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Maternal lipid profile differs by gestational diabetes physiologic subtype

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Abstract

Aim: To characterize lipid profiles in women with different gestational diabetes mellitus (GDM) physiologic subtypes.

Methods: We measured seven lipid markers (total cholesterol, LDL, HDL, triglycerides, non-esterified fatty acids (NEFA), ApoA, ApoB) in fasting plasma collected in a prospective cohort of 805 pregnant women during second trimester. We estimated insulin sensitivity and secretion using oral glucose tolerance test-based validated indices. We categorized GDM physiologic subtypes by insulin sensitivity and secretion defects defined as values below the 25th percentile among women with normal glucose tolerance (NGT), as previously established. We compared lipid markers across NGT and GDM subtypes. We explored associations between lipid markers and newborn anthropometry in the overall group and stratified by glucose tolerance status.

Results: Among 805 women, 67 (8.3%) developed GDM. Women with GDM had higher body mass index (BMI; 29.3 vs. 26.6 kg/m²), while ethnicity (97.3% vs. 97.0% European ancestry) and age (28 vs. 29 years) were similar. In comparison to women with NGT, women with GDM characterized by a predominant insulin sensitivity defect had significantly higher triglycerides

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Author Contributions

J.C.L., J.W. and M.-F.H. contributed to the study design, analysis and interpretation of data. All authors contributed to manuscript writing/editing and approved of the final submitted version.

Declarations of Interest

None.

(2.20 vs. 1.82, $P=0.002$), lower HDL (1.64 vs. 1.90, $P=0.01$) and higher NEFA (0.34 vs. 0.24, $P<0.0001$). GDM women with a predominant insulin secretion defect differed from women with NGT with respect to NEFA (0.32 vs. 0.24, $P=0.003$) while other lipid markers were similar. These associations remained significant after adjusting for maternal age and BMI. Greater maternal levels of NEFA were associated with higher birth weight z-scores in women with an insulin secretion defect (BMI-adjusted $r=0.58$, $P=0.01$). We did not find significant associations between other lipid markers and newborn anthropometry in other groups.

Conclusion: Women with GDM have distinct lipid profiles based on their GDM physiologic subtype which may not be apparent when investigating GDM as a single group.

Keywords

Gestational diabetes; Lipids; Non-esterified fatty acids; Insulin sensitivity; Insulin secretion

1. Introduction

Pregnancy markedly alters lipid metabolism. The first trimester is characterized by increased lipogenesis; later in pregnancy there is a shift to a more lipolytic state, in accordance with fetal growth requirements [1,2]. Previous studies have suggested that pregnancies with gestational diabetes mellitus (GDM) are characterized by more severe hyperlipidemia as compared to pregnancies in which normal glucose tolerance (NGT) is maintained, but reports are inconsistent regarding which lipids are elevated [3–6]. A possible explanation is that GDM is a heterogeneous condition. We previously demonstrated that GDM can be subdivided into three physiologic subtypes [7]. We also observed that women with an insulin-sensitivity defect had higher complication rates at delivery suggesting other fuels may contribute to macrosomia and related birth complications [7]. However, it is unknown if lipids differ among GDM physiologic subtypes. Thus, the purpose of this study was to characterize gestational lipid profiles among three physiologic GDM subtypes and compare them to NGT women.

2. Methods

2.1. Study Details

This study used data from the Genetics of Glucose regulation in Gestation and Growth (Gen3G) cohort, a prospective cohort study of maternal-child pairs followed from the first trimester through delivery, previously described [8]. The ethics committee at Centre Hospitalier Universitaire de Sherbrooke (CHUS) approved the study and all participants provided written informed consent.

2.2. Gestational Diabetes Mellitus Status

After excluding women with pre-existing diabetes at enrollment, we classified women as having GDM if their second trimester 75 g oral glucose tolerance test (OGTT) based on international criteria [9]. We used plasma glucose and insulin values obtained at 0, 60 and 120 min during OGTT to calculate insulin sensitivity [10] (Matsuda index) and secretion indices [11,12] (Stumvoll first-phase estimate). We defined women as having insulin-

sensitivity defect if their Matsuda insulin sensitivity index was below the 25th percentile of the Matsuda index distribution (GDM-Sensitivity); we defined women as having insulin-secretion defect if their Stumvoll insulin secretion was below the 25th percentile of the Stumvoll estimate distribution (GDM-Secretion) (both distributions determined among NGT women) as previously described [7]. We categorized women with both defects as GDM-mixed.

2.3. Measurements

We obtained 8-hour fasting samples during the second trimester visit for the following biomarkers: total cholesterol (TC), high-density lipoprotein-cholesterol (HDL), low-density lipoprotein-cholesterol (LDL), Apolipoprotein A and B (Apo A and B), triglycerides, non-esterified fatty acids (NEFA), fasting glucose, HbA1c and insulin using methods described elsewhere [8]. The CHUS biomedical laboratory measured fasting TC, HDL, and triglycerides using a colorimetric method (Johnson & Johnson Clinical Diagnostics) and calculated LDL using Friedewald's equation. We measured NEFA levels [minimum level of detection of 0.0014 mM; intra-assay coefficients of variation (CV) 0.8%] using a colorimetric assay (Wako Chemicals). We excluded three NGT women whose LDL >7.0 mmol/L as this was suggestive of familial hypercholesterolemia.

2.4. Statistical Analysis

We presented participants' characteristics as medians (interquartile range) for continuous measures and frequencies for categorical values. We used the rank-sum test to compare the GDM group overall with the NGT group. We used Kruskal-Wallis tests for continuous variables and Fisher exact tests for categorical variables to compare differences across groups. When P values from Kruskal-Wallis or Fisher exact tests were <0.05, we did pairwise comparisons between groups using Dunn's tests [13] or Fisher exact tests with Bonferroni correction of P values. We used linear regression to control for maternal BMI and age, transforming non-normally distributed dependent variables with either the natural-log or square-root transformations to achieve normality prior to regression. We assessed the relationships between NEFA and insulin sensitivity, secretion and oral disposition indices (controlling for second trimester maternal BMI) and all lipids and birth weight using pairwise and partial correlation analyses (controlling for BMI or gestational weight gain). We considered a two-tailed $P < 0.05$ statistically significant. We performed analyses using SAS version 9.4 (Cary, North Carolina) and Stata version 15 (Stata Corp LP, College Station, Texas).

3. Results

We included 805 women in our current analyses, 67 (8.3%) diagnosed with GDM (34 GDM-Sensitivity, 20 GDM-Secretion and 12 GDM-Mixed) and 739 who maintained normoglycemia. GDM-Sensitivity women had a significantly higher BMI as compared to NGT women [7]. There was a higher frequency of previous GDM or macrosomia among GDM-Secretion vs. NGT women (Table 1). Additional subtype differences in characteristics have been reported previously [7].

GDM women had higher median triglycerides and NEFA vs. NGT women: 1.97 vs. 1.82 ($P = 0.02$) and 0.33 vs. 0.24 ($P < 0.0001$), respectively (Table 1). Regarding GDM subtypes, we noted differences in NEFA, triglycerides, HDL, LDL and TC across groups (Table 1 and Fig. 1). Compared to NGT women, GDM-Sensitivity women had lower HDL (1.64 vs. 1.90, $P = 0.01$) and higher triglycerides (2.20 vs. 1.82, $P = 0.002$) and NEFA (0.34 vs. 0.24, $P < 0.0001$). GDM-Sensitivity women also had lower HDL and higher triglycerides than GDM-Secretion women (Fig. 1). These differences remained statistically significant after maternal BMI and age adjustment. In contrast, we did not find significant differences in most lipid markers in GDM-Secretion compared to NGT women except for NEFA (0.32 vs. 0.24, $P = 0.003$). NEFA levels in GDM-Mixed women were similar to other subtypes (Fig. 1) and demonstrated higher levels but not statistically different from NGT women. GDM-Mixed women had higher TC compared to NGT, and higher TC and LDL compared to GDM-Sensitivity and GDM-secretion (Fig. 1); these associations were maintained after adjustments for maternal age and BMI.

To further our understanding of the pathophysiologic role of NEFA in pregnancy, we evaluated the relationship between NEFA and the oral disposition index (DI_o). After controlling for BMI, we observed a weak negative partial correlation in the NGT group ($r = -0.14$, $P = 0.001$). We did not find significant correlations between NEFA and DI_o in overall GDM or in any GDM subtype. We did not find significant associations between NEFA and insulin sensitivity and secretion indices in women with NGT or GDM.

Finally, we explored links between lipids and newborn anthropometry: we did not find significant relationships between triglycerides, HDL, LDL, TC and birth weight by glycemic status (adjusted for maternal BMI or gestational weight gain). However, we found that greater maternal NEFA levels were associated with higher birth weight z-scores in GDM-Secretion subtype (adjusted for BMI or gestational weight gain, $r = 0.58$, $P = 0.01$).

4. Discussion

We found that the lipid profile differs between GDM and NGT women and demonstrated heterogeneity in lipid profiles between GDM physiologic subtypes. Women with GDM-Sensitivity defect had higher triglycerides and lower HDL when compared with NGT or GDM Secretion women. Conversely, the lipid profiles of women with GDM-Secretion defect did not differ significantly from those of NGT women, apart from NEFA which were elevated in all GDM subtypes. TC and LDL were significantly higher in the GDM-Mixed subtype compared to the GDM-Secretion and GDM-Sensitivity subtypes.

By investigating GDM subtypes, we demonstrated striking differences in lipid profiles that were not apparent when investigating GDM as one group, with most prominent differences for women with GDM-Sensitivity defect. Prior literature suggests that NEFA and triglycerides, after crossing the placenta, are released into fetal circulation and may serve as fetal fuel in addition to glucose [14,15]. We expected both to be higher in GDM-Sensitivity, since we observed higher birth weight in this subtype [7]. However, our finding of elevated NEFA across all GDM physiologic subtypes indicates other factors could be influencing

birth weight. Yet we did not observe any significant relationship between birth weight and triglycerides, HDL, LDL and TC, thus other lipids should be investigated.

Study strengths include using validated indices to estimate insulin sensitivity and secretion, and previously characterized GDM physiologic subtypes allowing us to observe distinct lipid profiles. Study limitations include most participants were of European origin, limiting generalizability, and small sample sizes contributing to limited power and lack of statistically significant differences. We did not assess glycemia at the end of pregnancy, but all GDM women were treated at the same multidisciplinary clinic and proportion of insulin treatment did not differ across GDM subtypes [7].

5. Conclusions

Our results demonstrate that GDM physiologic subtypes differ in their gestational lipid profiles, underscoring the presence of physiologic heterogeneity in GDM. Future studies should investigate small lipid fractions and metabolites in GDM physiologic subtypes and their role in fetal overgrowth.

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Abbreviations:

GDM	gestational diabetes mellitus
NGT	normal glucose tolerance
OGTT	oral glucose tolerance test
TC	total cholesterol
HDL	high-density lipoprotein-cholesterol
LDL	low-density lipoprotein-cholesterol
Apo A and B	apolipoprotein A and B
NEFA	non-esterified fatty acid
DI₀	oral disposition index

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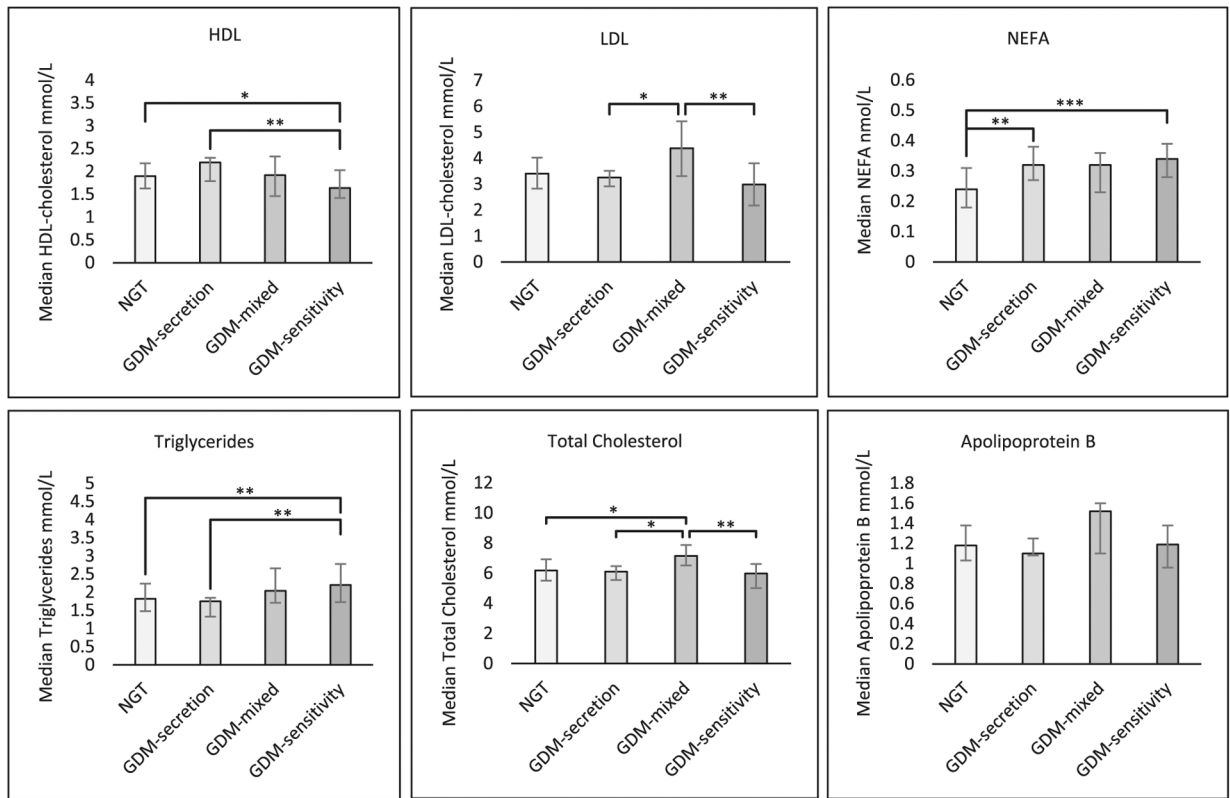


Fig. 1. Median and interquartile range for lipids, by physiological subtype. Pairwise comparisons between groups with statistically significant differences shown by brackets. * P -value < 0.05; ** P -value < 0.01; *** P -value < 0.0001; GDM: gestational diabetes mellitus; NGT: normal glucose tolerance; HDL: high-density lipoprotein-cholesterol; LDL: low-density lipoprotein cholesterol; NEFA: non-esterified fatty acid.

Table 1

Characteristics and lipid values of women with NGT or GDM, by physiological subtype.

	GDM-Secretion Median (IQR)/n (%)	P^b	GDM-Mixed Median (IQR)/n (%)	P^b	GDM-Sensitivity Median (IQR)/n (%)	P^b	NGT Median (IQR)/n (%)	P^d	All GDM Median (IQR) n	P^c
n^d	20 (29.9% of GDM)	-	12 (17.9% of GDM)	-	34 (50.7% of GDM)	-	739 (91.8% of total n)	-	67 (8.3% of total n)	-
Age (years)	30 (28–32)	-	31 (23–33)	-	28 (25–35)	-	28 (25–31)	0.16	29 (26–33)	0.05
% European descent	18 (90.0%)	-	12 (100.0%)	-	34 (100.0%)	-	714 (97.1%)	0.22	65 (97.0%)	0.95
% Nulliparous	5 (25.0%)	-	4 (33.3%)	-	17 (50.0%)	-	382 (51.8%)	0.07	27 (40.3%)	0.07
% Patient history of GDM or macrosomia	8 (40.0%)	0.001	3 (25.0%)	0.23	5 (14.7%)	0.61	62 (8.4%)	0.0001	16 (23.9%)	<0.0001
First-trimester visit										
Gestational age (weeks)	8.8 (7.5–10.8)	-	10.2 (8.2–12.1)	-	9.4 (7.2–11.6)	-	9.2 (8.1–11.4)	0.85	9.3 (7.4–12.0)	0.89
BMI (kg/m ²)	21.9 (20.7–26.0)	0.15	25.3 (21.3–29.8)	0.98	30.1 (26.9–37.7)	<0.0001	23.9 (21.5–27.5)	<0.0001	27.0 (22.0–32.4)	0.002
Second-trimester visit										
Gestational age (weeks)	26.4 (25.9–27.7)	-	26.2 (25.3–27.8)	-	26.3 (25.6–26.6)	-	26.2 (25.6–27.1)	0.65	26.3 (25.6–34.5)	0.91
BMI (kg/m ²)	25.0 (23.7–28.3)	0.21	28.7 (24.6–32.8)	0.65	32.3 (28.9–40.9)	<0.0001	26.6 (24.2–30.1)	<0.0001	29.3 (25.6–34.5)	0.0004
Insulin sensitivity (Matsuda)	7.4 (6.2–8.5)	>0.99	5.4 (4.7–5.5)	0.0003	2.9 (2.3–4.0)	<0.0001	7.9 (5.8–11.2)	<0.0001	4.6 (2.9–6.2)	<0.0001
Insulin secretion (Stumvoll)	594 (501–739)	<0.0001	748 (694–870)	<0.0001	1364 (1063–1677)	0.0003	1121 (934–1286)	<0.0001	969 (676–1368)	0.004
DI ₀	4491 (3947–5318)	<0.0001	4082 (3824–4374)	<0.0001	4038 (3655–4613)	<0.0001	8557 (6797–12,004)	<0.0001	4144 (3760– 4850)	<0.0001
Lipids										
Fasting NEFA (nmol/L)	0.32 (0.27–0.38)	0.01	0.32 (0.23–0.36)	0.17	0.34 (0.28–0.39)	<0.0001	0.24(0.18–0.31)	<0.0001	0.33 (0.26–0.38)	<0.0001
Triglycerides (mmol/L)	1.75 (1.33–1.85)	0.54	2.04 (1.71–2.66)	0.18	2.20 (1.73–2.78)	0.004	1.82 (1.48–2.24)	0.001	1.97 (1.58–2.61)	0.02
HDL cholesterol (mmol/L)	2.20 (1.79–2.30)	0.29	1.92 (1.46–2.33)	>0.99	1.64 (1.42–2.03)	0.01	1.90 (1.63–2.18)	0.01	1.89 (1.46–2.22)	0.28
LDL cholesterol (mmol/L)	3.26 (2.92–3.52)	0.29	4.39 (3.31–5.43)	0.07	2.99 (2.18–3.81)	0.06	3.41 (2.83–4.03)	0.01	3.24 (2.67–3.82)	0.19
Total cholesterol (mmol/L)	6.10 (5.55–6.47)	>0.99	7.15 (6.52–7.88)	0.01	5.98 (5.01–6.61)	0.28	6.18 (5.51–6.92)	0.01	6.23 (5.34–6.96)	0.68
Apolipoprotein A (mmol/L)	2.21 (2.08–2.36)	-	2.16 (2.03–2.35)	-	2.08 (1.84–2.20)	-	2.11 (1.93–2.30)	0.22	2.17 (1.91–2.24)	0.74

	GDM-Secretion Median (IQR)/n (%)	P^b	GDM-Mixed Median (IQR)/n (%)	P^b	GDM-Sensitivity Median (IQR)/n (%)	P^b	NGT Median (IQR)/n (%)	P^a	All GDM Median (IQR)	P^c
Apolipoprotein B (mmol/L)	1.10 (1.08–1.25)	–	1.52 (1.10–1.60)	–	1.19 (0.96–1.38)	–	1.18 (1.03–1.38)	0.18	1.14 (1.08–1.38)	0.98

NGT, normal glucose tolerance; GDM, gestational diabetes mellitus; IQR, interquartile range; GCT, glucose challenge test; DI₀, oral disposition index; NEFA, non-esterified fatty acid.

^aDifferences across groups were compared using the Kruskal-Wallis test for continuous variables and the Fisher exact test for categorical variables.

^bWhen the P -values from the Kruskal-Wallis test or Fisher exact test were <0.05 , pairwise comparisons were made between the NGT group and each of the GDM subtypes using Dunn's test or Fisher exact test. P -values were adjusted using the Bonferroni method and are presented in the third, fifth and seventh columns.

^cDifferences between the All GDM group and the NGT group were compared using the rank sum test with P -values given in the eleventh column.

^dMissing data on second-trimester LDL (n = 34) and second-trimester HDL (n = 28); all other variables had data missing for 5 participants. There were a total of 67 participants with GDM, however one participant could not be categorized into a subtype leaving a total of 66 available for subtype analyses.