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Ethnic differences in glucose disposal, hepatic insulin sensitivity, and endogenous glucose production among African American and European American women

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Abstract

Objective—Intravenous glucose tolerance tests (IVGTT) have demonstrated lower whole-body insulin sensitivity (S_I) among African Americans (AA) compared to European Americans (EA). Whole-body S_I represents both insulin-stimulated glucose disposal, primarily by skeletal muscle, and insulin's suppression of endogenous glucose production (EGP) by liver. A mathematical model was recently introduced that allows for distinction between disposal and hepatic insulin sensitivity. The purpose of this study was to examine specific indexes of insulin sensitivity among AA and EA women to determine whether lower whole-body insulin sensitivity in AA may be attributed to insulin action at muscle, liver, or both.

Methods—Participants were 53 non-diabetic, premenopausal AA and EA women. Profiles of EGP and indexes of Disposal S_I and Hepatic S_I were calculated by mathematical modeling and incorporation of a stable isotope tracer ($6,6\text{-}^2\text{H}_2$ glucose) into the IVGTT. Body composition was assessed by dual energy X-ray absorptiometry.

Results—After adjustment for percent fat, both Disposal S_I and Hepatic S_I were lower among AA ($p=0.009$ for both). Time profiles for serum insulin and EGP revealed higher peak insulin response and corresponding lower EGP among AA women compared to EA.

Conclusions—Indexes from a recently-introduced mathematical model suggest that lower whole-body insulin sensitivity among non-diabetic AA women is due to both hepatic and peripheral components. Despite lower Hepatic S_I , AA displayed lower EGP, resulting from higher post-challenge insulin levels. Future research is needed to determine the physiological basis of lower insulin sensitivity among AA and its implications for type 2 diabetes risk.

Keywords

mathematical modeling; intravenous glucose tolerance test; hepatic glucose production

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Author Contributions: BAG designed the study and supervised all aspects of the project. ACE analyzed the data and prepared the manuscript. JAA performed data analyses and provided input into manuscript preparation. WMG performed mathematical modeling of the data. FO served as study physician and provided conceptual advice. All authors contributed to interpretation and implications of these results as well as review of the manuscript.

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Introduction

Compared to European Americans (EA), African Americans (AA) are twice as likely to be diagnosed with type 2 diabetes (T2D) [1]. In particular, AA women have higher prevalence rates of T2D than any other gender-ethnic group [2]. The reasons for this disparity are not clear, but ethnic differences in tissue sensitivity to insulin may play a role.

Indexes of whole-body insulin sensitivity (S_I) and glucose effectiveness (S_g) can be estimated by mathematical modeling of glucose and insulin values from an intravenous glucose tolerance test (IVGTT) [3,4]. Previous studies that have estimated insulin sensitivity by the traditional one-compartment minimal model have consistently shown lower whole-body insulin sensitivity among healthy AA compared to EA, independent of body composition [5-8]. However, the one-compartment model is not ideal for comparing groups such as EA and AA that differ regarding the magnitude of the post-challenge insulin response [9] due to overestimation of the effects of glucose itself (S_g) in subjects with a higher insulin response [10,11]. Further, the one-compartment minimal model assesses insulin sensitivity only at the whole-body level. Whole-body insulin sensitivity comprises both insulin action to promote glucose uptake as well as to suppress endogenous glucose production by the liver [12]. Whether lower insulin sensitivity among AA represents impairment in glucose uptake or hepatic glucose production is not known. Discerning whether lower insulin sensitivity among AA relates to hepatic or peripheral glucose regulation could help elucidate the etiology behind racial/ethnic differences in T2D.

When a stable isotope tracer is included in the IVGTT, a two-compartment model can be used to provide an index of disposal-specific insulin sensitivity and a more reliable index of glucose effectiveness [13,14]. In 2005, researchers expanded on the traditional two-compartment minimal model by describing endogenous glucose production (EGP) from tracer-labeled IVGTT data [15]. In 2009, Tokuyama and colleagues further developed the two-compartment model framework by introducing model-derived insulin sensitivity indexes specific to hepatic glucose regulation [16]. Thus, these latest advances in two-compartment modeling offer the opportunity to provide novel data comparing disposal-specific insulin sensitivity, hepatic insulin sensitivity, and hepatic glucose production in AA and EA.

The aim of this study was to describe insulin action and hepatic glucose metabolism among non-diabetic, premenopausal women of AA and EA ancestry using a recently-introduced integrated model [16] with the goal of determining whether lower insulin sensitivity among AA may be attributed to either hepatic or peripheral sensitivity to insulin.

Methods

Participants

Fifty-three non-diabetic, premenopausal women were categorized by ethnicity as African American (AA) or European American (EA) based on self-report and assertion that both parents shared the same ethnicity as the participant. Normal glucose tolerance was verified by 2-hour oral glucose tolerance testing, and premenopausal status was confirmed by serum follicle stimulating hormone concentration < 35 IU/mL in addition to self-report of regular menstrual cycles. Exclusion criteria included history of polycystic ovary disease, hypoglycemia, and any medication known to influence glucose metabolism (including oral contraceptives). All participants provided oral and written consent, and the study was approved by the Institutional Review Board for Human Use at the University of Alabama at Birmingham (UAB).

Protocol

For three days prior to testing, participants were instructed to consume ~250 g carbohydrates. The evening before testing, they reported to the General Clinical Research Center (GCRC). After a 12-hour overnight fast, metabolic parameters were determined by insulin-modified intravenous glucose tolerance test (IVGTT). Body composition was assessed by dual energy X-ray absorptiometry (DXA; Lunar Prodigy, GE Healthcare, Madison, WI), and scans were analyzed with software version 1.5.

Intravenous Glucose Tolerance Test (IVGTT)

For the IVGTT, flexible intravenous catheters were placed in the antecubital spaces of both arms. Blood was sampled 3 times over 15 minutes, and averages of these three fasting samples were used to determine fasting glucose, insulin, and C-peptide concentrations. At time zero, a bolus of glucose (50% dextrose, 270 mg/kg, plus [6,6-²H₂]glucose, 30 mg/kg) was infused intravenously. Insulin (0.02 U/kg) was administered intravenously over a 5-minute period from 20-25 minutes after the glucose injection. Blood was sampled at the following times (minutes) following glucose administration: 2, 3, 4, 5, 6, 8, 10, 12, 15, 19, 20, 21, 22, 24, 26, 28, 30, 35, 40, 45, 50, 55, 60, 70, 80, 100, 120, 140, 180, 210, 240, for a total of 34 samples). Sera were stored at -85°C until laboratory analysis of glucose and insulin.

AIR_g, the integrated incremental area under the curve for insulin during the first 10 min of the IVGTT, was calculated using the trapezoidal method. An integrated two-compartment mathematical model [16] was used to estimate the time course of endogenous glucose production (**EGP**) in mg/kg/min. EGP primarily reflects hepatic glucose production [17,18]. Indexes of insulin sensitivity specific to glucose disposal (**Disposal S_i**) and hepatic insulin action (**Hepatic S_i**) were calculated [16]. The same model was used to derive indexes of disposal and hepatic glucose effectiveness (**Disposal S_g** and **Hepatic S_g**) as estimates of the ability of glucose itself to promote disposal and suppress EGP, respectively. Details of the integrated two-compartment minimal model have been previously described by Tokuyama et al [16], and indexes from the model are summarized by the following equations:

$$EGP(t) = k_{out} [1 - H_2(t)] G_L(t)$$

$$Disposal S_g = S_g^{2*} = V_1 (k_p + k_{21}k_{02} / (k_{02} + k_{12}))$$

$$Disposal S_i = S_i^{2*} = V_1 s_k k_{21} k_{12} / (k_{02} + k_{12})^2$$

$$Hepatic S_g = hS_g^2 = -\partial EGP(t) / \partial G(t) = k_{in} / 4G_b$$

$$Hepatic S_i = hS_i^2 = \partial^2 EGP(t) / \partial G(t) \partial I(t) = S_i^{2*} * hS_g^2 / (S_g^{2*} + hS_g^2)$$

IC_{50} is representative of insulin action required for 50% inhibition of EGP. H_2 represents a function of insulin's inhibitory effect on EGP; $H_2 (= x(t)/[IC_{50} + x(t)])$, and G_L signifies available glucose in the liver.

V_1 is the volume of the accessible compartment, s_k is a parameter of insulin action, and k_p , k_{02} , k_{12} , and k_{21} are rate constant parameters.

Basal glucose is quantified by G_b , while k_{in} and k_{out} are rate constants of hepatic glucose production and hepatic glucose loss, respectively.

Data were modeled using SCIENTIST software (version 2.01; St. Louis, MO). For each participant, Disposal S_I and Hepatic S_I from the 2-compartment model were summed to yield a value for **Total S_I** . A surrogate index for fasting hepatic insulin resistance was calculated by multiplying basal EGP \times fasting insulin concentration as previously described [19].

Laboratory Analyses

All analyses were performed in the Core Laboratories of the General Clinical Research Center (GCRC), Nutrition Obesity Research Center (NORC), and Diabetes Research and Training Center (DRTC). Glucose was measured in 10 μ L sera using the Ektachem DT II analyzer (Johnson and Johnson Clinical Diagnostics, Rochester, NY). This analysis had a mean intra-assay coefficient of variation (CV) of 0.61%, and a mean inter-assay CV of 1.45%. Insulin was measured by radioimmunoassay (Linco Research Inc., St. Charles, MO; now Millipore Corporation, Billerica, MA). This assay has a sensitivity of 3.35 μ IU/ml, mean intra-assay CV of 3.49%, and mean interassay CV of 5.57%. C-peptide was measured by radioimmunoassay (Siemens Healthcare Diagnostics, Los Angeles, CA) in duplicate 25 μ L aliquots. Sensitivity for this assay is 0.318 ng/mL, mean intra-assay CV is 3.57%, and mean interassay CV is 5.59%. Serum concentrations of free fatty acids (FFA) were measured with "NEFA-C" assays (Wako Diagnostics, Richmond, VA [20]). This assay has an intra-assay CV was 3.89%, and the inter-assay CV is 5.87%. Minimum assay sensitivity was 0.0014mEq/L. [6,6- 2 H $_2$]glucose enrichment for each of the 34 blood samples listed above were analyzed by gas chromatography mass as previously described [21]. Briefly, serum samples were deproteinized, evaporated, and prepared with N,O-bis[Trimethylsilyl]trifluoroacetamide (BSTFA) and 1% trimethylchlorosilane (TMSC). Derivatives were analyzed on an Agilent 6890 gas chromatograph coupled to a 5973 mass spectrometer in Electron Impact mode. This analysis uses a standard curve prepared with in-house control serum samples and monitoring of M+0 and M+2 ions. Mole fractions were calculated from total area counts. CV of the [6,6- 2 H $_2$]glucose among the fasting samples was 1.75%.

Statistical Analysis

Ethnic group differences for age, body mass index (BMI), k_{in} , k_{out} , and IC_{50} were compared by nonparametric Mann Whitney-U tests, percentages of overweight subjects in each group were compared by the Fisher exact test, and independent t-tests were used to determine between-group differences for all other variables of interest. Variables were Log_{10} transformed for normality when appropriate. ANCOVA analyses were performed to examine group differences in S_I and S_g indexes with adjustment for % body fat.

Mann Whitney-U tests and independent t-tests were used to identify ethnic differences in glucose, insulin, and EGP at individual time points over the duration of the IVGTT. Composite scores for glucose, insulin, and EGP time courses were calculated as incremental area under the curve (AUC) by the trapezoidal method.

Statistical tests were performed with SPSS software (version 19.0; Chicago, IL) and GraphPad Prism (version 5.0; La Jolla, CA). All tests were two-sided with a Type I error rate of 0.05.

Results

Participant characteristics and metabolic parameters are displayed as mean \pm SD by ethnic group in Table 1. Groups were similar in age, body weight, and body composition. Both Hepatic S_I and Disposal S_I were lower for AA ($p=0.009$ for both), and these ethnic differences intensified with adjustment for body composition. On average, Hepatic S_I accounted for approximately 30% Total S_I in both EA (range: 24.5 – 39.6) and AA (range 22.2 – 42.2). Hepatic S_g did not differ between ethnic groups, but Disposal S_g was significantly lower among AA independent of body composition ($p=0.020$). AIR $_g$ was more than two-fold higher among AA ($p<0.001$), and after adjustment for percent body fat, the surrogate index of fasting hepatic insulin resistance [19] was higher for AA ($p=0.048$). Time profiles for serum total glucose, glucose enrichment, insulin, C-peptide, and EGP are shown in Figure 1. Total glucose concentrations did not differ between groups at any point during the test (panel A), but AA demonstrated higher peak insulin levels (panel C) ($p<0.05$ for minutes 3-22 following the glucose challenge). AUC for insulin was also greater for AA compared to EA ($p=0.01$). C-peptide measurements were available for 50 of the 53 participants. Panel D displays higher post-challenge C-peptide concentrations among AA ($p<0.05$ for minutes 4-10). EGP among AA was significantly lower ($p<0.05$) from minutes 2 to 22 of testing (Panel E). Although total AUC of EGP did not differ between ethnic groups, AUC for the first 30 minutes of testing was significantly lower for AA $p=0.016$). Serum concentrations of FFA were similar between groups at baseline, and circulating FFA did not differ between groups at any point of blood sampling (data not shown).

Discussion

It is well-established that AA are at higher risk than EA for developing T2D. Previous studies quantifying insulin sensitivity by IVGTT and the traditional one-compartment minimal model [22] have repeatedly reported lower whole-body S_I among AA participants, independent of body composition [5-8]. However, T2D is a disease involving multiple organs and tissues, and whole-body S_I represents a composite of both peripheral and hepatic components. Whether lower S_I among AA relates to lower Disposal S_I (insulin-mediated glucose uptake), lower Hepatic S_I (suppression of EGP in response to insulin concentration), or both has not previously been investigated. The results presented here indicate that *both* Disposal S_I and Hepatic S_I were lower among a cohort of healthy, premenopausal AA vs EA women. Additionally, a surrogate index of basal hepatic insulin resistance was higher among AA. However, despite the lower Hepatic S_I , AA demonstrated lower model-derived EGP following a glucose challenge. Lower EGP among AA was concomitant with higher insulin and C-peptide concentrations. The physiological basis for lower insulin sensitivity in AA remains to be determined.

This study provides a novel contribution to the literature by comparing specific indexes of S_I between EA and AA. Previous studies describing lower whole-body S_I among AA by one-compartment modeling [5-8] were unable to tease apart specific effects of insulin on glucose disposal and glucose production, and inherent limitations of the one-compartment model warrant caution about its estimation of glucose effectiveness (S_g) [9-11]. Incorporation of a stable isotope tracer of glucose into the IVGTT in combination with two-compartment modeling provides a more accurate estimate of S_g as well as discrimination of disposal-specific S_I and S_g [12-14]. Based on a description of endogenous glucose kinetics by Krudys et al [15], Tokuyama et al recently expanded modeling of the glucose system to

provide new indexes of Hepatic S_I and Hepatic S_g that correspond to the effects of insulin and glucose on EGP specifically [16]. Using this new integrated mathematical model, we report for the first time that both Disposal S_I and Hepatic S_I appear lower in AA compared to EA. Moreover, although the ability of glucose per se to suppress EGP (Hepatic S_g) did not differ between groups, the action of glucose to facilitate its own disposal (Disposal S_g) appeared lower among AA.

Lower Disposal S_I and Disposal S_g among AA women may relate to inherent differences in aspects of skeletal muscle function. A previous study among premenopausal women reported lower muscle oxidative capacity among AA vs. EA, as well as an independent correlation between in vivo mitochondrial function and whole-body S_I [23]. Our group also recently demonstrated that disposal-specific S_I was independently associated with systemic markers of oxidative stress in AA but not EA [24]. In contrast, infiltration of skeletal muscle by lipid was related to decreased insulin sensitivity among EA but not AA [25-27]. Taken together, these observations suggest that mitochondrial dysfunction and consequent oxidative stress may compromise muscle function and consequent glucose disposal in AA women.

The source of lower Hepatic S_I among AA is not clear. Previous studies in animals [28] and humans [29,30] have related compromised hepatic insulin resistance to accumulation of visceral adipose tissue, with increased free fatty acids into portal circulation as a proposed mechanism [30,31]. However, AA have been shown to have lower hepatic triglyceride content [32,33], and it is well-established that AA women tend to deposit less fat as visceral adipose [5,34-37]. Among our cohort, neither fasting nor post-challenge FFA concentrations differed between AA and EA, and ethnic differences in Hepatic S_I intensified with adjustment for percent body fat. Although it is possible that hepatic triglycerides or portal FFA may have contributed to differences between groups, these analyses were beyond the scope of this study. Thus, while it seems likely that lower Hepatic S_I among AA is attributable to factors other than body composition, the source of this difference awaits further study.

Despite significantly lower Hepatic S_I , time course comparisons revealed lower EGP among AA during the early period of testing. Little data are available regarding ethnic differences in EGP, but one study examining basal hepatic glucose production among obese adolescents similarly reported lower EGP in AA compared to EA [38]. Although the physiological mechanism for lower EGP in AA is not known, higher insulin is likely influential. The model-derived estimate of EGP involves both IC_{50} and insulin response. IC_{50} did not significantly differ between the two groups, suggesting that greater suppression of EGP resulted from the relatively robust early insulin response of AA. Fasting insulin concentration was higher in AA vs EA in this sample, and average AIRg was approximately two-fold higher in AA, as previously reported [6,8,39]. First-phase insulin secretion has been identified as a determinant of EGP [40], and higher C-peptide concentrations in AA concurrent with higher AIRg suggests that early insulin secretion was higher in AA vs EA [41]. Further research is indicated to explain the inter-relationships of fasting and post-challenge insulin and EGP as well as the physiological significance of these relationships for diabetes risk within healthy, non-diabetic individuals.

Major strengths of this study were incorporation of a stable isotope tracer and application of recent advances in mathematical modeling to differentiate peripheral and hepatic insulin sensitivity. However, this mathematical model has yet to be validated against model-independent measures of insulin sensitivity, so our analysis may be subject to limitations inherent to model-derived indexes of insulin sensitivity [9,12]. Additional limitations included modest sample size and cross-sectional design. Although we adjusted for percent

body fat and confirmed equal distribution of normal weight and overweight women in each group, we lacked statistical power for further subgroup analysis of normal weight vs overweight participants.

In conclusion, present results indicate that lower whole-body insulin sensitivity previously reported among healthy AA may be due to both disposal and hepatic components. Further research is indicated to determine why AA women exhibit lower insulin sensitivity and higher risk for T2DM despite their lower post-challenge EGP, and future studies using model-independent methods are needed to determine the physiological basis of lower insulin sensitivity among AA.

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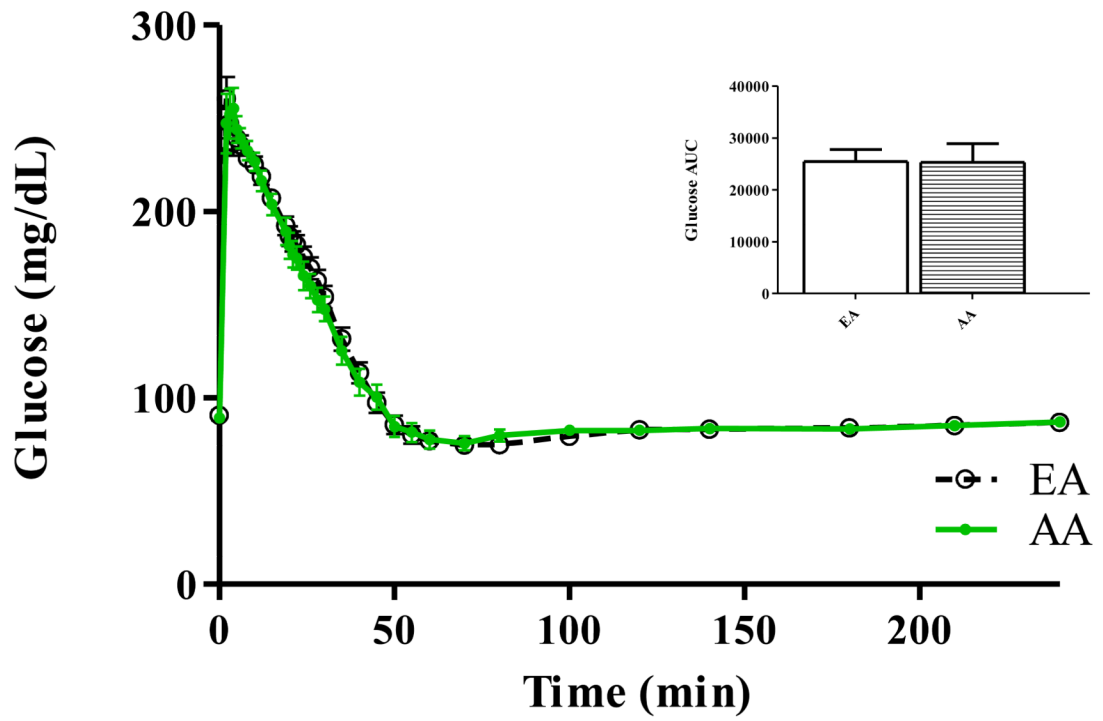
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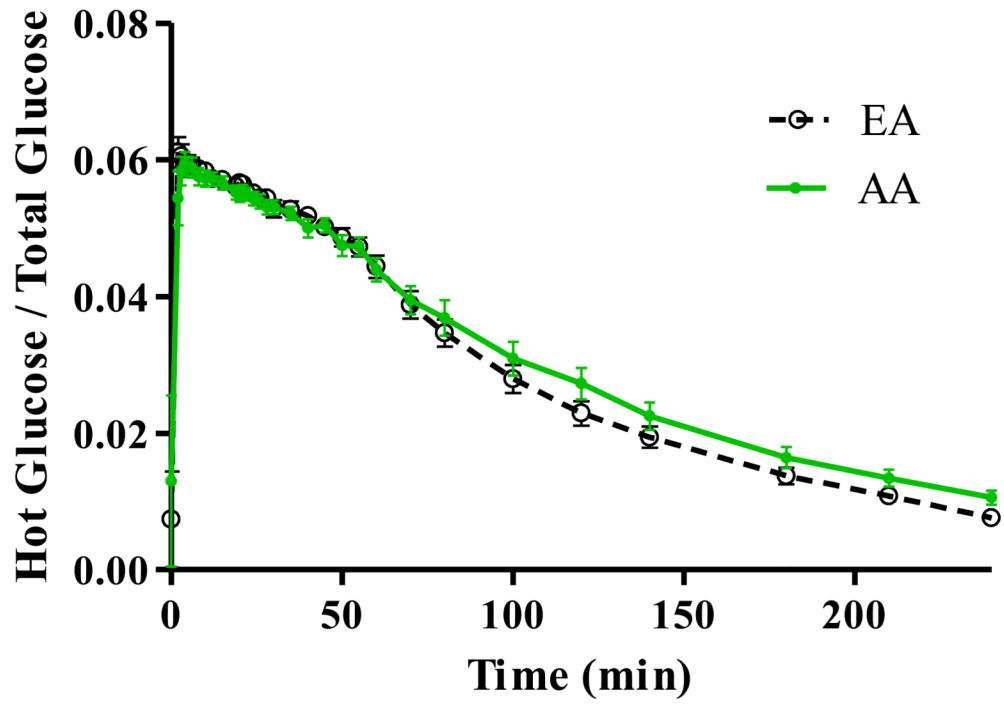
Abbreviations

AA	African Americans
EA	European Americans
IVGTT	intravenous glucose tolerance tests
EGP	endogenous glucose production
T2D	type 2 diabetes
S_I	insulin sensitivity
S_g	glucose effectiveness
AIR_g	acute insulin response to glucose
BMI	body mass index
FFM	fat-free mass
FM	at mass
FFA	free fatty acids

A)

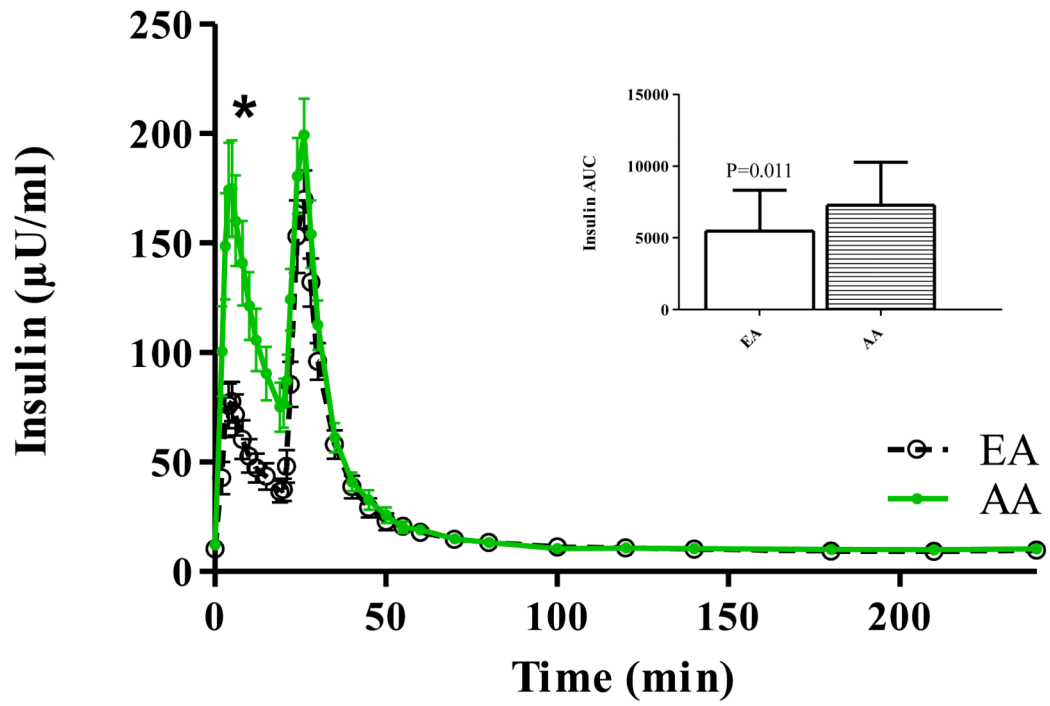
Glucose time course



B)**Enrichment ratio time course**

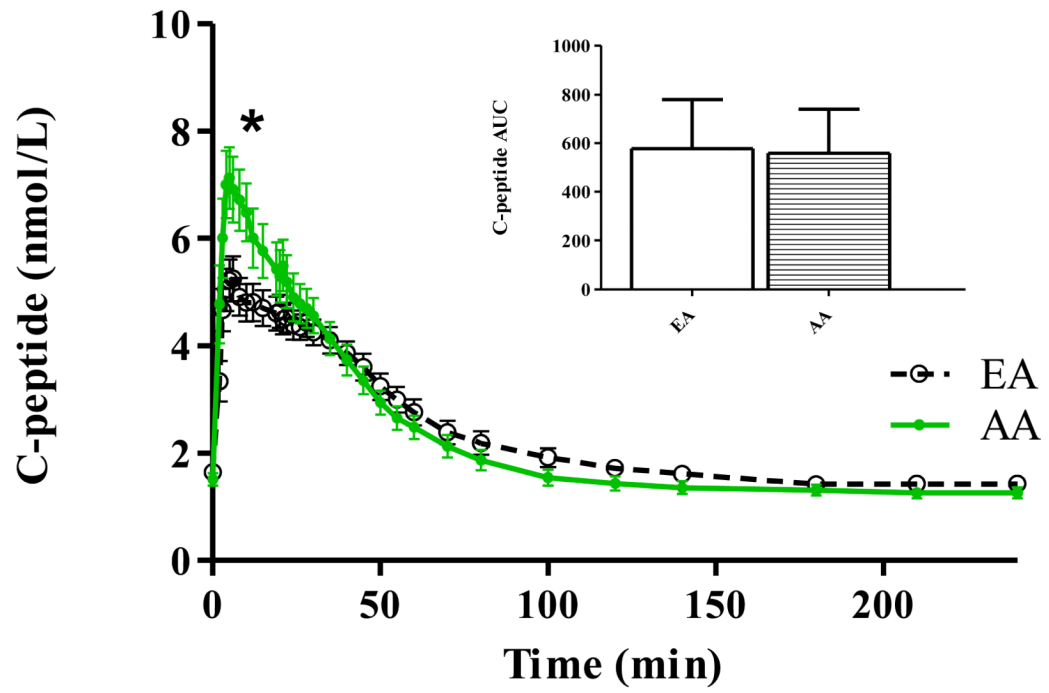
C)

Insulin time course



D)

C-peptide time course



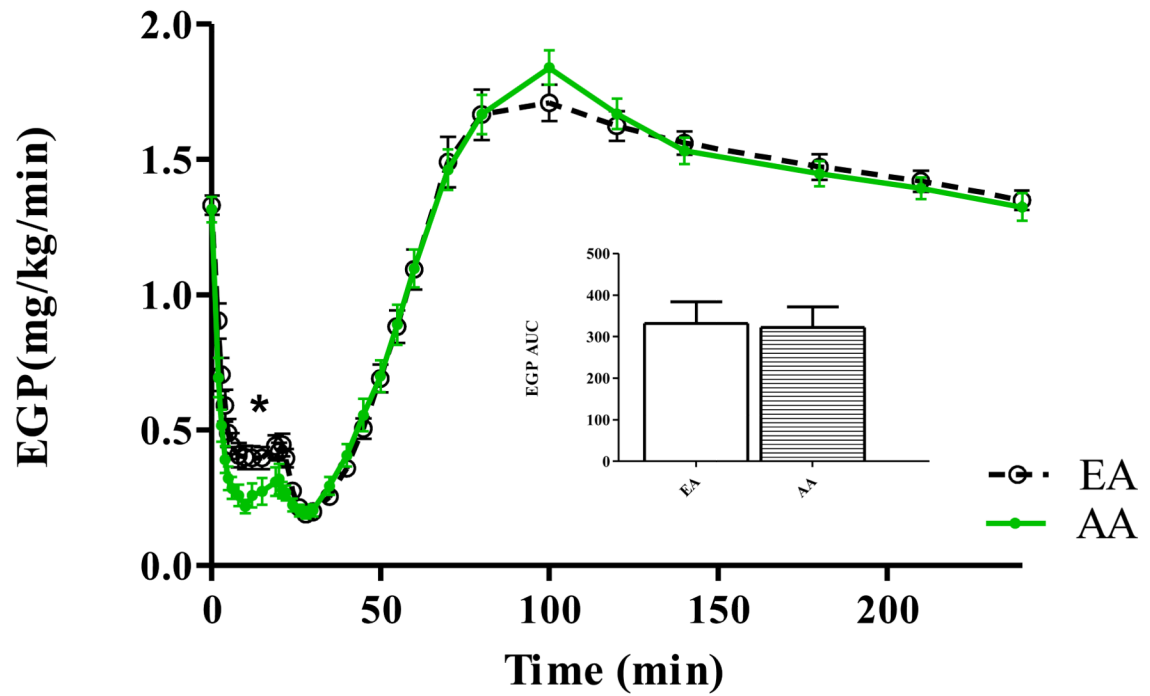
E)**Endogenous glucose production time course**

Figure 1. Time courses for glucose (A), isotope enrichment (B), insulin (C), C-peptide (D), and EGP (E). *P < 0.05.

Table 1
Participant characteristics and metabolic parameters

	EA (n = 30)	AA (n = 23)	P-value	P-value adjusted for %FAT
Age	26.02 ± 3.45 (19.3 – 32.9)	24.27 ± 4.13 (18.0 – 29.9)	0.141	
Weight (kg)	69.86 ± 12.82 (52.6 – 103.0)	73.27 ± 16.05 (51.9 – 111.8)	0.442	
BMI (kg/m ²)	25.22 ± 4.33 (18.7 – 35.2)	26.62 ± 5.84 (18.5 – 38.6)	0.440	
BMI > 25 (%)	(43.3%)	(43.5%)	1.000	
FFM (kg)	41.14 ± 5.09 (32.6 – 49.7)	42.72 ± 4.32 (36.1 – 50.9)	0.212	
FM (kg)	25.56 ± 9.79 (9.8 – 50.9)	27.06 ± 13.36 (10.9 – 57.8)	0.954	
Percent body fat (%)	35.92 ± 7.91 (16.4 – 49.9)	35.31 ± 10.34 (21.0 – 52.0)	0.814	
k_{in} (mg/kg/min)	2.66 ± 0.40 (2.23 – 3.82)	2.63 ± 0.45 (1.47 – 3.73)	0.872	
k_{out} (min ⁻¹)	0.027 ± 0.004 (0.020 – 0.038)	0.027 ± 0.005 (0.021 – 0.043)	0.693	
IC ₅₀ (min ⁻¹)	0.013 ± 0.002 (0.011 – 0.018)	0.016 ± 0.012 (0.010 – 0.071)	0.844	
Basal FFA (mEq/L)	0.50 ± 0.16 (0.13 – 0.79)	0.56 ± 0.15 (0.38 – 0.96)	0.143	0.141
Basal glucose (mg/dL)	90.71 ± 6.32 (77.0 – 105.0)	88.86 ± 7.97 (79.0 – 112.6)	0.311	0.326
Basal insulin (μU/mL)	10.29 ± 4.56 (4.0 – 26.0)	12.18 ± 4.26 (7.0 – 23.2)	0.062	0.027
Basal C-peptide (nmol/L)	1.64 ± 0.48 (0.64 – 2.66)	1.52 ± 0.52 (0.66 – 2.72)	0.359	0.328
AIRg (μU/mL × 10 min)	481.45 ± 384.28 (66.8 – 1958.5)	1178.03 ± 799.80 (262.0 – 3080.7)	<0.001	<0.001
Basal EGP (mg/kg/min)	1.33 ± 0.20 (1.12 – 1.91)	1.32 ± 0.23 (0.73 – 1.86)	0.702	0.737
Basal EGP × basal insulin	13.59 ± 6.02 (5.47 – 35.17)	16.10 ± 6.64 (7.68 – 35.87)	0.112	0.048
Disposal S _g (×10 ² /min)	0.736 ± 0.235 (0.337 – 1.342)	0.586 ± 0.189 (0.249 – 1.214)	0.020	0.008
Hepatic S _g (×10 ² /min)	0.57 ± 0.09 (0.460 – 0.871)	0.538 ± 0.109 (0.315 – 0.838)	0.156	0.152
Disposal S _I (×10 ⁴ /min/μU/ml)	10.51 ± 4.54 (3.49 – 20.53)	7.46 ± 3.81 (1.66 – 20.31)	0.009	0.002
Hepatic S _I (×10 ⁴ /min/μU/ml)	4.46 ± 1.71 (2.29 – 10.22)	3.42 ± 1.34 (0.94 – 6.52)	0.009	0.005
Total S _I (×10 ⁴ /min/μU/ml)	14.97 ± 6.13 (5.77 – 30.75)	10.88 ± 4.97 (2.60 – 26.13)	0.009	0.002

(Mean ± SD and Range); BMI = body mass index, FFM = fat-free mass, FM = fat mass, FFA = free fatty acids, AIRg = acute insulin response to glucose, EGP = endogenous glucose production, k_{in} = rate constant of hepatic glucose production, k_{out} = rate constant of hepatic glucose loss, IC₅₀ = insulin action required to suppress EGP by 50%, S_g = glucose effectiveness index, S_I = insulin sensitivity index, Total S_I = (Disposal S_I + Hepatic S_I)