



Improved Detection of Abnormal Glucose Tolerance in Africans: The Value of Combining Hemoglobin A_{1c} With Glycated Albumin

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OBJECTIVE

In African-born Blacks living in America, we determined by BMI category 1) prevalence of abnormal glucose tolerance (Abnl-GT) and 2) diagnostic value and reproducibility of hemoglobin A_{1c} (HbA_{1c}), fructosamine, and glycated albumin (GA).

RESEARCH DESIGN AND METHODS

Participants ($n = 416$; male, 66%; BMI 27.7 ± 4.5 kg/m² [mean \pm SD]) had an oral glucose tolerance test with HbA_{1c}, GA, and fructosamine assayed. These glycemic markers were repeated 11 ± 7 days later. Abnl-GT diagnosis required 0 h ≥ 5.6 mmol/L (≥ 100 mg/dL) and/or 2 h ≥ 7.8 mmol/L (≥ 140 mg/dL). Thresholds for HbA_{1c}, GA, and fructosamine were the values at the 75th percentile for the population (39 mmol/mol [5.7%], 14.2%, and 234 μ mol/L, respectively).

RESULTS

Abnl-GT prevalence in the nonobese was 34% versus 42% in the obese ($P = 0.124$). Reproducibility was excellent for HbA_{1c} and GA (both $\kappa \geq 0.8$), but moderate for fructosamine ($\kappa = 0.6$). Focusing on HbA_{1c} and GA in the nonobese, we found as single tests the sensitivities of HbA_{1c} and GA were 36% versus 37% ($P = 0.529$). Combining HbA_{1c} and GA, sensitivity increased to 58% because GA identified 37% of Africans with Abnl-GT not detected by HbA_{1c} (P value for both tests vs. HbA_{1c} alone was <0.001). For the obese, sensitivities for HbA_{1c}, GA, and the combined tests were 60%, 27%, and 67%, respectively. Combined test sensitivity did not differ from HbA_{1c} alone ($P = 0.25$) because GA detected only 10% of obese Africans with Abnl-GT not detected by HbA_{1c}.

CONCLUSIONS

Adding GA to HbA_{1c} improves detection of Abnl-GT in nonobese Africans.

Mathematical models predict that between 2019 and 2045, sub-Saharan Africa will experience a 143% increase in the prevalence of abnormal glucose tolerance (Abnl-GT) (1). This increase in Abnl-GT, which encompasses both prediabetes and type 2 diabetes (T2D), is the highest anticipated increase in the world (1). Slowing this upward trajectory requires strategies for the diagnosis of Abnl-GT that are feasible and effective in Africa.

Another challenge is that 60–80% of Africans living with Abnl-GT are undiagnosed (1). Hence, Africa has the highest prevalence in the world of people living with

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undiagnosed Abnl-GT (1). Even when undetected, complications from Abnl-GT progress. Furthermore, Abnl-GT is associated with greater susceptibility and higher mortality from infectious diseases, including tuberculosis and coronavirus disease 2019 (2,3). Therefore, lowering the prevalence of Abnl-GT may assist in decreasing the scope and consequences of several important infectious diseases. However, the International Diabetes Federation reports that <20% of African countries have in-country data on the prevalence of Abnl-GT (1).

The challenges of collecting data are magnified by the fact that routine tests used for Abnl-GT screening, such as fasting plasma glucose (FPG) and hemoglobin A_{1c} (HbA_{1c}), have poor diagnostic sensitivities ($\leq 50\%$) in Africans (4,5). Thus, even when screening programs exist, Africans with Abnl-GT are often misdiagnosed as normal (6).

As a screening test, FPG is suboptimal. Fasting is difficult to achieve. People often travel many miles and wait for extended periods before being seen in clinics. In addition, people carry their food with them so they can easily eat when they are hungry. As going to a medical clinic and being fasted is not standard practice in many medical clinics in African countries, our focus is on the identification of effective nonfasting screening tests. Furthermore, blood samples kept at room temperature for even 30 min undergo extracorporeal glycolysis, resulting in the reporting of spuriously low glucose concentrations (7). HbA_{1c} is also problematic because, even in the absence of factors that adversely affect it, such as nutritional deficiencies, hemoglobinopathies, and anemia, HbA_{1c} detects <50% of Africans with Abnl-GT (6). The diagnostic sensitivity of HbA_{1c} is also <50% in African Americans, Whites, Hispanics, and Arab immigrants to the U.S. (8–10).

Due to the inadequate diagnostic performance of FPG and HbA_{1c}, attention has turned to fructosamine and glycated albumin (GA), both of which are nonfasting markers of glycemia. Fructosamine reflects the concentration of all circulating glycated proteins, including GA, which is formed by the nonenzymatic attachment of glucose to albumin (11).

A previous study of 236 African-born Blacks enrolled in the Africans in America study revealed that combining GA with HbA_{1c} doubled the detection of

prediabetes in the nonobese (12). The prevalence of Abnl-GT in the nonobese needs special attention because in low- and middle-income countries globally and in Africa, the prevalence of T2D is rapidly rising in the nonobese (1,13,14). To pursue improved detection of hyperglycemia in Africans, we increased recruitment to the Africans in America cohort and broadened our detection goal from prediabetes to Abnl-GT. With this larger cohort of 416 African-born Blacks living in America, our objectives were to determine by BMI category: 1) the prevalence of Abnl-GT, and 2) the diagnostic value and reproducibility of HbA_{1c}, GA, and fructosamine.

RESEARCH DESIGN AND METHODS

Population

The Africans in America cohort assesses the cardiometabolic health of African-born Blacks living in the U.S. (15,16). Recruitment is by newspaper advertisements, flyers, community event presentations, and relevant websites. The National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK) Institutional Review Board (ClinicalTrials.gov identifier NCT00001853; Bethesda, MD) approved the study. Written informed consent is obtained prior to enrollment.

During a telephone screening interview, potential enrollees must state they were born in sub-Saharan Africa to two Black parents who were also born in sub-Saharan Africa. Additionally, they must self-identify as healthy and deny a history of diabetes.

A total of 451 African-born Blacks living in metropolitan Washington, DC, successfully completed the telephone interview and had an outpatient screening visit at the National Institutes of Health (NIH) Clinical Center (Supplementary Fig. 1). A history, physical, electrocardiogram, and routine blood tests were performed. Thirty-three individuals did not proceed to an oral glucose tolerance test (OGTT). Exclusion criteria were: anemia ($n = 10$), elevated liver transaminases ($n = 1$), hypothyroidism ($n = 1$), intravenous access issues ($n = 4$), and scheduling conflicts ($n = 17$). Two individuals were excluded after the OGTT because hemoglobin electrophoresis revealed hemoglobin type AF. The percent hemoglobin F in one participant was 15% and for the other person 25%.

Hemoglobin F at these levels interferes with the determination of HbA_{1c} by the high-performance liquid chromatography (HPLC) used in this study.

OGTTs

OGTTs were performed in 416 individuals (male, 66%; aged 39 ± 10 years [mean \pm SD], range 20–65 years; BMI 27.7 ± 4.5 kg/m², range 18.2–42.2 kg/m²).

After a 12-h fast, participants came to the NIH Clinical Center at 7:00 A.M. On arrival, women provided a urine sample for pregnancy testing, and all were negative. As described previously, weight, height, and waist circumference (WC) were measured (15). Weight was obtained with a calibrated digital scale (Scale-Tronix 5702; Welch Allyn, Skaneateles Falls, NY). Height was measured with a wall stadiometer (Seca 242; Seca Corp., Hanover, MD). BMI was calculated as weight in kilograms divided by height in meters squared. BMI categories were defined according to World Health Organization guidelines (17). WC was measured at the superior border of the iliac crest at the end of expiration using a stretch-resistant tape measure with the person standing with feet hip-width apart and weight evenly distributed. The mean of three values was recorded.

Baseline blood samples were obtained for HbA_{1c}, hemoglobin electrophoresis, glucose, insulin, GA, and fructosamine measurements. This was followed by an OGTT (TRUTOL 75 g; Custom Laboratories) with samples taken at 0.5, 1, and 2 h for glucose and insulin concentrations. Abnl-GT was defined as FPG ≥ 5.6 mmol/L and/or 2-h glucose ≥ 7.8 mmol/L (1).

After the OGTT, a computed tomographic scan (Siemens and SOMATOM Force Scanner) was performed to measure visceral adipose tissue (VAT).

Reproducibility of HbA_{1c}, GA, and Fructosamine Values

To determine the reproducibility of HbA_{1c}, GA, and fructosamine, 36% (150 out of 416) of enrollees returned to the Clinical Center 11 ± 7 days after the OGTT.

Participants were divided into two groups (A and B) (Supplementary Fig. 1).

Group A comprised the first 281 consecutively enrolled individuals. In group A, repeat studies were planned only if the initial OGTT met the glucose criteria for T2D (1). Seventeen out of the 18 individuals newly diagnosed with T2D returned

for a second visit, and 1 declined. All 17 individuals had HbA_{1c} levels determined, but GA and fructosamine concentrations were available only in the last 5 consecutively enrolled participants.

Group B consisted of the 135 subsequently enrolled individuals. All were invited, independent of glucose tolerance status at the initial OGTT, for a repeat study. Two individuals declined repeat studies. For one individual who did return, the blood sample obtained for HbA_{1c} clotted. Therefore, for group B, duplicate HbA_{1c}, GA, and fructosamine were available for 132, 133, and 133 individuals, respectively.

Overall, repeat values were available for HbA_{1c}, GA, and fructosamine in 149 (17 plus 132), 138 (5 plus 133), and 138 (5 plus 133) individuals, respectively.

Metabolic Parameters

Degree of glycemia was measured by using the trapezoidal rule for area under the curve (AUC) for glucose during the OGTT. Insulin was assessed by the Matsuda Index, insulin secretion by AUC for insulin divided by AUC for glucose, and β -cell function by the Oral Disposition Index (18,19).

Assays

Glucose and hs-CRP were measured in plasma and insulin in serum with a Roche Diagnostics cobas 6000 analyzer.

Hemoglobin, hematocrit, white blood cells, and mean corpuscular volume were measured in EDTA-anticoagulated whole blood using a Sysmex XE-5000.

HbA_{1c} by HPLC

HbA_{1c} values were determined with two different NGSP-certified instruments using HPLC technology manufactured by BioRad Laboratories. HbA_{1c} samples from the first 139 enrollees were measured on the VARIANT II instrument. The next 277 participants had HbA_{1c} measurements performed on a D-10 instrument. The correlation (R^2) and mean bias between the VARIANT II and D-10 instruments were 0.9934 and 0.07 (1.21%), respectively.

Fructosamine and GA

Fructosamine and GA were measured in plasma on the cobas 6000. For fructosamine, a colorimetric nitroblue tetrazolium assay was used. Interassay coefficient of variation for fructosamine

was 2.9% at 308 μ mol/L and 2.6% at 521 μ mol/L (12). For GA, the Lucica GA-L enzymatic assay, provided by Asahi Kasei Pharma Corporation (Tokyo, Japan), was used. Albumin was measured with bromocresol purple. GA is reported as percent of albumin concentration. The interassay coefficient of variation for GA was 1.6% at 15.6% and 1.8% at 35.2%.

Determination of Diagnostic Thresholds for HbA_{1c}, GA, and Fructosamine

For detecting Abnl-GT by HbA_{1c}, the standard threshold of 39 mmol/mol (5.7%) was used (1). To identify a diagnostic threshold for GA and fructosamine, the procedure established by the Atherosclerosis Risk in Communities (ARIC) investigators was followed (20). As the diagnostic threshold for detecting Abnl-GT using HbA_{1c} was 39 mmol/mol (5.7%) and corresponded to the upper quartile of our population distribution, we chose the upper quartile for GA (14.2%) and fructosamine (234 μ mol/L).

Statistical Analyses

Unless otherwise stated, data are presented as mean \pm SD. Analyses included one-way ANOVA with Bonferroni corrections for multiple comparisons, χ^2 tests, unpaired t tests, McNemar test for matched pairs, Net Reclassification Improvement (NRI), and κ -statistic for diagnostic reproducibility. κ -Statistic categories were: slight (0–0.20), fair (0.21–0.40), moderate (0.41–0.60), substantial (0.61–0.80), and excellent (0.81–1.0) (21). P values \leq 0.05 were considered significant. Analyses were performed with Stata 16.

RESULTS

The African regions of origin of the participants were: West 53% (220 out of 416), Central 18% (74 out of 416), and East 29% (12 out of 416). Three participants from Southern African countries were included in the Central African group. Characteristics by African region of origin are provided in Supplementary Table 1. Of note, there was no difference by African region in sex distribution, age, or age at immigration. West Africans had resided in the U.S. the longest, 14 ± 11 years ($P = 0.009$). Hemoglobin did not differ by African region. The prevalence of heterozygous hemoglobinopathies (i.e., sickle cell trait and hemoglobin C trait) were

highest in West and Central Africa (both $P = 0.002$).

There was no difference by African region in body size, glucose tolerance, insulin resistance, or insulin secretion (Supplementary Table 1). As these parameters did not vary significantly by African region of origin, participants from West, Central, and East Africa were combined into a single group and evaluated by BMI category and glucose tolerance status (Table 1).

Metabolic Characteristics by BMI Category

Seventy-five percent were nonobese (303 out of 416) (BMI <30 kg/m²) and 25% obese (113 out of 416) (BMI ≥ 30 kg/m²). The nonobese were younger than the obese participants (38 ± 10 vs. 41 ± 10 ; $P = 0.001$). In addition, the nonobese had lower BMI, lower WC, and less VAT (Table 1). Raw data are presented in Table 1; even after adjustment for age, the significant differences between the nonobese and obese remained.

Additionally, FPG, 2-h glucose, and AUC-glucose were lower in the nonobese group. Similarly, the nonobese were less insulin resistant, had lower insulin secretion, and a higher Oral Disposition Index.

However, the prevalence of Abnl-GT (34% vs. 42%; $P = 0.124$) was similar in the nonobese and obese, respectively; and the absolute number of nonobese Africans with Abnl-GT ($n = 104$) was nearly double the number of obese with Abnl-GT ($n = 48$). Among the nonobese who had Abnl-GT, 30% (31 out of 104) had a BMI <25 kg/m², and 70% (73 out of 104) had a BMI ≥ 25.0 kg/m² and <30 kg/m².

Reproducibility of Nonfasting Markers of Glycemia

Reproducibility by κ -statistic for HbA_{1c}, GA, and fructosamine was 0.85, 0.83, and 0.60, respectively. This degree of reproducibility is excellent for HbA_{1c} and GA, but only moderate for fructosamine (21). Therefore, only results for HbA_{1c} and GA are presented.

HbA_{1c} and GA Individually and Combined

The term GA-alone represents the added value provided by GA. Some case subjects with Abnl-GT were detected by HbA_{1c} and not GA (Fig. 1, in red). Some case subjects were detected by GA and not

Table 1—Participant characteristics according to BMI category¹

Parameter ²	Total (n = 416)	Nonobese (n = 303) (73%)	Obese (n = 113) (27%)	P value ³
Age (years)	39 ± 10	38 ± 10	41 ± 10	0.001
BMI (kg/m ²)	27.7 ± 4.5	25.5 ± 2.7	33.5 ± 2.9	<0.001
BMI if Abnl-GT present (kg/m ²)	28.8 ± 4.5	26.3 ± 2.3	34.1 ± 3.1	<0.001
WC (cm) (n = 415)	91 ± 12	86 ± 9	103 ± 9	<0.001
VAT (cm ³) (n = 406)	99 ± 69	83 ± 61	143 ± 71	<0.001
Fasting glucose (mmol/L)	5.1 ± 0.8	5.1 ± 0.5	5.4 ± 1.2	<0.001
Glucose at 2 h (mmol/L)	7.3 ± 2.4	7.1 ± 2.1	7.9 ± 2.9	0.001
AUC-glucose (n = 414)	544 ± 128	533 ± 109	572 ± 166	0.007
ISI (n = 412)	5.43 ± 3.64	6.01 ± 3.73	3.89 ± 2.87	<0.001
Insulin secretion (n = 412)	0.54 ± 0.36	0.49 ± 0.30	0.67 ± 0.45	<0.001
Oral Disposition Index (n = 412)	2.25 ± 0.99	2.34 ± 0.98	2.01 ± 0.97	0.002
Abnl-GT (%)	37 (152/416)	34 (104/303)	42 (48/113)	0.124
Diabetes (%)	7 (28/416)	5 (16/303)	11 (12/113)	0.053
Prediabetes (%) ⁴	32 (124/388)	31 (88/287)	36 (36/101)	0.356

¹Nonobese: BMI <30.0 kg/m²; obese: BMI ≥30.0 kg/m². ²Data are mean ± SD or percentages. ³Comparisons were by unpaired *t* tests for continuous variables and χ^2 for categorical variables. ⁴Denominators are all individuals without diabetes (normal glucose tolerance and prediabetes).

HbA_{1c} (Fig. 1, in blue). Some were identified by both tests (Fig. 1, in purple). GA-alone refers to the individuals with Abnl-GT detected by GA and not HbA_{1c} (Fig. 1, in blue).

Sensitivities and Specificities

Nonobese

Sensitivity for the diagnosis of Abnl-GT by HbA_{1c} and GA was similar, 36% and 37%, respectively (*P* = 0.999). The sensitivity of the combined tests was 58%, which was significantly greater than HbA_{1c} alone (*P* < 0.001) (Fig. 2A and Supplementary Table 2). Sensitivity was higher for the combined tests because GA-alone identified 37% (22 out of 60) of the nonobese not detected by HbA_{1c}

(Fig. 1). Specificities for the diagnosis of Abnl-GT singly by HbA_{1c} and by GA and then both combined were 80%, 75%, and 60%, respectively (Supplementary Table 2). The NRI for HbA_{1c} plus GA versus HbA_{1c} alone was 0.238 (95% CI 0.018, 0.459; *P* = 0.034).

Obese

For the diagnosis of Abnl-GT, the sensitivity of HbA_{1c} at 60% was greater than the sensitivity of 27% for GA (*P* < 0.001). As GA-alone detected only 10% (3 out of 32) of the obese not detected by HbA_{1c}, combining HbA_{1c} with GA did not improve detection of Abnl-GT (*P* = 0.250) (Fig. 2B and Supplementary Table 2). Specificities for the diagnosis of Abnl-GT by HbA_{1c} and GA singly and combined

were: 80%, 94%, and 74%, respectively (Supplementary Table 2). The NRI for HbA_{1c} versus GA versus HbA_{1c} alone was 0.419 (95% CI 0.141, 0.696; *P* = 0.003).

Total Cohort

Diagnostic sensitivities of HbA_{1c} and GA were 43% versus 34% (*P* = 0.086). At 61%, the sensitivity of the combined tests was significantly greater than when only HbA_{1c} was used (*P* < 0.001) (Fig. 2C and Supplementary Table 2). By identifying 28% (26 out of 92) of Africans with Abnl-GT not detected by HbA_{1c}, GA-alone contributed to the higher sensitivity of the combined tests in the nonobese category. Specificities for the diagnosis of Abnl-GT by HbA_{1c} and GA singly and combined were 80%, 80%, and 63%, respectively (Supplementary Table 2). The NRI for HbA_{1c} and GA versus HbA_{1c} alone was 0.270 (95% CI 0.091, 0.448; *P* = 0.003).

Characteristics of Individuals With Abnl-GT Diagnosed by GA-Alone Versus HbA_{1c}

Participants with Abnl-GT detected by GA-alone were younger than those detected by HbA_{1c} (41 ± 9 vs 47 ± 10 years; *P* = 0.015). BMI, WC, and VAT were also lower (Fig. 3). Differences in body size did not change with age adjustment; therefore, raw data are presented (Fig. 3 and Supplementary Table 3). In contrast to body size measurements, insulin resistance, insulin secretion, and Disposition Index did not differ by diagnostic test (Supplementary Table 3). Similarly, prevalence of heterozygous hemoglobinopathy

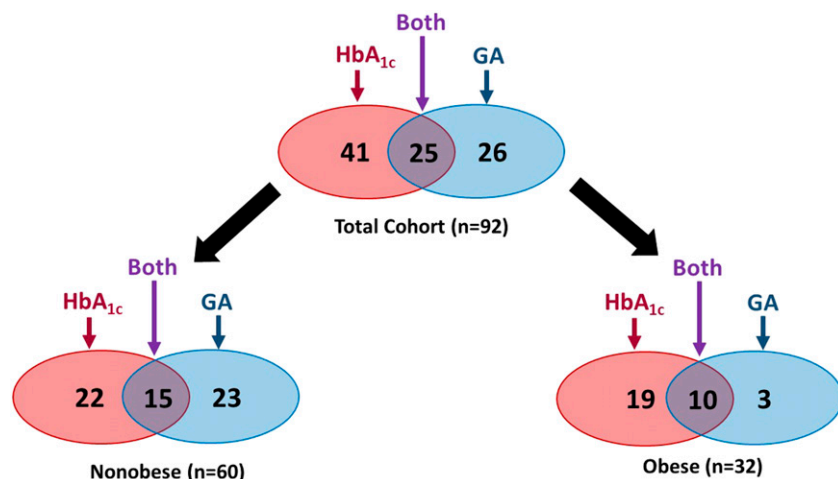


Figure 1—Successful diagnostic test by BMI category in participants with Abnl-GT. The diagnostic color coding: red for HbA_{1c}, blue for GA, and purple for both tests.

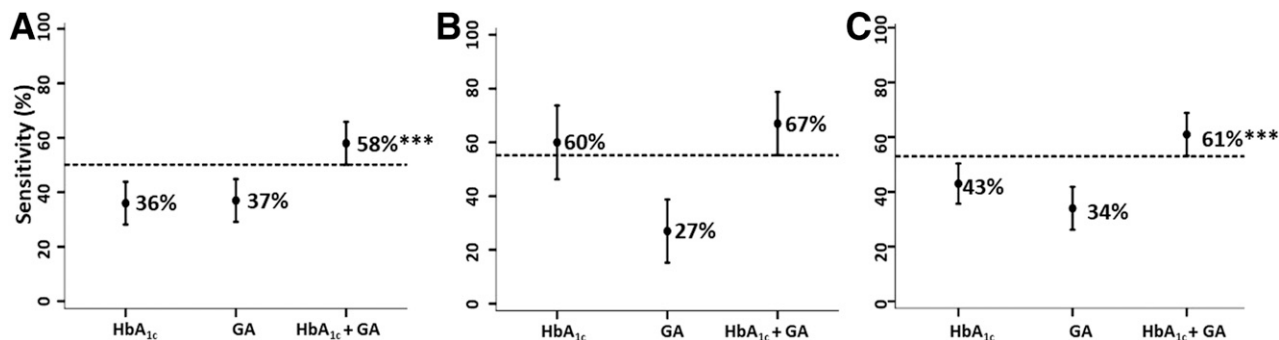


Figure 2—Sensitivities for the diagnosis of Abnl-GT by BMI category for HbA_{1c} and GA singly and combined. *A*: Nonobese: sensitivities for HbA_{1c}, 36%; GA, 37%; and HbA_{1c} plus GA, 58%. Corresponding specificities were 80%, 75%, and 60%, respectively. *B*: Obese: sensitivities for HbA_{1c}, 60%; GA, 27%; and HbA_{1c} plus GA, 67%. Corresponding specificities were 80%, 94%, and 74%, respectively. *C*: Whole cohort: sensitivities for HbA_{1c}, 43%; GA, 34%; and HbA_{1c} plus GA, 61%. Corresponding specificities were 80%, 80%, and 63%, respectively. Data are mean (95% CI). ****P* < 0.001.

(sickle cell trait or hemoglobin C trait) (27% vs. 24%; *P* = 0.789) and albumin (4.04 ± 0.25 vs. 3.98 ± 0.28; *P* = 0.367) did not differ by diagnostic test (Supplementary Table 3).

CONCLUSIONS

This investigation made three findings that could influence the approach to Abnl-GT in Africa. First, there were significant differences by BMI category in the diagnostic capabilities of GA and HbA_{1c} such that the combination leads to improved detection of Abnl-GT in the nonobese. Second, the prevalence of Abnl-GT was similar in nonobese and obese Africans. Third, in contrast to fructosamine, both HbA_{1c} and GA provided highly reproducible results.

Overall, the combination of GA and HbA_{1c} identified more Africans with Abnl-GT than HbA_{1c}. This was because GA increased detection of the nonobese with Abnl-GT by 33% (Fig. 1). As adding GA did not improve the detection of Abnl-GT in the obese, the use of GA could be reserved for the nonobese.

The prevalence of Abnl-GT in African immigrants to the U.S. was 37% and similar in the nonobese and obese. Therefore, clinicians caring for Africans should not identify an individual as “low risk” because they are nonobese (15,22). In short, the threshold for BMI-related risk for Abnl-GT may be lower in Africans than for African Americans, which suggests the need for tailored therapeutic approaches (23). Furthermore, Abnl-GT in the nonobese is observed in India and in many other low- and middle-income countries globally (13,14).

Reproducibility of diagnostic markers is another important finding. With studies done 11 ± 7 days apart and a κ-statistic ≥0.8 for both HbA_{1c} and GA, their reproducibility was excellent. Hence, the biological variability and the interassay variation for both HbA_{1c} and GA are low. In contrast, the fructosamine assay is less optimal and subject to more analytic variables than GA (24). Therefore, it is not surprising that duplicate fructosamine studies revealed only moderate reproducibility (κ-statistic of 0.6). Suboptimal

reproducibility may explain why for Abnl-GT, fructosamine is a poor diagnostic test (25).

Relationship Between GA and BMI

Africans with Abnl-GT detected by GA-alone had lower BMI, WC, and VAT than their counterparts detected by HbA_{1c} (Fig. 3). This is consistent with the observation that GA correlates inversely with BMI, WC, and VAT (12,26). In fact, He et al. (27) found that for every 1 kg/m² increase in BMI, GA decreased by 0.13%. This inverse relationship is often attributed to increased catabolism of albumin from obesity-related chronic inflammation or insulin resistance (28,29). Interestingly, degree of insulin resistance and β-cell function did not differ by diagnostic test (Supplementary Table 3). Therefore, the etiology of Abnl-GT (insulin resistance vs. relative β-cell failure) cannot be ascertained by whether diagnosis was by HbA_{1c} or GA.

BMI Category to Guide Use of GA

In detecting Abnl-GT, the diagnostic utility of GA appears to depend on BMI category. We speculate that for normal-weight people, GA-alone may be sufficient to detect Abnl-GT. In overweight people, GA should be combined with HbA_{1c} while HbA_{1c} may be satisfactory in obese people.

Studies from East Asia reveal that the performance of GA as a single diagnostic test was equivalent to or better than HbA_{1c} (30,31). GA may be an effective diagnostic test for Abnl-GT because the BMI in East Asians with T2D is typically <25 kg/m² (31).

In the Africans in America cohort, two-thirds of the nonobese participants with

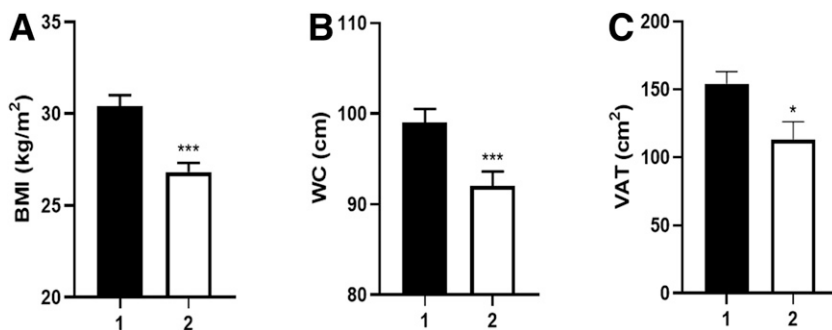


Figure 3—Participant characteristics with Abnl-GT according to diagnostic test: *A*: BMI, *B*: WC, *C*: VAT. In each panel: 1 is HbA_{1c} or HbA_{1c} and GA; 2 is GA-alone. Data presented as mean ± SE. **P* < 0.05; ****P* < 0.001.

Abnl-GT were overweight (BMI 25.0–29.9 kg/m²), and one-third were normal weight (<25.0 kg/m²). Considering that the Africans in America cohort represents mainly the overweight with Abnl-GT, we found that combining HbA_{1c} with GA optimized detection.

For obese people with Abnl-GT, HbA_{1c} alone may be sufficient. In a South African study of mixed-ancestry adults in whom the mean BMI of the group with T2D was 32.5 kg/m², Zemlin et al. (32) reported that the combination of GA and HbA_{1c} was no better than HbA_{1c} alone. Our findings were similar in the obese participants in the Africans in America cohort (Fig. 2B). However, NRI suggests that the combined tests may be beneficial in the obese as well (Supplementary Table 4). While this would be an excellent development, there is concern that the assumptions made in the calculation of NRI may be overly optimistic in predicting improvement (33).

Feasibility of Obtaining GA

The enzymatic method used to measure GA has been evaluated in many clinical studies and is approved for use in the U.S., Japan, Korea, Indonesia, and China (11,30). The assay is reproducible, precise, and easily performed on automated analyzers that can measure glucose or electrolytes. Analyzers of this type are widely available in both clinical and research settings in Africa (M. Nyirenda, personal communication).

Methodology for Determining Diagnostic Thresholds for GA

Thresholds for FPG, 2-h glucose, and HbA_{1c} were determined by their relationship to diabetic retinopathy (34). The methods used to determine diagnostic thresholds for GA rely on glucose concentrations obtained during the OGTT. The two most common approaches are: 1) identifying the ability of GA to predict Abnl-GT by calculating the area under the receiver operating characteristic curve and applying the Youden Index to define the optimal cut point (30–32), and 2) use the GA cutoff at the upper 75th, 95th, or 97.5th percentile of the population being evaluated (20). For example, the ARIC investigators determined the upper fraction of their cohort who had HbA_{1c} ≥5.7%. As 5.7% occurred as the cutoff for the 75th percentile, they used the GA threshold at the 75th percentile

(20). With these two approaches, GA thresholds range from 13 to 16% depending on whether the outcome is prediabetes, T2D, or a combination (i.e., Abnl-GT) (20,30–32). We used the approach taken by the ARIC investigators, and our threshold for Abnl-GT was 14.2%.

To systematically resolve which diagnostic threshold to use for GA, the way forward may be a two-step process with the establishment of an International Working Group followed by an International Consensus Panel. The International Working Group could survey existing studies and determine: 1) end point criteria for GA (retinopathy or glucose or other), 2) the statistical methodology for determining threshold (AUC–receiver operating characteristic or percentile), and 3) size and diversity of the cohorts needed to test recommendations. Within these cohorts, BMI and race/ethnicity would be specifically considered. If the International Working Group decides that current data are insufficient, parameters for study design for future determination by International Consensus Panel could be set.

Strengths and Limitations

Our study is the largest exploring the diagnostic value of nonfasting markers of glycemia in African-born Blacks. In addition, it is the first investigation to explore the diagnostic reproducibility of GA, fructosamine, and HbA_{1c}. Furthermore, studies evaluating the diagnostic efficacy of these markers often state they cannot rule out the effect of confounding factors (6). However, our study provides documentation of the performance of GA, fructosamine, and HbA_{1c} in the absence of cirrhosis, thyroid disease, hemoglobinopathies, anemia, nutritional deficiencies, hypoalbuminemia, and renal failure.

The study used a convenience sample. However, for three reasons, the Africans in America cohort appears to be representative of African-born Blacks living in the U.S. First, as most participants were from West African countries, the sample size of 416 was large enough to reflect known immigration patterns (35). Second, compared with East Africa, the prevalence of sickle cell trait and hemoglobin C trait was higher in West and Central Africans. Therefore, the sample size was large enough to detect known genetic differences by African region of

origin (36). Third, the prevalence of T2D was 7%, which was comparable to the 8% prevalence of T2D in African-born Blacks living in Canada (37). Similar data are not available in the U.S.

The potential limitations of our investigation are intrinsic to our study design. One limitation is the applicability of our results to populations in African countries. However, we designed the study to use resources available in America to provide a proof of concept about the potential value of GA in Africans. Therefore, our investigation provides justification: 1) for the funding of studies to be conducted in Africa, and 2) for the study to be used for power analyses for population-based prospective studies that have optimization of GA and HbA_{1c} thresholds built into their study design. To assist in these analyses, we provided the total number of participants with Abnl-GT and then for the two components of Abnl-GT, diabetes and prediabetes (Table 1). The current study was not large enough to examine diabetes and prediabetes separately.

Another challenge for African countries is that in the absence of point-of-care options for GA, initial studies will have to be conducted in urban or semiurban areas where there is access to clinics that have both automated analyzers and the opportunity to detect confounders such as nutritional deficiencies, infections, and hemoglobinopathies.

In addition, we focused on sensitivity over specificity. Results for specificity for each testing paradigm are provided only in the legend to Fig. 2 and Supplementary Table 2. However, the sensitivity of HbA_{1c} as a single test in the nonobese is <40%. Therefore, HbA_{1c} is not a viable option for effective screening for a disease that is reaching epidemic proportions in Africa. After improved detection is achieved, future studies can determine if the benefits of more optimal detection offset the lower specificity.

Conclusion

Between 2019 and 2045, a steep increase in Abnl-GT prevalence in Africa, especially in the nonobese, is anticipated. Innovative and improved diagnostic tools may be the best way to limit the epidemic, improve data collection, and inform treatment paradigms. GA is a nonfasting, easily obtainable, highly reproducible test that, in combination with HbA_{1c}, provides valuable

diagnostic information about Abnl-GT in nonobese Africans living in the U.S. Data from the Africans in America cohort provide a proof of concept and can serve as primary data for sample size calculations for population-based prospective studies in Africa. Research on the diagnostic value of GA may lead to better screening, earlier interventions, and ultimately less medical and social consequences from Abnl-GT.

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