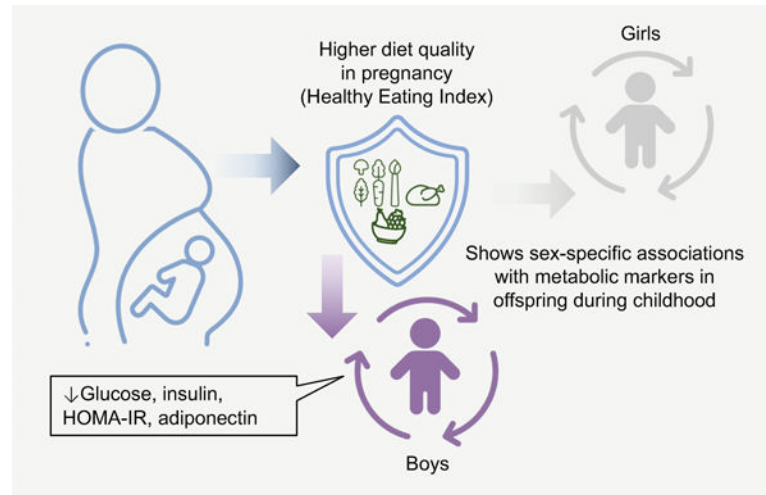
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9.3% (95% CI -16.1, -2.0). Among girls, there was an inverse association of maternal HEI with insulin and a positive association with LDL. However, following covariate adjustment, all estimates among girls were attenuated to the null.

Conclusions/interpretation—Greater compliance with the USA Dietary Guidelines via the HEI may improve the maternal–fetal milieu and decrease susceptibility for poor metabolic health among offspring, particularly boys. Future studies are warranted to confirm these associations and determine the underlying mechanisms.

Graphical Abstract



Keywords

Childhood glucose metabolism; Healthy Eating Index; Maternal dietary quality; Pregnancy

Introduction

Maternal nutritional status during pregnancy is a key contributor to the intrauterine environment and fetal development. Both historical and contemporary studies have exemplified the importance of macronutrient intake and balance, as well as intake of specific nutrients (e.g., folate, calcium, iron), for a range of offspring health outcomes, including obesity and related sequelae [1]. However, assessment of individual nutrients fails to capture the overall quality of diet and the interplay between nutrients and non-nutrient components in foods [2]. Of increasing interest are diet indices, such as the Healthy Eating Index-2010 (HEI), that measure dietary patterns marked by higher consumption of vegetables, fruit, fish and unsaturated fats, in conjunction with lower intakes of red and processed meat and saturated fats [3–5].

Few studies have assessed the HEI among pregnant women, and most have focused on associations with offspring body composition [6–13]. Biomarkers of metabolic risk, particularly alterations in insulin resistance and secretion, are also important to consider as differences in such biomarkers have been found in children and adults within normal ranges of weight [14, 15]. Sex differences in body composition, glucose homeostasis,

insulin signalling and lipid metabolism typically diverge during adolescence [16, 17], but have been detectable as early as at birth [18, 19], and may be attributable to differences in fetal response to the gestational environment [19–21]. Further, studies of murine and human placentas have shown clear sex-specific differences in gene expression with respect to maternal diet during pregnancy [22–24].

The current study assessed whether diet quality during pregnancy influences offspring metabolic health. We calculated the HEI [25], based on the 2010 USA Dietary Guidelines, and examined associations with biomarkers of glucose homeostasis, the adipoinular axis, lipid metabolism and measures of body composition among offspring in early childhood (4–7 years of age), a sensitive life stage for development of obesity and risk of chronic disease [26, 27]. Given documented sex differences in the outcomes of interest (body composition and metabolic biomarkers) in youth [17], and evidence that fetal response to the in utero environment differs for male vs female fetuses [24], we hypothesised that a higher-quality diet during pregnancy is associated with lower adiposity and biomarkers of metabolic risk, and that these associations differ by offspring sex.

Methods

Study participants and design

The Healthy Start study is a multi-ethnic pre-birth cohort of mother–offspring pairs who were enrolled at <24 weeks from prenatal clinics at the University of Colorado Hospital between 2009 and 2014. Women were excluded if they had prior diabetes, a history of prior premature birth with a gestational age (GA) <25 weeks or fetal death, asthma with active steroid management, serious psychiatric illness or multiple gestations. During pregnancy, women completed questionnaires on demographic characteristics, personal and family medical histories, and behaviours during pregnancy. Offspring returned for an in-person visit when they were between 4 and 7 years of age, during which body composition, anthropometry and a fasting blood sample were obtained. All women provided written informed consent and children 7 years of age or older provided assent. The study protocol and procedures were approved by the Colorado Multiple Institutional Review Board. The Healthy Start study is registered as an observational study at [ClinicalTrials.gov](https://clinicaltrials.gov/ct2/show/study/NCT02273297) (registration no. NCT02273297).

Mother–offspring pairs were eligible for the current analysis if women had at least one dietary recall during pregnancy ($N=1366$), and had an offspring who attended an in-person child visit at age 4–7 years ($N=907$). We excluded offspring without air displacement plethysmography or blood assay data ($N=134$), and women missing components necessary to calculate the HEI ($N=12$). The final sample included 761 mother–offspring pairs (Fig. 1). Participants who were included were similar to those excluded, except those included had a slightly older maternal age (~1 year) and a greater percentage reported a household income \$70,000 (36.7% vs 27.9%).

Maternal dietary intake and the HEI

Beginning in the first trimester (GA<13 weeks), maternal diet was assessed via the Automated Self-Administered 24 h Dietary Recall (ASA24) [28]. Details on the frequency of ASA24 completion by GA and offspring sex can be found in electronic supplementary material (ESM) Table 1. The mean number of ASA24s per woman was between three and five. A National Cancer Institute SAS macro (<https://epi.grants.cancer.gov/hei/developing.html>; accessed June 2021) was used to generate the HEI total scores from averaged recalls for each participant, which has been extensively reviewed [4]. Briefly, the model estimates the distribution of usual HEI scores based on a multivariate distribution (covariates: pre-pregnancy BMI, gravidity and smoking status) of usual intakes, which allows for the assessment of diet quality over time [4]. The HEI is a measure of diet quality used to assess compliance with USA Dietary Guidelines that are intended to be met over time and not necessarily every day [4, 29]. The HEI has 12 components assessed on a density basis, e.g. per 4184 kJ (1000 kcal) or as a percentage of kilojoules. Alcohol was not included in the HEI because all participants consumed under 13 g of alcohol per 4184 kilojoules for each recall, which is below the threshold for inclusion of calories from alcohol.

Offspring body composition and metabolic biomarkers

Fat mass (FM) and fat-free mass (FFM) were measured using whole body air displacement plethysmography (BodPod, Life Measurement, USA) with the Pediatric Option [30]. Offspring weight was measured using an electronic scale. Triceps, subscapular and mid-thigh skin-fold thicknesses were measured using Lange Skin-fold Calipers (Beta Technology, USA) to the nearest 1.0 mm by trained nurses. Body composition measurements for each participant were taken in triplicate and the mean of the two closest measures was used for analyses. Skinfolds were summed (sum of skinfolds) as a measure of subcutaneous adiposity. Age-specific BMI z scores were calculated according to the World Health Organization growth reference [31, 32].

Fasting triacylglycerols (TAGs), total cholesterol, HDL, LDL and glucose were measured using manufacturer-prepackaged enzymatic kits and the AU400e Chemistry Analyzer (Olympus America, USA). Insulin was measured using a radioimmune assay, and leptin and adiponectin were measured using a Multiplex assay kit, all by Millipore (USA). We calculated the ratio of TAGs:HDL as an indicator of an atherogenic lipid profile and a strong correlate of insulin resistance [33, 34], $1/(\text{fasting insulin})$ as a measure of insulin sensitivity [35] and an updated HOMA-IR [36]. The inter-assay CVs of these biomarkers were all <6.0%.

Covariate assessment

Maternal race/ethnicity, educational attainment, parity and smoking status during pregnancy were self-reported via questionnaire. Physical activity level was assessed twice during pregnancy with the Pregnancy Physical Activity Questionnaire as metabolic equivalent tasks (METs) in hours/week and averaged as a global measure of physical activity during pregnancy [37]. We calculated maternal pre-pregnancy BMI (kg/m^2) based on pre-pregnancy weight obtained from medical records among women who received primary care from

an affiliated University of Colorado doctor (89%) or from self-report (11%), and height measured at the first research visit. Early childhood dietary data were collected via two ASA24 dietary recalls (one weekend and one weekday), with parents as a proxy, and nutrient and caloric intakes were derived using the Nutrition Data System for Research software package. We derived the child HEI scores using the same procedure as for the mothers, except the 2015 USA Dietary Guidelines were used to align with the timing of data collection. Offspring physical activity was measured using wGT3X-BT ActiGraph accelerometers (Pensacola, FL, USA) [38] and categorised based on youth-specific cut-points [39]. Details on sample sizes for covariates are reported in ESM Tables 2, 3.

Statistical analysis

Prior to the main analysis, we examined bivariate associations of perinatal characteristics chosen based on prior knowledge with HEI. To create the most parsimonious model, we used bivariate analysis to select only those characteristics associated with HEI and offspring metabolic and body composition outcomes. To better understand differences in metabolic markers by offspring sex, we compared the mean differences in biomarkers and adiposity indicators in early childhood between boys and girls. The p values for an interaction by sex were not statistically significant for some of the outcomes (e.g., insulin, leptin, HDL) when tested in an unadjusted or covariate-adjusted model (Model 2, see below) that included HEI and sex as main and interaction terms. However, we observed substantially different results between the sexes, which, in conjunction with our a priori research interest, informed our decision to present all results stratified by sex.

For the primary analysis, we took the mean of food group servings across multiple recalls in pregnancy for each participant in light of prior evidence indicating that maternal diet quality is consistent across pregnancy [40, 41], and because the HEI includes food components that are consumed both episodically and regularly. In a previous study within this cohort, a score of >57, representing the upper two quintiles of the HEI distribution, was associated with neonatal adiposity [42]. Given the similarity of this threshold to HEI >58, which represented the upper two quintiles among participants in the present study, and to HEI 59, which corresponds with mean diet quality for Americans [43], we dichotomised the score as >57 vs 57 given its biological relevance among Healthy Start mother–child dyads and to facilitate comparability across studies in this cohort [42].

We used linear regression models to estimate the association of maternal HEI score >57 vs 57 with offspring glucose concentration, insulin, 1/insulin, HOMA-IR, adiponectin, leptin, cholesterol, TAGs, HDL, LDL, TAGs:HDL ratio, percentage fat mass (%FM), FM, BMI z score, and sum of skinfolds. Offspring insulin concentration, 1/insulin, HOMA-IR, adiponectin, TAGs and TAGs:HDL ratio, and measures of FM, and sum of skinfolds, were natural log transformed due to non-normal distributions. We present the results within the text as per cent change, calculated as: %change = $(\exp(\beta)-1) \times 100$.

Multivariable models were specified as follows. Model 1 adjusted for mean GA at ASA24 to account for differences in GA at dietary recalls. Model 2 included mean GA at ASA24, and confounders to the relationship between maternal diet and offspring health (maternal race/ethnicity, age, education, pre-pregnancy BMI and prenatal smoking habits, and child's age

at outcome assessment). Model 3: potential mediators such as maternal physical activity in pregnancy, gestational weight gain status and offspring birthweight. Model 4: all covariates in Model 2 plus child HEI score and physical activity levels, which may be related to maternal diet during pregnancy through backdoor paths. In our interpretation of results, we focus on estimates from Model 2, and assess for consistency in the direction, magnitude and precision of estimates across models to ensure robustness of associations. Sample sizes of covariates are reported in ESM Tables 2, 3.

To enhance interpretation of findings and to understand the interrelatedness of metabolic biomarkers and adiposity indicators in early childhood, we calculated Spearman correlations among markers of body measures and metabolic biomarkers in offspring.

Assessment of residuals from multivariable models indicated a normal distribution. Given the collinearity of outcomes in this study, testing for multiple comparisons was likely overly conservative [44]. However, to account for potential false findings, we tested mean differences in levels of biomarkers (log transformed as appropriate) by maternal HEI >57 vs ≤57 and adjusted p values using the bootstrap method, with 20,000 resamples performed with replacement within each sex [45, 46].

Sensitivity analyses

First, to check for differences in HEI during early (<27 gestational weeks) vs late pregnancy (≥27 gestational weeks), we compared mean scores among women who had data at both time-points ($N=424$) and calculated Pearson correlations. In this subsample of 220 boys and 204 girls, we assessed the association between HEI and offspring outcomes during early and late pregnancy. In addition, to allow for potential changes in diet quality across pregnancy, we assessed associations of having an HEI score >57 in both early and late pregnancy (vs not) with offspring outcomes. Second, although the HEI is based on ratios of nutrient intake per total energy, we adjusted models for total energy intake kilocalories to ensure our findings were not solely driven by energy intake. Third, to assess for a dose–response relationship between HEI and offspring metabolic and body composition outcomes we modelled HEI as a continuous variable, as well as in quintiles, and tested for a linear trend by fitting the median HEI score of each quintile as a continuous variable. Fourth, to ensure that our findings were independent of the diet change that often accompanies a diagnosis of gestational diabetes mellitus (GDM), we excluded these women ($N=17$ among boys, $N=18$ among girls).

Finally, although some studies suggest that HEI >80 represents a high-quality diet [47], there is no validated HEI threshold in pregnant women. Only a small proportion of women in our sample had an HEI >80 (2.10%), and therefore we were underpowered to perform stratified analyses using this cut-off.

Results

The mean (SD) GA in weeks at time of recall for women who completed only one dietary assessment was 19.6 (4.8) (data not shown). For women who completed more than one dietary recall ($N=636$), 97% completed ASA24s in the second (GA 13–26 weeks) and third

trimesters (GA 27 weeks to delivery). On average, women had an HEI score of 55.0 (SD 13.3) throughout pregnancy, and 43.0% had a score >57, a threshold associated with lower adiposity at birth in this cohort. On average, women with a score >57 consumed fewer carbohydrates, less total fat and slightly less protein compared with women with a score 57 (257.9 g [SD 51.1] vs 262 g [SD 56.1] of carbohydrates, 76.9 g [SD 13.8] vs 82.2 g [SD 15.57] of total fat and 82.7 g [SD 12.9] vs 81.8 g [SD 13.0] of protein).

Table 1 shows the sample sizes, mean age, distribution of child race/ethnicity, biomarkers and adiposity indicators between boys and girls at 4–7 years. The mean age at body composition measurement and blood draw was 4.8 (SD 0.69) in boys and 4.8 (SD 0.71) in girls. There were no racial/ethnic differences in the distribution of boys and girls. Boys had slightly higher levels of glucose mmol/l (4.62 [SD 0.37] vs 4.52 [SD 0.33]; p value <0.001) and as expected were larger overall. Among girls, we observed higher leptin and TAG concentrations and a slightly greater sum of skinfolds.

At delivery, the women were 28.3 (SD 6.1) years of age. A higher HEI score was associated with higher education, lower pre-pregnancy BMI, not smoking during pregnancy and lower physical activity (Table 2). Within each offspring sex there were similar patterns of associations when assessing maternal characteristics and HEI score during pregnancy. The HEI score was slightly higher among women of male vs female offspring (55.9 [SD 13.3] vs 54.1 [SD 13.3]; p=0.07), although there were no differences in the proportion of women with HEI>57 by sex (p=0.37). Associations of maternal characteristics with offspring outcomes generally tracked within each sex (ESM Tables 2, 3). For instance, offspring of Hispanic women and women with lower educational status or higher pre-pregnancy BMI had higher concentrations of glucose and HOMA-IR.

Among boys, there was an inverse association of maternal HEI with offspring glucose (−0.11; 95% CI −0.20, −0.02), insulin (%change −15.3; 95% CI −24.6, −5.0), HOMA-IR (%change −16.3; 95% CI −25.7, −5.6), adiponectin (%change −9.3; 95% CI −16.1, −2.0) and TAGs:HDL (%change −0.3; 95% CI −0.6, 0.01) in unadjusted models (Table 3); however, the upper CI for the estimate of TAGs:HDL crossed the null. These associations were materially unchanged after adjusting for confounders or potential mediators, though estimates for glucose became only marginally significant. Following further adjustment for the child's HEI score and physical activity levels, the magnitudes of associations for glucose, adiponectin and TAGs:HDL were slightly attenuated and no longer reached the threshold of statistical significance.

When modelling HEI as quintiles or as a continuous variable, we observed similar findings to HEI >57, with a significant linear trend indicating a dose–response relationship between higher-quality diet and lower glucose, insulin and adiponectin levels among boys (ESM Table 4). Associations of HEI as quintiles and continuous variables among girls are reported in ESM Table 5. After adjusting p values for multiple comparisons, we did not observe substantial differences in our findings for the insulin-based metrics (insulin, 1/insulin and HOMA-IR bootstrap p values <0.05) (ESM Fig. 1).

Among girls (Table 4), there was an inverse association of maternal HEI with offspring insulin (%change -12.0 ; 95% CI $-21.9, -0.9$) and a positive association with $1/\text{insulin}$ (%change -13.6 ; 95% CI $0.9, 28.0$) and LDL (5.45 ; 95% CI $0.90, 9.99$) in unadjusted models. All estimates were attenuated to the null after accounting for covariates (Table 3, Models 1–3). ESM Fig. 2 shows correlations among the biomarkers and adiposity indicators among offspring at 4–7 years. Insulin was positively correlated with %FM and negatively correlated with adiponectin within boys and girls. Among boys, adiponectin was negatively correlated with BMI z score, whereas among girls adiponectin was positively correlated with %FM and BMI z score.

Sensitivity analyses

The mean HEI score in early pregnancy (N=682) was 61.1 (10.1) and in late pregnancy (N=441) was 63.0 (10.3). Among women who had data at both time-points (N=424), the mean HEI score in early pregnancy was 62.3 (9.9) and in late pregnancy was 63.2 (10.3), with a global mean change of 0.03 (0.18) and a percentage change of 1.5%. HEI scores were positively correlated in early and late pregnancy (Pearson correlation: 0.46; $p < 0.001$). In analyses of early and late pregnancy HEI scores, each assessed separately as predictors of the offspring outcomes, we found consistent direction and magnitude of associations (albeit attenuated to the null due in part to the reduced sample size), including the inverse association between HEI score and biomarkers of glucose–insulin homeostasis among boys only (ESM Table 6). Additionally, findings were similar when we assessed associations of having consistently high HEI score (>57) across early and late pregnancy with offspring outcomes.

When controlling for total maternal kilocalories (tested in Model 2), we found no substantial change, such that HEI remained significantly associated with glucose ($p=0.02$), insulin ($p=0.02$), $1/\text{insulin}$ ($p=0.02$), HOMA-IR ($p=0.03$), adiponectin ($p=0.01$) and TAGs:HDL ratio ($p=0.03$) among boys and there were no significant associations among girls. Excluding women with GDM slightly attenuated the estimates and reduced statistical significance, with some evidence for a greater impact on insulin-based metrics. However, the overall trend in direction of association across all outcomes remained unchanged.

Discussion

Using longitudinal data collected in 761 mother–child pairs from gestation to early childhood, we examined associations of maternal diet quality during pregnancy, as measured by the HEI, with a panel of metabolic biomarkers and measures of body composition among offspring 4–7 years of age. Higher maternal diet quality during pregnancy was associated with a more favourable glucose–insulin homeostasis and lipid profile in male offspring, as indicated by lower concentrations of glucose, insulin and HOMA-IR, and the ratio of TAGs:HDL, even after accounting for potential confounders and mediators. Maternal HEI score was not associated with metabolic biomarkers or body composition among girls after accounting for maternal and perinatal characteristics.

In the present analysis, we found that already at 5 years of age, sex dimorphisms in markers of glucose–insulin homeostasis and body composition were detectable. Our most noteworthy

finding is the apparent protective effect of higher maternal diet quality on the glucose–insulin homeostasis and lipid profile of boys. Interestingly, higher maternal diet quality was associated with lower levels of adiponectin in boys. Among adults, lower adiponectin is often associated with worse metabolic outcomes such as type 2 diabetes [48]. Prior studies in neonates and children have found positive associations of adiponectin with weight, weight gain and sum of skinfolds in childhood [49–51]. However, in childhood and during periods of rapid development (e.g., 4–7 years of age), the role of adiponectin is less clear. Indeed, in the current study, we found that among boys adiponectin was inversely associated with BMI *z* score and among girls it was positively associated with BMI *z* score. Although the exact roles of adipokines in early childhood are unknown and warrant future study, adiponectin may play a role in weight gain during early childhood that differs by sex. Thus, our findings of higher diet quality in pregnancy and lower offspring adiponectin may represent a protective process by which a healthful diet in pregnancy reduces risk for later increased weight gain in boys.

To date, most studies that examined maternal diet quality via the HEI, or HEI adaptations, assessed associations with outcomes at birth [11, 12, 42, 52, 53]. In general, our findings in male offspring align with existing literature documenting inverse associations of maternal HEI score with neonatal FM [11, 42], and markers of insulin resistance in cord blood [52], though sex-specific estimates were not reported. For instance, in a study among 35 Spanish mother–offspring pairs, low maternal diet quality in the first trimester was associated with higher glucose, insulin and HOMA-IR in cord blood [12]. However, not all studies have found associations with offspring metabolic health, either at birth [53] or in early childhood [8]. In addition, interpretations across studies should be considered cautiously given differences in participant demographics and cultural practices [12, 52], timing of offspring outcome assessment (birth vs childhood) and use of birthweight as a marker of metabolic risk.

The lower atherosclerotic and glycaemic components of the HEI diet [54] may influence fetal programming of metabolism and glucose homeostasis [55, 56] and increase susceptibility to metabolic traits in childhood. In the current study, despite no difference in the diet quality of women who delivered male vs female offspring, maternal diet was associated with offspring metabolic traits in a sex-specific manner. Data from animal models indicate that males are more susceptible to overnutrition in the gestational period and early life, which may arise from the effects of hormones, epigenetics and differences in placental size, shape and efficiency [6, 21, 57, 58]. Further, studies of the epigenome and transcriptome of murine and human placentas have shown clear sex-specific differences in gene expression with respect to maternal diet during pregnancy [22–24]. The ability of the placenta to adapt to changes in the maternal–fetal milieu [59] may contribute to the sex-specificity of our findings. A prevailing hypothesis is that male fetuses respond to the maternal milieu with fewer functional changes in the placenta, resulting in persistent fetal growth and development, regardless of whether that environment is optimal or substandard. In contrast, female fetuses show greater placental adaptation to maternal exposures, resulting in greater placental gene and protein changes and increased survival rate when confronted with adverse events such as nutrient deprivation [59–61]. Thus, it may be that the placentas of female fetuses are less impacted by the maternal environment, whether positively or

negatively; and, while male fetuses demonstrate fewer adaptations, their increased sensitivity to maternal exposures results in greater receipt of benefits when, for instance, a healthy dietary pattern is consumed. However, this latter concept is speculative, and the underlying mechanisms are still under investigation.

Maternal HEI and offspring body composition

In the current study we did not find associations between maternal diet quality and offspring body composition. Prior studies have reported an inverse association of HEI with neonatal adiposity and birthweight [11, 12, 42, 62]. One study that followed offspring into adolescence and adulthood (ages 12–23 years) reported no association of HEI with risk of obesity during any time-period [63]. This could suggest that the effect of diet quality in pregnancy on offspring adiposity may diminish with increasing age. One potential explanation for our relatively null findings between maternal diet quality and early childhood adiposity is increased intra- and inter-individual variability in body composition during these transitional years. Continued follow-up will allow us to assess growth trajectories of participants as they age to shed light on this hypothesis.

Limitations of the data

We measured the HEI, which was designed to reflect the USA Dietary Guidelines, and thus our findings are likely more generalisable to women in the USA. The calculation of the HEI was based on data from dietary recalls collected over the course of pregnancy, which may suffer recall bias. However, this method of dietary assessment is considered valid to estimate relative dietary intake in large studies (and, therefore, inter-individual rank is preserved). On average, slightly more recalls were completed in the second and third trimesters (i.e., mean of 3.7 recalls during the first trimester, 3.8 recalls during the second trimester and 4.4 recalls during the third trimester) and thus our findings may reflect the effect of dietary quality in these time-periods. We provided a series of sequentially adjusted multivariable models that account for maternal and child confounders, mediators and lifestyle factors. Although these adjustments did attenuate some estimates, the association of higher maternal diet quality with markers of glucose–insulin homeostasis among boys remained significant.

Conclusions and future directions

In the current study, higher maternal diet quality during pregnancy was associated with a more favourable glucose–insulin homeostasis and lipid profile in male offspring. Future studies focused on identifying dietary intake thresholds among pregnant women should explore numerous cut-offs and consider whether thresholds are specific to the outcome of interest, i.e., offspring health vs maternal health. There are many mediating lifestyle factors that link maternal diet to offspring health which could be points of intervention, and which warrant further exploration in future analyses. The onset of childhood obesity and associated metabolic traits are occurring at increasingly early ages [64, 65], highlighting the gestational period as a critical window during which prevention efforts could have long-lasting impacts. The relevance of sex in susceptibility to and severity of disease is highly complex, and whether a particular sex has a more appropriate response to diet quality during gestation that persists into adulthood is likely dependent on myriad factors [58]. Given that pregnancy represents a window of opportunity for change that may result in

sustained healthy behaviours for both mother and child, increased emphasis on adherence to dietary patterns during pregnancy that align with the HEI and USA Dietary Guidelines may improve the maternal–fetal milieu.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Data availability

Data described in the manuscript, code book and analytic code will be made available upon request pending reasonable application and approval.

Abbreviations

ASA24	Automated Self-Administered 24-h Dietary Recall
FM	Fat mass
%FM	Percentage fat mass
GA	Gestational age
GDM	Gestational diabetes mellitus
HEI	Healthy Eating Index-2010
TAG	Triacylglycerol

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Research in context

What is already known about this subject?

- Maternal diet quality during pregnancy is important for fetal development
- Recent data indicate there are sex differences in the sensitivity of the fetus to the intrauterine environment

What is the key question?

- Is maternal diet quality in pregnancy, as indicated by the Healthy Eating Index-2010, associated with offspring metabolic biomarkers and body composition at age 4–7 years, and are the associations sex-dimorphic?

What are the new findings?

- Higher maternal Healthy Eating Index score was associated with lower fasting glucose, insulin and adiponectin among male offspring at age 4–7 years

How might this impact on clinical practice in the foreseeable future?

- Greater compliance with the dietary guidelines for all pregnant women may improve the maternal-fetal milieu and decrease susceptibility for poor metabolic health among offspring, particularly male offspring

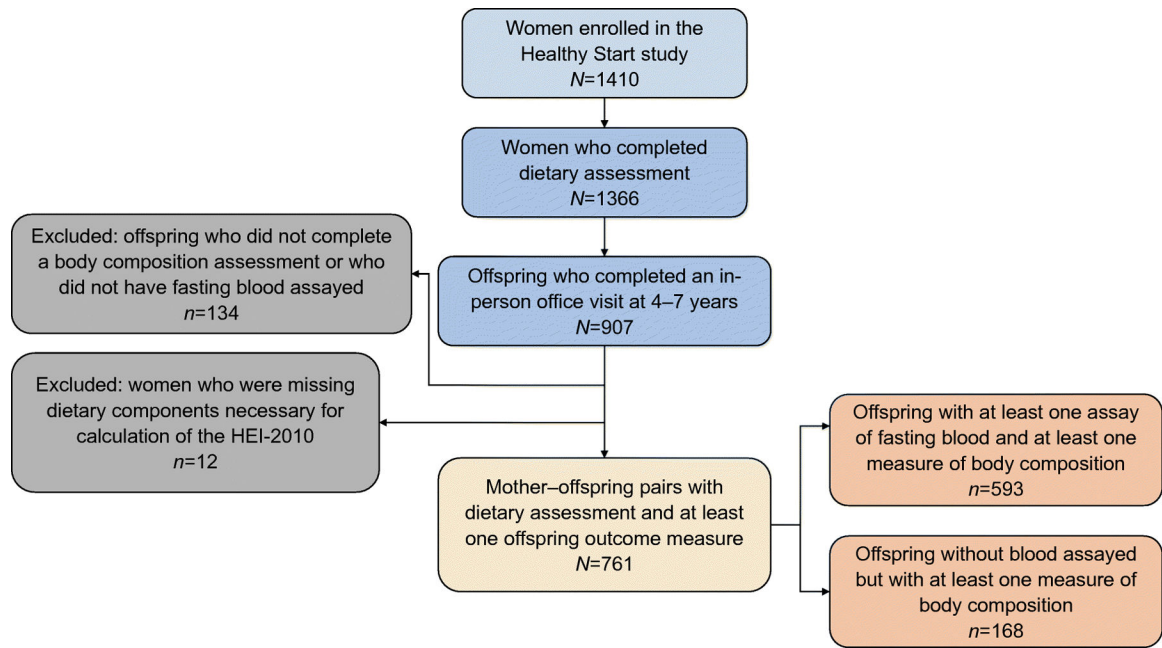


Fig. 1.
Participant flow chart

Table 1

Child characteristics, metabolic biomarkers and body composition by sex

Characteristic	Boys		Girls		<i>p</i> ^a
	<i>n</i>	Mean ± SD	<i>n</i>	Mean ± SD	
Age ^b	396	4.8 ± 0.69	365	4.8 ± 0.71	0.67
Race/ethnicity (<i>N</i> , %) ^c					0.63
Non-Hispanic white	201	50.8	179	49.0	
Non-Hispanic black	43	10.9	51	14.0	
Hispanic	101	25.5	89	24.4	
Non-Hispanic other ^d	51	12.9	46	12.6	
Metabolic biomarkers					
Glucose, mmol/l	272	4.62 ± 0.37	231	4.52 ± 0.33	<0.001
Insulin, pmol/l ^e	284	3.63 ± 0.49	251	3.67 ± 0.47	0.26
1/insulin, pmol/l ^e	284	-3.63 ± 0.49	251	-3.68 ± 0.47	0.26
HOMA-IR, % ^e	263	-0.35 ± 0.49	215	-0.32 ± 0.46	0.47
Adiponectin, µg/ml ^e	168	2.33 ± 0.26	119	2.36 ± 0.26	0.32
Leptin, µg/l	167	3.83 ± 1.00	196	6.75 ± 1.91	<0.001
Cholesterol, mmol/l	286	36.8 ± 5.62	243	37.2 ± 5.64	0.38
TAGs, mmol/l ^e	285	0.043 ± 0.004	244	0.444 ± 0.003	0.01
HDL, mmol/l	219	12.83 ± 1.36	181	12.71 ± 1.51	0.40
LDL, mmol/l	270	21.27 ± 5.23	227	21.99 ± 4.50	0.10
TAGs:HDL ratio ^e	218	0.0034 ± 0.0005	181	0.0036 ± 0.0005	0.02
Body composition					
%FM	345	19.91 ± 6.68	335	19.72 ± 6.89	0.70
FM, kg ^e	345	1.23 ± 0.44	335	1.15 ± 0.52	0.04
BMI <i>z</i> score	389	0.22 ± 1.11	358	0.05 ± 0.90	0.02
Sum of skinfolds, mm ^e	390	3.39 ± 0.28	360	3.50 ± 0.26	<0.001

^a *p* value based on generalised linear models^b Age at blood draw and assessment of body composition^c *p* value based on χ^2 ^d Other includes Asian, American Indian/Alaska natives, Hawaiian/Pacific Islanders^e Estimates represent geometric means; *p* value based on log-transformed outcome

Table 2
Differences in maternal HEI total score during pregnancy by maternal characteristics and offspring sex

Characteristic	All			Male			Female		
	Total n ^a	Mean ± SD	p ^a	n	Mean ± SD	p ^a	n	Mean ± SD	p ^a
HEI total score	761	55.0 ± 13.3		396	55.9 ± 13.3		365	54.1 ± 13.3	
Score < 57	434	45.7 ± 7.9	–	219	46.2 ± 7.8	–	215	45.1 ± 8.1	–
Score > 57	327	67.5 ± 7.5		177	67.9 ± 7.8		150	67.1 ± 7.1	
Age at index pregnancy (years)			<0.001			<0.001			<0.001
16–24	222	48.8 ± 12.5		102	47.2 ± 12.3		120	50.1 ± 12.5	
25–29	192	53.3 ± 12.8		96	56.1 ± 12.3		96	50.5 ± 12.7	
30–34	224	59.7 ± 12.3		132	59.8 ± 12.3		92	59.7 ± 12.4	
35	123	60.6 ± 11.9		66	61.2 ± 11.7		57	59.9 ± 12.2	
Race/ethnicity			<0.001			<0.001			0.01
Non-Hispanic white	419	58.3 ± 13.3		218	59.5 ± 13.0		201	56.9 ± 13.5	
Non-Hispanic black	115	46.6 ± 10.3		57	47.5 ± 10.4		58	45.7 ± 10.3	
Hispanic	186	53.8 ± 12.5		99	53.6 ± 13.1		87	54.0 ± 11.8	
Non-Hispanic other ^b	41	51.5 ± 12.2		22	51.6 ± 10.3		19	51.3 ± 14.3	
Education			<0.001			<0.001			<0.001
High school or less	223	49.3 ± 12.1		100	49.7 ± 12.5		123	48.9 ± 11.8	
Some college/associate degree	182	51.8 ± 12.4		95	51.9 ± 12.2		87	51.6 ± 12.8	
College graduate	166	59.1 ± 12.3		90	59.2 ± 11.7		76	59.0 ± 13.1	
Graduate degree	190	61.4 ± 12.4		111	62.2 ± 12.6		79	60.3 ± 12.2	
Pre-pregnancy BMI (kg/m ²) ^c			0.05			0.09			0.21
Underweight (<19.0)	19	58.6 ± 14.5		9	56.8 ± 16.6		10	60.3 ± 13.1	
Normal (19.0–24.9)	377	55.8 ± 13.7		188	57.2 ± 13.9		189	54.3 ± 13.4	
Overweight (25.0–29.9)	206	54.3 ± 12.7		113	54.4 ± 12.6		93	54.1 ± 12.9	
Obese (≥ 30.0)	159	53.8 ± 12.8		86	54.7 ± 12.3		73	52.8 ± 13.4	
Gestational weight gain ^d			0.70			0.42			0.16
Inadequate	129	54.9 ± 12.6		63	54.0 ± 11.3		66	55.8 ± 13.7	
Adequate	95	58.1 ± 15.0		50	59.7 ± 15.8		45	56.3 ± 14.0	

Characteristic	All			Male			Female		
	Total	n ^a	Mean ± SD	n	Mean ± SD	p ^a	n	Mean ± SD	p ^a
Excessive	475		55.1 ± 13.2	247	56.5 ± 13.1	0.58	228	53.6 ± 13.1	0.90
GDM									
No	677		54.8 ± 13.3	352	55.6 ± 13.4	0.50	325	54.0 ± 13.2	
Yes	34		56.1 ± 12.7	17	57.8 ± 10.8		17	54.4 ± 14.4	
Smoking during pregnancy						<0.001			<0.001
No	713		55.9 ± 13.0	376	56.5 ± 13.1		337	55.2 ± 12.9	
Yes	48		42.5 ± 11.3	20	43.8 ± 11.5		28	41.5 ± 11.2	
Physical activity status (moderate-to-vigorous)						<0.001			0.09
<150 min/week	300		57.2 ± 12.8	158	58.7 ± 13.1	0.01	142	55.6 ± 12.3	
150 min/week	461		53.6 ± 13.5	238	54.0 ± 13.2		223	53.2 ± 13.8	

^aTests for significant differences between mean maternal HEI score and maternal characteristics in the entire sample and stratified by offspring sex were based on generalised linear models; *p* value represents Type III main effects for categorical variables. For ordinal variables (age, education level, pre-pregnancy weight status, gestational weight gain status and birthweight categories) the *p* value represents a test for linear trend; dash (–) indicates no test for statistical difference

^bOther includes Asian, American Indian/Alaska natives, Hawaiian/Pacific Islanders

^cDivision of Nutrition, Physical Activity, and Obesity, National Center for Chronic Disease Prevention and Health Promotion

^dInstitute of Medicine 2009 guidelines

Table 3

Association of maternal HEI score >57 vs ≤57 during pregnancy with childhood biomarkers of glucose homeostasis, the adipoinular axis, lipids and body composition among boys (*n*=396)

HEI score >57 vs ≤57 Outcome	MODEL 1 (GA at diet recall)			MODEL 2 (confounders)			MODEL 3 (mediators)			MODEL 4 (concurrent child lifestyle)		
	<i>n</i>	β (CI)	<i>p</i>	<i>n</i>	β (CI)	<i>p</i>	<i>n</i>	β (CI)	<i>p</i>	<i>n</i>	β (CI)	<i>p</i>
Glucose, mmol/l	272	-0.11 (-0.20, -0.02)	0.01	272	-0.10 (-0.19, -0.00)	0.04	223	-0.08 (-0.18, 0.02)	0.11	199	-0.09 (-0.20, 0.01)	0.09
Insulin, pmol/l ^a	284	-0.17 (-0.28, -0.05)	<0.01	284	-0.13 (-0.25, -0.01)	0.04	232	-0.15 (-0.29, -0.02)	0.03	206	-0.15 (-0.29, -0.01)	0.03
1/insulin, pmol/l ^a	284	0.17 (0.05, 0.28)	<0.01	284	0.13 (0.01, 0.25)	0.04	232	0.15 (0.02, 0.29)	0.03	206	0.15 (0.01, 0.29)	0.03
HOMA-IR, % ^a	263	-0.18 (-0.30, -0.06)	<0.01	263	-0.13 (-0.26, -0.00)	0.046	216	-0.15 (-0.29, -0.01)	0.04	192	-0.15 (-0.29, -0.01)	0.03
Adiponectin, µg/ml ^a	168	-0.10 (-0.18, -0.02)	0.01	168	-0.11 (-0.19, -0.03)	0.01	137	-0.12 (-0.21, -0.02)	0.02	122	-0.08 (-0.17, 0.02)	0.13
Leptin, µg/l	167	0.22 (-0.09, 0.53)	0.17	167	0.34 (0.00, 0.68)	0.047	139	0.47 (0.11, 0.83)	0.01	121	0.39 (0.02, 0.76)	0.04
Cholesterol, mmol/l	286	0.17 (-4.91, 5.26)	0.95	286	-0.74 (-6.33, 4.86)	0.80	234	-0.64 (-6.58, 5.30)	0.83	211	-0.20 (-6.65, 6.25)	0.95
TAGs, mmol/l ^a	285	-0.02 (-0.10, 0.06)	0.55	285	-0.05 (-0.13, 0.04)	0.28	233	-0.07 (-0.16, 0.03)	0.15	211	-0.02 (-0.11, 0.08)	0.74
HDL, mmol/l	219	0.99 (-0.41, 2.39)	0.16	219	1.12 (-0.39, 2.64)	0.15	180	1.29 (-0.23, 2.81)	0.10	162	1.10 (-0.68, 2.88)	0.23
LDL, mmol/l	270	-0.89 (-5.78, 3.99)	0.72	270	-1.14 (-6.48, 4.20)	0.68	221	-0.75 (-6.38, 4.87)	0.79	199	1.04 (-5.04, 7.11)	0.74
TAGs:HDL ratio ^a	218	0.00 (-0.01, 0.00)	0.06	218	0.00 (-0.01, 0.00)	0.02	179	0.00 (-0.01, 0.00)	0.01	162	0.00 (-0.01, 0.00)	0.17
%FM	345	0.96 (-0.46, 2.38)	0.18	345	0.83 (-0.62, 2.29)	0.26	282	0.45 (-1.10, 2.00)	0.57	244	0.91 (-0.80, 2.61)	0.30
FM, kg ^a	345	0.05 (-0.05, 0.14)	0.31	345	0.04 (-0.06, 0.14)	0.43	282	0.01 (-0.09, 0.11)	0.84	244	0.02 (-0.09, 0.13)	0.71
BMI z score	389	-0.18 (-0.40, 0.04)	0.11	389	-0.04 (-0.26, 0.18)	0.73	317	-0.14 (-0.37, 0.08)	0.22	274	-0.14 (-0.38, 0.10)	0.26
Sum of skinfolds, mm ^a	390	-0.03 (-0.09, 0.03)	0.27	390	-0.03 (-0.09, 0.03)	0.30	318	-0.04 (-0.10, 0.03)	0.27	275	-0.05 (-0.12, 0.02)	0.16

Model 1 adjusts for mean GA at diet recall throughout pregnancy

Model 2 (confounders): GA at diet recall; maternal race/ethnicity, age, education, pre-pregnancy BMI and prenatal smoking habits; and child's age

Model 3 (mediators): Model 2 + maternal physical activity during pregnancy and gestational weight gain status, and offspring birthweight

Model 4 (concurrent child lifestyle): Model 2 covariates + child's HEI score and physical activity levels

^aEstimates represent geometric means; *p* value derived from models where outcomes were log transformed due to non-normal distributions

Table 4

Association of maternal HEI score >57 vs ≤57 during pregnancy and childhood biomarkers of glucose homeostasis, the adipoinular axis, lipids and body composition among girls ($n=365$)

HEI score >57 vs ≤57 Outcome	MODEL 1 (GA at diet recall)			MODEL 2 (confounders)			MODEL 3 (mediators)			MODEL 4 (concurrent child lifestyle)		
	n	β (CI)	p	n	β (CI)	p	n	β (CI)	p	n	β (CI)	p
Glucose, mmol/l	231	-0.01 (-0.10, 0.07)	0.75	231	0.00 (-0.08, 0.09)	0.93	196	0.02 (-0.07, 0.11)	0.65	163	0.01 (-0.10, 0.12)	0.81
Insulin, pmol/l ^a	251	-0.13 (-0.25, -0.01)	0.03	251	-0.08 (-0.20, 0.04)	0.20	210	-0.03 (-0.16, 0.10)	0.60	172	-0.10 (-0.25, 0.06)	0.21
1/insulin, pmol/l ^a	251	0.13 (0.01, 0.25)	0.03	251	0.08 (-0.04, 0.20)	0.20	210	0.03 (-0.10, 0.16)	0.60	172	0.10 (-0.06, 0.25)	0.21
HOMA-IR, % ^a	215	-0.13 (-0.25, 0.00)	0.05	215	-0.08 (-0.21, 0.05)	0.22	182	-0.05 (-0.19, 0.09)	0.50	148	-0.07 (-0.24, 0.09)	0.39
Adiponectin, µg/ml ^a	119	0.03 (-0.07, 0.12)	0.60	119	0.00 (-0.10, 0.10)	0.99	94	-0.02 (-0.14, 0.11)	0.79	80	-0.05 (-0.15, 0.06)	0.37
Leptin, µg/l	196	0.01 (-0.54, 0.57)	0.96	196	0.06 (-0.52, 0.64)	0.84	159	0.19 (-0.47, 0.85)	0.57	130	0.03 (-0.66, 0.72)	0.93
Cholesterol, mmol/l	243	4.80 (-0.72, 10.32)	0.09	243	3.29 (-2.50, 9.07)	0.27	204	1.91 (-4.32, 8.13)	0.55	172	4.38 (-2.60, 11.36)	0.22
TAGs, mmol/l ^a	244	0.00 (-0.08, 0.08)	0.94	244	-0.02 (-0.10, 0.06)	0.64	206	-0.01 (-0.10, 0.08)	0.81	174	-0.03 (-0.13, 0.08)	0.62
HDL, mmol/l	181	0.05 (-1.70, 1.79)	0.96	181	0.39 (-1.42, 2.20)	0.67	154	-0.08 (-2.05, 1.89)	0.94	120	-0.56 (-2.84, 1.72)	0.63
LDL, mmol/l	227	5.45 (0.90, 9.99)	0.02	227	3.59 (-1.12, 8.31)	0.14	191	4.01 (-1.08, 9.11)	0.12	160	3.77 (-2.33, 9.88)	0.23
TAGs:HDL ratio ^a	181	0.00 (0.00, 0.00)	0.80	181	0.00 (-0.01, 0.00)	0.43	154	0.00 (0.00, 0.00)	0.74	120	0.00 (-0.01, 0.00)	0.92
%FM	335	-1.38 (-2.87, 0.10)	0.07	335	-1.13 (-2.55, 0.29)	0.12	277	-1.09 (-2.63, 0.45)	0.17	229	-1.10 (-2.85, 0.65)	0.22
FM, kg ^a	335	-0.06 (-0.17, 0.05)	0.29	335	-0.05 (-0.17, 0.06)	0.34	277	-0.03 (-0.14, 0.09)	0.66	229	-0.05 (-0.19, 0.10)	0.52
BMI z score	358	0.00 (-0.19, 0.19)	0.98	358	0.02 (-0.17, 0.21)	0.82	299	0.00 (-0.20, 0.20)	0.99	242	0.09 (-0.15, 0.33)	0.46
Sum of skinfolds, mm ^a	360	-0.01 (-0.06, 0.05)	0.84	360	0.00 (-0.06, 0.05)	0.89	299	0.01 (-0.05, 0.07)	0.71	245	0.01 (-0.06, 0.08)	0.73

Model 1 adjusts for mean GA at diet recall throughout pregnancy

Model 2 (confounders): GA at diet recall; maternal race/ethnicity, age, education, pre-pregnancy BMI and prenatal smoking habits; and child's age

Model 3 (mediators): Model 2 + maternal physical activity during pregnancy and gestational weight gain status, and offspring birthweight

Model 4 (concurrent child lifestyle): Model 2 covariates + child's HEI score and physical activity levels

^aEstimates represent geometric means; *p* value derived from models where outcomes were log transformed due to non-normal distributions