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Gestational Glucose Tolerance and Maternal Metabolic Profile at 3 Years Postpartum

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Abstract

OBJECTIVE—To estimate the independent effect of gestational impaired glucose tolerance, defined as a single abnormal oral glucose tolerance test (OGTT) value, on metabolic dysfunction at 3 years postpartum.

METHODS—We used multiple linear regression to measure associations between glucose testing during pregnancy and metabolic markers at 3 years postpartum in Project Viva, a prospective cohort study of maternal and infant health. We compared metabolic measures at 3 years postpartum among four groups: normal glucose challenge test (less than 140 mg/dL, n=461); abnormal glucose challenge test but normal glucose tolerance test (GTT) (n=39); impaired glucose tolerance (IGT) (a single abnormal GTT value, n=21); and gestational diabetes mellitus (GDM) (n=16).

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RESULTS—Adjusting for age, race, parity, parental history of diabetes, and maternal BMI at 3 years postpartum, we found women with GDM had lower adiponectin (11.2 ng/mL vs. 20.7 ng/mL) and higher homeostatic model assessment – insulin resistance (3.1 vs. 1.3) and waist circumference (91.3 cm vs. 86.2 cm) compared with women with IGT or normal glucose tolerance. Women in both the IGT and GDM groups had lower high-density lipoprotein (GDM: 44.7 mg/dL; IGT: 45.4/dL vs normal glucose tolerance 55.8 mg/dL) and higher triglycerides (GDM: 136.1 mg/dL; IGT: 140.1 mg/dL, vs. normal glucose tolerance: 78.3), compared with women in the normal glucose tolerance group. We found the highest values for Hemoglobin A1c (GDM: 5.1%; IGT 5.3%, normal glucose tolerance 5.1%) and high-sensitivity c reactive protein (GDM 1.4 mg/dL IGT: 2.2 mg/dL; NGT 1.0 mg/dL) among women with IGT.

CONCLUSION—GDM and IGT during pregnancy are associated with persistent metabolic dysfunction at 3 years postpartum, independent of other clinical risk factors.

Introduction

Metabolic dysfunction causes substantial morbidity and mortality among women. Among women, pregnancy complications predict metabolic disease risk. Women with gestational diabetes (GDM) have an 17 to 63% risk of developing type 2 diabetes within 5 to 16 years of the index pregnancy(1), and recent studies have linked a history of gestational diabetes with cardiovascular risk(2, 3).

The diagnosis of GDM presumes a threshold value above which women are at increased risk of pregnancy complications; however, recent work shows that adverse pregnancy outcomes rise continuously with increasing fasting glucose values(4). Maternal metabolic risk may similarly rise with glucose values. Indeed, recent studies have linked gestational impaired glucose tolerance with subsequent diabetes and cardiovascular disease risk(3, 5–9). However, it is not known whether impaired glucose tolerance during pregnancy predicts metabolic dysfunction independent of clinical risk factors such as body mass index and family history.

The aim of our study was to determine whether a history of GDM or gestational impaired glucose tolerance (IGT) is predictive of maternal metabolic dysfunction, independent of recognized clinical risk factors. We hypothesized that we would find a monotonic relationship between degree of gestational glucose tolerance and metabolic dysfunction at 3 years postpartum. To test this hypothesis, we compared measures of metabolic dysfunction at 3 years postpartum among women with normal glucose tolerance (NGT), abnormal glucose loading test results but normal GTT values, gestational impaired glucose tolerance (IGT), or GDM, in Project Viva, a prospective cohort study of maternal and infant health.

Materials and Methods

We performed an unplanned secondary analysis of participants in Project Viva, a longitudinal cohort study of maternal and child health(10). Women were recruited for Project Viva at their first prenatal visit at one of eight obstetrical offices of a multispecialty group practice in Eastern Massachusetts from 1999 to 2002. To be eligible for the study, potential participants were required to be fluent in English, <22 weeks gestation at study entry, and have a singleton pregnancy All participants provided written informed consent, and the Institutional Review Board of Harvard Pilgrim Health Care approved all procedures.

Assessment of gestational glucose tolerance

Obstetrical care providers assessed gestational glucose tolerance among women in our cohort according to the following guidelines: At 26 to 28 weeks' gestation, all mothers

underwent a non-fasting 50g oral glucose challenge test (GCT). Women with a result of 140 mg/dL or higher underwent a 100g oral glucose tolerance test (OGTT), administered the morning after an overnight fast. Normal results were defined by Carpenter-Coustan criteria: fasting <95 mg/dL, 1 hour <180 mg/dL, 2 hour < 155 mg/dL, 3 hour < 140 mg/dL. Gestational glucose tolerance was categorized as normal (GLT < 140 mg/dL), Abnormal GLT, normal GTT (GLT \geq 140 mg/dL, GTT with no abnormal results), gestational impaired glucose tolerance (IGT) (GLT \geq 140 mg/dL, GTT with only 1 abnormal result) (8, 11), or gestational diabetes (GLT \geq 140 and GTT with 2 or more abnormal results).

Assessment of metabolic parameters at 3 years postpartum

Women returned at 3 years postpartum for a physical examination that included anthropometric measurements and a blood sample. Methodology for anthropometric measures has been previously described elsewhere(12). We tested all blood samples (N=537) for HbA1c, sex hormone binding globulin (SHBG), C-reactive protein (CRP), and the adipokines leptin(13) and adiponectin(14). We identified as fasting those participants who did not eat or drink anything other than water for 8 hours before blood samples were obtained (N=166). We tested fasting blood samples for insulin, glucose, total cholesterol, LDL cholesterol, HDL cholesterol, triglycerides, IL 6, ghrelin(15) and PYY(16). When we compared women who provided fasting samples with those who provided non-fasting samples, we found no differences in age, race, parity, family history of DM, gestational glucose tolerance, or in any outcome variables measured in both fasting and non-fasting participants. Laboratory methods for assessment of metabolic markers in this study have been previously described (12, 17).

Study covariates

Participants reported sociodemographic variables including parity, race/ethnicity, and personal history of type 1 or type 2 diabetes at the initial study visit during prenatal care. They reported parental history of type 2 diabetes at three-year follow-up visit. Women missing data on study covariates were excluded.

Data Analysis

We used analysis of variance and chi square tests to measure bivariate associations between sociodemographic characteristics and gestational glucose tolerance. We used multiple linear regression to model the relation between gestational glucose tolerance category and metabolic markers at three-years. Because BMI may have a non-linear association with metabolic markers, we used linear, quadratic and 3-knot cubic spline models(18) to adjust for maternal BMI 3 years postpartum, retaining the more complex model if the log likelihood ratio test p value was < 0.05. Because inclusion of quadratic and 3-knot quadratic spline terms did not improve model fit, we modeled BMI as a linear variable. We further adjusted for maternal age, race, parity, and parental history of type 2 diabetes to ascertain the predictive role of gestational glucose tolerance independent of clinical risk factors for metabolic disease. Adjustment for breastfeeding duration and weight change from pre-pregnancy to the three-year visit did not materially change our results, and they were therefore excluded from our model.

The Shapiro-Wilk test and visual inspection of regression residuals suggested that normality should not be assumed in several cases. Log transformation of HOMA-IR, insulin, sex hormone binding globulin, triglycerides, C-reactive protein and IL6 improved normality of regression residuals. We present p values for the partial F test to assess the joint null hypothesis of equality across all of the glucose tolerance categories(19).

To facilitate interpretation of both the magnitude and clinical significance of differences among gestational glucose tolerance groups, we present results as predicted means and 95% confidence intervals for the mean. We present both unadjusted mean values and adjusted predicted mean values for participants of average postpartum BMI (26.2) who were white, age 35 to <40, had two children, and had no parental history of type 2 diabetes.

Data analyses were performed using SAS 9.2. Two-tailed P values < 0.05 were considered statistically significant.

Results

Of 5055 women screened for Project Viva, 4208 were eligible, and 2670 enrolled (Figure 1). Among the 2128 Project Viva participants who gave birth, 1579 met criteria for a three-year follow-up examination with their children by virtue of completing a pregnancy dietary questionnaire and consenting for child follow-up. Of these women, 611 met criteria for the current analysis because they attended the three-year visit, had not delivered another child since the birth of the index child three-years previously, denied a diagnosis of type 1 or type 2 diabetes early in the index pregnancy and provided a blood sample. We excluded women with missing data for three-year lab results (n=25), breastfeeding duration at 1 year (n=8), BMI at 3 years postpartum (n=14), or gestational glucose tolerance (n=27), leaving 537 women for analysis. At this three-year visit, 166 women provided a fasting blood sample.

Glucose challenge test results were normal for 85.9% (N=461, 95%CI 82.9–88.8%) of women in our cohort. Among the 76 women with a GCT \geq 140, 39 had all normal values on the 100g GTT, 21 had one abnormal value (IGT), and 16 met the diagnostic criteria for GDM. Maternal age, parity, body mass index at 3 years postpartum, and race were similar among the glucose tolerance groups. Women with a parental history of type 2 diabetes were more likely to have had GDM (Table 1).

Most women in our cohort gained weight from prior to the index pregnancy to the three-year visit: the median weight gain was 2.2 kg (Interquartile range -0.4 to 5.3 kg), and 26.5% (95%CI 22.8–30.4%) of women had gained 5 kg or more during this interval. Most participants had normal glucose tolerance at 3 years postpartum by ADA criteria(20). Among 535 women for whom HbA1C was obtained, 14 were at increased risk of diabetes (A1C 5.7–6.4%) and one, with an A1C of 6.58%, met criteria for ADA diabetes (A1C \geq 6.5%). In addition, one participant self-reported a diagnosis of type 2 diabetes. Among 164 women for whom fasting glucose was obtained, 2 had impaired fasting glucose (FPG 100–125 mg/dl) and none met criteria for diabetes (FPG > 126 mg/dl).

As we hypothesized, women with both IGT and GDM had more adverse metabolic profiles than women with normal GCT results or with abnormal GCT but normal values on the GTT. These patterns were similar in unadjusted models and in models adjusted for BMI at 3 years postpartum, parity, age, self-reported race and parental history of diabetes (Tables 2, 3 and 4). Women with GDM had lower adiponectin and higher HOMA-IR and waist circumference compared with women with IGT or normal glucose tolerance (Figure 2a, HOMA-IR GDM: 2.7 vs. NGT:1.3; adiponectin GDM 13.1 ng/mL vs. NGT 21.2 ng/mL; waist circumference GDM 91.3 cm vs. NGT 86.2 cm; partial F test p < 0.05 for all models). Women in both the IGT and GDM groups had lower HDL and higher triglycerides, compared with women in the NGT group (Figure 2b: HDL GDM: 44.7 mg/dL; IGT: 45.4 mg/dL vs NGT 55.8 mg/dL, partial F test p =0.07; triglycerides GDM: 136.1 mg/dL; IGT: 140.1 mg/dL, vs. NGT: 78.3 mg/dL; partial F test p <0.01). We had hypothesized that we would find a monotonic association between metabolic dysfunction and degree of glucose intolerance; however, we found the highest values for Hemoglobin A1c and CRP among

women with IGT (Figure 2c: Hemoglobin A1c GDM: 5.1%; IGT 5.3%, NGT 5.1%, partial F test $p<0.01$; hsCRP GDM 1.4 mg/dL IGT: 2.2 mg/dL; normal 1.0 mg/dL, partial F test $p<0.01$). We found no pattern of association between gestational glucose tolerance category and sex hormone binding globulin, total cholesterol, fasting IL6, leptin, ghrelin or PYY (Tables 2 and 3).

Discussion

In this community-based prospective cohort study, we found that both gestational diabetes and gestational impaired glucose tolerance were associated with an adverse metabolic profile at 3 years postpartum, independent of body mass index and parental history of diabetes.

Strengths of our study include its prospective assessment of gestational glucose tolerance and standardized assessment of three-year outcomes. Nevertheless, our results must be interpreted within the context of the study design. Our population was healthy, resulting in low rates of gestational diabetes and impaired glucose tolerance. Among the 91 women ages 30–39 for whom we had data on waist circumference, blood pressure, serum lipids and glucose, only 5 (5.5%, 95% CI 1.8–12.4%) met criteria for the metabolic syndrome, compared with 15% of women in this age range in the general US population(21). In addition, the number of participants with fasting blood samples limited power to detect subtle differences among glucose tolerance groups, and we were not able to define metabolic syndrome in the full cohort. Further studies in larger populations will be needed to validate our findings. Nevertheless, our study size is comparable to several other studies that have assessed metabolic markers among postpartum women with a history of GDM(2, 22). We did not measure post-glucose load insulin or glucose in our population, and therefore we were unable to compare indices of glycemia, insulin sensitivity and beta-cell function. Nevertheless, our study included postpartum measures of adiponectin, which is highly correlated with beta cell dysfunction during pregnancy(23) and with 2-h post OGTT in the postpartum period(24).

Our results confirm and extend earlier work linking gestational glucose tolerance with an adverse maternal metabolic profile in later life. Several authors have reported an increased risk of impaired glucose tolerance and type 2 diabetes among women with abnormal glucose screening results in pregnancy in the setting of both normal OGTT(7) and one abnormal GTT result (5, 6, 9, 25). Moreover, both IGT and GDM have been associated with the metabolic syndrome at 3 months postpartum(8). Other authors have reported associations between GDM and markers of metabolic dysfunction after pregnancy. At a mean of 2 years postpartum, Costacou et al reported adverse associations between history of GDM (N=22) and waist circumference, hemoglobin A1c, and HOMA-IR, compared with women without a history of pregnancy complications (N=29)(22). Heitritter et al similarly compared women with a GDM history (N=23) with normal controls (N=23) at a mean of 4 years postpartum. Women in the GDM group had higher diastolic blood pressure, mean arterial pressure, heart rate, fasting glucose, HOMA, triglycerides, CRP, IL-6, and PAI-1 and lower adiponectin than women in the control group.

No studies to our knowledge have measured associations between IGT and LDL, inflammatory markers or adipokines, or with other metabolic markers beyond 3 months postpartum. We found that women with impaired glucose tolerance during pregnancy had elevations of triglycerides, hemoglobin A1c and CRP, as well as lower HDL, after adjustment for current body mass index and parental history of diabetes. Women with a history of GDM had triglyceride and HDL levels that were similar to those with IGT, but they had higher HOMA-IR and waist circumference, as well as lower adiponectin levels.

These adverse profiles of intermediate markers among women with pregnancy dysglycemia imply increased risk for cardiovascular disease, which is consistent with findings in a recent population-based cohort study (3). In that study, compared with women who did not undergo glucose tolerance testing during pregnancy and therefore were presumed to have had normal glucose screening test results, women with both IGT and GDM were more likely to experience cardiovascular events (IGT OR 1.19, 95% CI 1.02–1.39; GDM OR 1.66, 95% CI 1.30–2.13).

Compared with women with normal glucose testing during pregnancy, we found that women with a history of gestational glucose intolerance had unfavorable markers of glucose and lipid homeostasis and inflammation. These findings persisted with adjustment for current body mass index, suggesting that normal or overweight women with a history of IGT may be at risk for metabolic dysfunction at 3 years postpartum. These women may therefore benefit from dietary changes, physical activity, and/or screening for metabolic syndrome. Current guidelines recommend screening women with a history of GDM for type 2 diabetes (26, 27).

In conclusion, in a prospective study of maternal and infant health, we found that maternal gestational glucose intolerance and gestational diabetes were both associated with adverse metabolic profile at 3 years postpartum, independent of other clinical risk factors.

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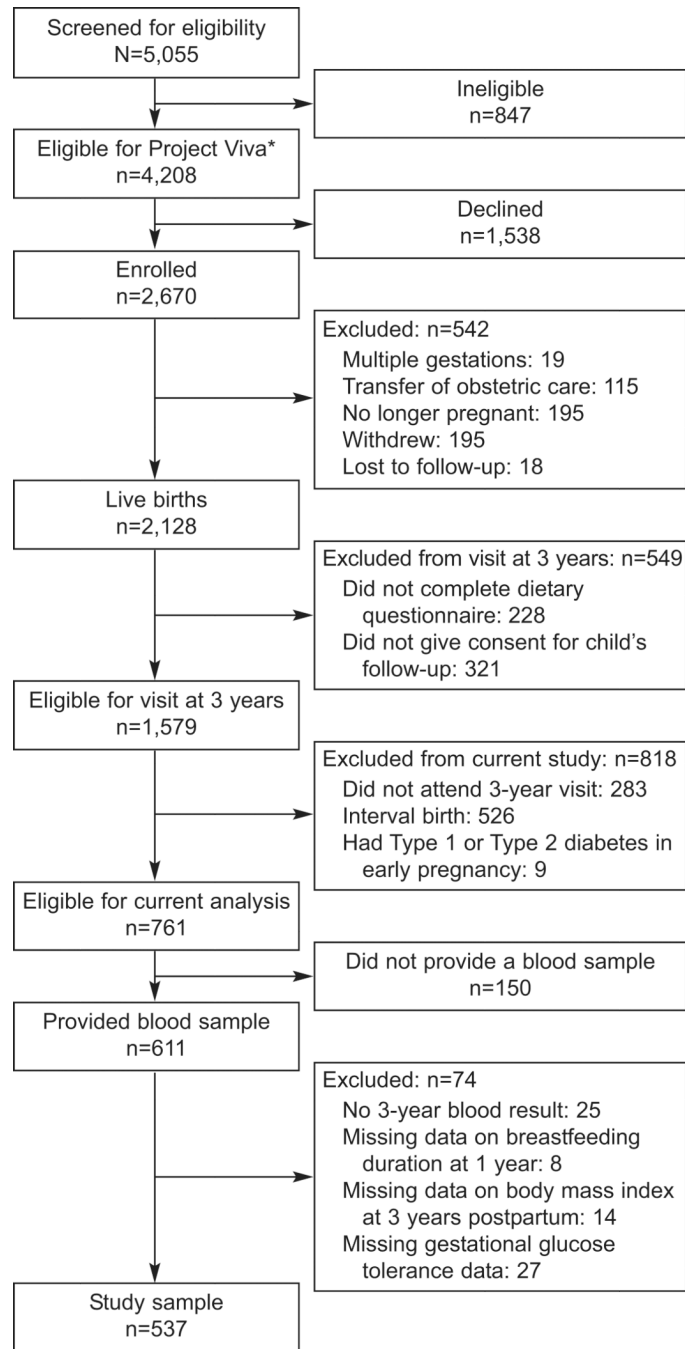


Figure 1.

Flow of patients through the current study, which is a secondary analysis of Project Viva, a longitudinal study of maternal and child health. *Eligibility requirements included singleton gestation, ability to answer questions in English, plans to remain in the area through delivery, and gestational age less than 22 weeks at initial prenatal clinical appointment.

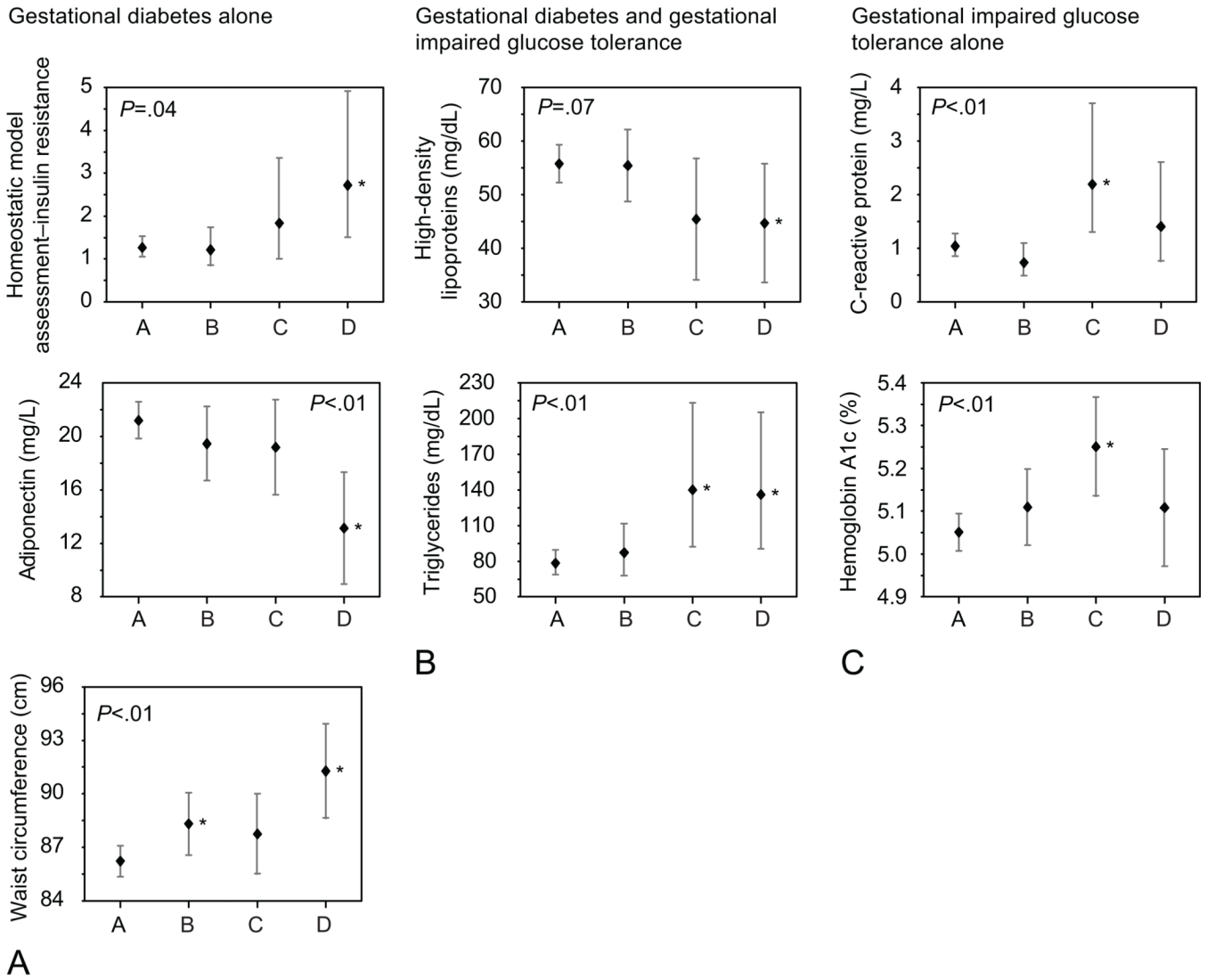


Figure 2. Mean predicted values for a white participant between 35 and less than 40 years of age with a body mass index (BMI) of 26.2 (the study population mean), who has two children and no parental history of diabetes. *P*-values on graph are for partial F test. The figure shows adverse metabolic markers that differ with gestational diabetes mellitus (GDM) alone (column A), both gestational impaired glucose tolerance and GDM (column B), or gestational impaired glucose tolerance alone (column C). In each graph's x-axis, A, B, C, and D represent the following groups: Group A, 50-g screen less than 140 mg/dL; B, 50-g screen 140 mg/dL or more with normal oral glucose tolerance test (OGTT); C, 50-g 140 mg/dL or more with one abnormal OGTT value; D, gestational diabetes mellitus (GDM: two or more abnormal OGTT values). Predicted mean, 95% confidence limit of the mean, adjusted for BMI, age, race, parity, and parental history of diabetes mellitus. **P*<.05 compared with normal glucose tolerance group.

Table 1

Baseline characteristics of study population, by pregnancy glucose tolerance category. Data from 578 participants in Project Viva who presented for follow-up at three years postpartum without an intervening birth.

	Normal		Abnormal GCT, normal GTT		Impaired glucose tolerance during pregnancy		Gestational diabetes		Missing gestational glucose category ²		p ¹
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	
N	461		39		21		16		28		
BMI at 3 year pp visit (kg/m ²)	25	(5)	26	(5)	25	(4)	27	(6)	27	(6)	0.15
Age at 3 year pp visit	37	(5)	39	(5)	38	(4)	38	(5)	38	(6)	0.51
	N	%	N	%	N	%	N	%	N	%	
Race											0.45
Asian	18	(82)	0	(0)	1	(5)	2	(9)	1	(5)	
Black	68	(83)	5	(6)	4	(5)	3	(4)	2	(2)	
Hispanic	29	(74)	5	(13)	2	(5)	0	(0)	3	(8)	
Other	22	(85)	0	(0)	0	(0)	1	(4)	3	(12)	
White	333	(81)	32	(8)	15	(4)	10	(2)	19	(5)	
Parity											0.78
1	131	(79)	11	(7)	7	(4)	7	(4)	10	(6)	
2	227	(83)	17	(6)	10	(4)	6	(2)	12	(4)	
3+	112	(80)	14	(10)	5	(4)	3	(2)	6	(4)	
Parental History DM											0.02
Yes	65	(74)	6	(7)	4	(5)	7	(8)	6	(7)	
No	405	(83)	36	(7)	18	(4)	9	(2)	22	(4)	
Missing BMI at 3 years ²											0.12
Yes	9	(60)	3	(20)	1	(7)	0	(0)	2	(13)	
No	461	(82)	39	(7)	21	(4)	16	(3)	26	(5)	

¹ ANOVA p value for continuous variables, chi square p value for categorical variables.

² Participants missing pregnancy glucose tolerance (N=27) or BMI at 3 years (N=14) were excluded from the analysis.

Markers of glucose and lipid homeostasis at three-years postpartum, by gestational glucose tolerance category. Predicted values¹ from multiple linear regression models. Data from 537 participants in Project Viva who presented for follow-up at three years postpartum without an intervening birth.

Table 2

	N	Normal		Abnormal GCT, normal GTT		Gestational impaired glucose tolerance		Gestational diabetes		P ³
		Mean (95%CI)		Mean (95%CI)		Mean (95%CI)		Mean (95%CI)		
Glucose metabolism										
Hemoglobin A1c%, N		460		38		20		15		
Unadjusted	533	5.10 (5.07–5.12)		5.18 (5.09–5.26)		5.32 ⁴ (5.20–5.44)		5.24 (5.10–5.37)		<.001
MV-Adjusted	533	5.05 (5.01–5.09)		5.11 (5.02–5.20)		5.25 ⁴ (5.14–5.37)		5.11 (4.97–5.24)		<.01
HOMA IR ² , N		142		12		4		5		
Unadjusted	163	1.3 (1.2–1.5)		1.3 (0.9–1.8)		1.6 (0.9–3.1)		3.1 ⁴ (1.7–5.5)		0.04
MV-Adjusted	163	1.3 (1.1–1.5)		1.2 (0.8–1.7)		1.8 (1.0–3.4)		2.7 ⁴ (1.5–4.9)		0.04
Insulin, u/mL ²										
Unadjusted	163	7.5 (6.8–8.4)		6.9 (4.8–9.9)		8.7 (4.7–16.2)		15.3 ⁴ (8.8–26.7)		0.09
MV-Adjusted	163	6.8 (5.8–8.1)		6.3 (4.5–8.7)		8.8 (5.1–15.3)		12.1 ⁴ (7.1–20.8)		0.12
Glucose, mg/dL										
Unadjusted	163	72.6 (70.3–74.8)		76.1 (68.3–83.8)		77.5 (64.1–90.9)		82.8 (70.8–94.8)		0.3
MV-Adjusted	163	76.5 (72.3–80.6)		79.0 (71.0–87.0)		82.9 (69.4–96.4)		89.0 ⁴ (75.8–102.3)		0.19
Sex hormone binding globulin ² , N		459		39		20		16		
Unadjusted	534	48.7		44.9		45.1		50.8		0.86

	N	Normal Mean (95%CI)	Abnormal GCT, normal GTT		Gestational impaired glucose tolerance		Gestational diabetes Mean (95%CI)	p ³
			Mean (95%CI)	Mean (95%CI)	Mean (95%CI)	Mean (95%CI)		
MV-Adjusted	534	50.0 (44.8–55.8)	48.8 (36.1–55.7)	50.3 (33.4–61.1)	59.2 (42.3–82.9)	0.77		
Lipid metabolism								
Total cholesterol, mg/dL, N	142	142	12	4	5			
Unadjusted	163	176.8 (171.9–181.6)	185.1 (168.3–201.8)	195.5 (166.5–224.5)	164.0 (138.1–189.9)	0.34		
MV-Adjusted	163	175.8 (166.6–185.0)	180.5 (162.9–198.0)	194.7 (165.1–224.4)	165.9 (136.8–194.9)	0.48		
LDL cholesterol, mg/dL								
Unadjusted	163	106.2 (101.7–110.7)	113.4 (98.0–128.8)	126.0 (99.3–152.6)	90.9 (67.1–114.8)	0.22		
MV-Adjusted	163	102.7 (94.4–111.0)	106.4 (90.6–122.3)	126.1 (99.3–152.8)	92.8 (66.6–119.0)	0.26		
HDL cholesterol, mg/dL								
Unadjusted	163	54.5 (52.5–56.4)	54.0 (47.2–60.7)	47.3 (35.5–59.0)	41.8 ⁴ (31.3–52.3)	0.09		
MV-Adjusted	163	55.8 (52.2–59.3)	55.4 (48.7–62.2)	45.4 (34.1–56.7)	44.7 ⁴ (33.6–55.8)	0.07		
Triglycerides, mg/dL ²								
Unadjusted	163	73.6 (68.6–78.9)	85.1 (67.0–108.1)	128.8 ⁴ (85.1–195.0)	133.3 ⁴ (92.0–193.2)	0.001		
MV-Adjusted	163	78.3 (68.7–89.2)	87.1 (68.0–111.7)	140.1 ⁴ (92.1–213.0)	136.1 ⁴ (90.3–205.2)	<.01		

¹Data presented are mean predicted values. MV-adjusted results are predicted means for a white participant age 35–<40 with a BMI of 26.2, the study population mean, who has two children and no parental history of diabetes.

²Results are geometric means.

³ Partial F test p among categories

⁴ Effect estimate $p < .05$ vs. NGT group

Table 3

Inflammatory markers and adipokines at three-years postpartum, by gestational glucose tolerance category. Predicted values¹ from multiple linear regression models. Data from 537 participants in Project Viva who presented for follow-up at three years postpartum without an intervening birth.

	N	Normal		Abnormal GCT, normal GTT		Gestational impaired glucose tolerance		Gestational diabetes		P ³
		Mean (95%CI)		Mean (95%CI)		Mean (95%CI)		Mean (95%CI)		
Inflammatory markers										
hsCRP, mg/dL ²										
Unadjusted	536	0.8 (0.7-0.9)	460	0.6 (0.4-0.9)	39	21	16	1.2 (0.6-2.4)	<.01	
MV-Adjusted	536	1.0 (0.8-1.3)	1.0	0.7 (0.5-1.1)	0.7	2.2 ⁴	1.4 (0.8-2.6)	<.01		
IL6, pg/mL ² , N										
Unadjusted	163	1.1 (0.9-1.2)	142	1.6 (0.9-2.6)	12	4	5 (0.7-3.3)	0.44		
MV-Adjusted	163	0.9 (0.7-1.1)	0.9	1.2 (0.7-2.0)	1.2	1.1	1.3 (0.6-3.1)	0.38		
Adipokines										
Leptin ng/mL, N										
Unadjusted	531	8.6 (8.0-9.1)	456	10.0 (8.1-11.9)	39	20	16 (6.5-12.4)	0.25		
MV-Adjusted	531	8.3 (7.7-8.9)	8.3	8.6 (7.4-9.9)	8.6	9.4	8.1 (6.2-9.9)	0.56		
Ghrelin pg/mL, N										
Unadjusted	165	769.1 (717.8-820.4)	144	699.8 (514.2-885.4)	11	5	459.5 ⁴ (184.2-734.8)	0.16		
MV-Adjusted	165	819.8 (739.8-899.7)	819.8	762.4 (603.0-921.8)	762.4	663.8	562.5 ⁴ (306.6-818.5)	0.11		
Adiponectin ng/mL, N										
		457	39	21	16					

	N	Normal		Abnormal GCT, normal GTT		Gestational impaired glucose tolerance		Gestational diabetes		p ³
		Mean (95%CI)	Normal	Mean (95%CI)	Abnormal GCT, normal GTT	Mean (95%CI)	Gestational impaired glucose tolerance	Mean (95%CI)	Gestational diabetes	
Unadjusted	533	20.7 (19.9–21.5)	19.1 (16.4–21.7)	18.0 (14.4–21.7)	11.2 ⁴ (7.1–15.4)	<.001				
MV-Adjusted	533	21.2 (19.8–22.6)	19.4 (16.7–22.2)	19.2 (15.6–22.7)	13.1 ⁴ (9.0–17.3)	<.001				
PYY ² pg/mL, N		143	12	5	5					
Unadjusted	165	58.5 (55.4–61.7)	55.2 (45.8–66.5)	51.8 (38.8–69.1)	71.3 (53.5–95.1)	0.42				
MV-Adjusted	165	59.9 (54.3–66.1)	55.8 (46.1–67.5)	52.0 (39.1–69.2)	71.8 (52.5–98.4)	0.38				

¹Data presented are mean predicted values. MV-adjusted results are predicted means for a white participant age 35–40 with a BMI of 26.2, the study population mean, who has two children and no parental history of diabetes.

²Results are geometric means.

³Partial F test p among categories

⁴Effect estimate p<.05 vs. NGT group

Anthropometry at three-years postpartum, by gestational glucose tolerance category. Predicted values¹ from multiple linear regression models. Data from 537 participants in Project Viva who presented for follow-up at three years postpartum without an intervening birth.

Table 4

	N	Normal		Abnormal GCT, normal GTT		Gestational impaired glucose tolerance		Gestational diabetes		P ³
		Mean (95%CI)		Mean (95%CI)		Mean (95%CI)		Mean (95%CI)		
Anthropometry										
3-year postpartum weight retention, kg, N		460		39		21		16		
Unadjusted	536	2.3 (1.7–3.0)		3.1 (0.9–5.2)		4.9 (1.9–7.8)		-0.1 (-3.5–3.3)		0.16
MV-Adjusted	536	1.6 (0.5–2.7)		1.9 (-0.3–4.1)		3.5 (0.7–6.4)		-1.6 (-4.9–1.8)		0.12
1-year postpartum weight retention, kg, N		399		32		17		16		
Unadjusted	464	0.9 (0.3–1.4)		0.5 (-1.4–2.4)		2.4 (-0.2–5.0)		-0.2 (-2.9–2.5)		0.56
MV-Adjusted	464	-0.2 (-1.2–0.8)		-0.5 (-2.5–1.5)		1.0 (-1.7–3.6)		-1.3 (-4.2–1.5)		0.65
Subscapular:Triceps skin fold ratio, N		460		39		21		14		
Unadjusted	534	0.8 (0.7–0.8)		0.8 (0.7–0.9)		0.8 (0.7–0.9)		0.9 ⁴ (0.8–1.0)		0.08
MV-Adjusted	534	0.7 (0.7–0.8)		0.8 (0.7–0.9)		0.8 (0.7–0.9)		0.9 ⁴ (0.8–1.0)		0.07
Waist circumference, cm, N		461		39		21		16		
Unadjusted	537	86.0 (84.9–87.1)		90.6 ⁴ (86.8–94.5)		89.6 (84.3–94.9)		93.3 ⁴ (87.3–99.4)		0.01
MV-Adjusted	537	86.2 (85.4–87.1)		88.3 ⁴ (86.6–90.1)		87.8 (85.5–90.0)		91.3 ⁴ (88.6–93.9)		<.001

¹Data presented are mean predicted values. MV-adjusted results are predicted means for a white participant age 35–40 with a BMI of 26.2, the study population mean, who has two children and no parental history of diabetes.

²Results are geometric means.

³ Partial F test p among categories

⁴ Effect estimate $p < .05$ vs. NGT group