Comparison of Telomere Length in Black and White Teachers From South Africa: The Sympathetic Activity and Ambulatory Blood Pressure in Africans Study

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Objective: Telomere length is a marker of biological aging that has been linked to cardiovascular disease risk. The black South African population is witnessing a tremendous increase in the prevalence of cardiovascular disease, part of which might be explained through urbanization. We compared telomere length between black South Africans and white South Africans and examined which biological and psychosocial variables played a role in ethnic difference in telomere length. Methods: We measured leukocyte telomere length in 161 black South African teachers and 180 white South African teachers aged 23 to 66 years without a history of atherothrombotic vascular disease. Age, sex, years having lived in the area, human immunodeficiency virus (HIV) infection, hypertension, body mass index, dyslipidemia, hemoglobin A1c, C-reactive protein, smoking, physical activity, alcohol abuse, depressive symptoms, psychological distress, and work stress were considered as covariates. Results: Black participants had shorter (median, interquartile range) relative telomere length (0.79, 0.70–0.95) than did white participants (1.06, 0.87–1.21; p < .001), and this difference changed very little after adjusting for covariates. In fully adjusted models, age (p < .001), male sex (p = .011), and HIV positive status (p = .023) were associated with shorter telomere length. Ethnicity did not significantly interact with any covariates in determining telomere length, including psychosocial characteristics. Conclusions: Black South Africans showed markedly shorter telomeres than did white South African counterparts. Age, male sex, and HIV status were associated with shorter telomere length. No interactions between ethnicity and biomedical or psychosocial factors were found. Ethnic difference in telomere length might primarily be explained by genetic factors. Key words: cardiovascular disease, ethnicity, genetics, psychological stress, telomere.

Telomeres are deoxyribonucleic acid (DNA)–protein complexes at the ends of eukaryotic chromosomes providing protection against genome instability—promoting events with potentially catastrophic health outcomes (1). During somatic cell division, the ribonucleoprotein enzyme telomerase compensates for terminal telomeric DNA losses by lengthening the ends of telomeres through RNA-templated addition of telomeric sequences (2). Repair processes do not fully balance telomere attrition, so telomeres grow shorter with repeated cell divisions.

Mean telomere length of replicating somatic cells is an indicator of biological age and is modified by many factors including genetic, socioeconomic, cardiometabolic, life-style, and psychosocial ones (2,3). Specifically, shorter telomere length has been linked to male sex (4), low socioeconomic status (SES) (5), and poorer health habits like smoking (5), low physical activity (6), and high alcohol consumption (7). In terms of psychosocial measures, shorter telomeres have been linked to depressive symptoms (8), psychological distress (9), and work-related stress (10). Oxidative stress and low-grade inflammation may contribute to telomere attrition in multiple diseases including atherosclerotic vascular diseases (11) like coronary heart disease (12) and ischemic stroke (13). Accordingly, the metabolic cardiovascular risk factors hypertension (14), obesity (15), dyslipidemia (16), and elevated blood glucose (16) have been linked to shortened telomeres.

Telomere length has not previously been investigated in the black population of South Africa. Moreover, studies on telomere length in African Americans are sparse, whereby the paradigm that links telomere length with CVD risk is based mainly on data from non-African populations (24). In population-based studies, African Americans showed longer (5,24–27) and shorter (28) age- and sex-adjusted telomere length compared with white Americans. However, African Americans tend to show ancestry primarily from Western Africa with a genetic makeup that is quite different from black South Africans (29).

INTRODUCTION

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Telomere length has not previously been investigated in the black population of South Africa. Moreover, studies on telomere length in African Americans are sparse, whereby the paradigm that links telomere length with CVD risk is based mainly on data from non-African populations (24). In population-based studies, African Americans showed longer (5,24–27) and shorter (28) age- and sex-adjusted telomere length compared with white Americans. However, African Americans tend to show ancestry primarily from Western Africa with a genetic makeup that is quite different from black South Africans (29).
Therefore, the primary aim of this study was to compare leukocyte telomere length between black South African teachers and white South African teachers with an equal sex distribution and socioeconomic background and free of clinically overt atherothrombotic vascular disease. To the extent that there would be an ethnic difference in telomere length, the secondary aim of our study was to explore whether sociodemographic factors, HIV infection, and cardiometabolic, lifestyle, and psychosocial characteristics (2–11,14–17,21) would explain an ethnic difference in telomere length.

**MATERIALS AND METHODS**

**Study Participants**

The participants of this study were recruited as part of the Sympathetic Activity and Ambulatory Blood Pressure in Africans (SABPA) study conducted between February 2008 and May 2009 (SABPA I). Data presented here were collected between February 2011 and May 2012 as part of the follow-up investigation (SABPA II) when leukocyte telomere length was measured. The study protocol applies with the Declaration of Helsinki (30) and was approved by the ethics review board of the North-West University, Potchefstroom Campus (00036/07/S6). All participants provided written informed consent before participation. The SABPA I study had a target population comparative design and recruited 341 black and white South African teachers aged 25 to 65 years, working in the Dr Kenneth Kaunda Education district in the North West Province, South Africa. This selection assured a homogenous sample from a similar socioeconomic class. Exclusion criteria for SABPA I included pregnancy, lactation, and vaccination or blood donation within 3 months before participation (22).

There were 359 teachers participating in SABPA II study (attrition rate, 12.2%), whereby a similar proportion of black and white South Africans (p = .44) and of men and women (p = .42) were lost to follow-up.

For the purpose of the present study, we excluded three participants with a history of atherothrombotic vascular disease (myocardial infarction, stroke, and a history of atherothrombotic vascular disease (myocardial infarction, stroke, hemoglobin A1c (HbA1c; a marker of dyslipidemia). HbA1c was determined by a turbidometric inhibition immunoassay method from EDTA plasma (Integra 400; Roche, Basel, Switzerland). Ultra high-sensitivity CRP was measured in serum with a turbidimetric method (Unicel DXC 800; Beckman and Coulter, Germany).

**Human Immunodeficiency Virus**

HIV infection was determined with a rapid antibody test in plasma (First response kit; PMC Medical, Daman, India) and confirmed with the Pareczhak test (BHAT Bio-Tech, Bangalore, India).

**Health Behaviors**

Participants who indicated that they currently smoked at least one cigarette per day were categorized as current smokers and all others as noncurrent smokers. Serum levels of γ-GT activity were measured with an enzymatic colorimetric assay (Cobas Integra 400 plus) and used as a marker of alcohol abuse with cutoffs of at least 55 U/L for men and at least 38 U/L for women (33). For the assessment of physical activity, participants wore the Actiheart physical activity monitor (CamNtech Ltd, Upper Pendrill Court, Papworth Everard, Cambridgeshire, CB233UY, UK) apparatus for 7 days continuously. Total energy expenditure in kilocalories per day was computed, taking resting metabolic rate into account.

**Psychosocial Measures**

We used the nine-item Patient Health Questionnaire to measure the frequency of depressive symptoms during the prior 2 weeks (34). Each item is rated on a 4-point scale (0 = “not at all,” 1 = “several days,” 2 = “more than half the days,” 3 = “nearly every day”) giving a possible global score of 0 to 27. We additionally tested for a threshold effect of depressive symptom severity applying a cutoff of at least 10 defining “moderate depression”; this cutoff was previously shown to have a sensitivity of 88% and a specificity of 88% for major depression assessed by a structured interview (34).

With the 28-item General Health Questionnaire, we quantified perceived psychological distress in general related to symptoms of depression, anxiety, somatic complaints, and social withdrawal over the past few weeks (35). Each item is rated on a 4-point scale (1 = “not at all,” 2 = “no more than usual,” 3 = “rather more than usual,” 4 = “much more than usual”); typical items are “Felt that you are ill?” and “Been getting scared or panicky for no good reason?” We applied the binary scoring method with the two least symptomatic answers scoring “0” and the two most symptomatic answers scoring “1.” The global score provides a severity measure of total psychological distress (35).

The 20-item Teacher Stress Inventory was applied to measure work stress (36). Participants are requested to answer the question: “As a teacher, how great a source of stress are these factors to you?” Participants are then presented a list of potential work-related stressors (e.g., “difficult class,” “pressure from principal and educational officials”). Each item is rated on a 5-point scale ranging from 1 (“no stress”) to 5 (“extreme stress”), yielding a global score of 20 to 100.

**Measurement of Leukocyte Telomere Length**

Leukocyte genomic DNA was extracted using a NucleoSpin 96 Blood Core kit (Machery Nagel, Düren, Germany) according to the manufacturer’s instructions.
and stored at −20°C. All samples were diluted to a concentration of 10 ng/μl before the analyses. To prepare the reference DNA samples, all isolated DNAs were mixed together in equal proportions representing the average of all analyzed patients. We assessed telomere length by multiplex quantitative real-time polymerase chain reaction (Q-PCR) as previously described (37). Five concentrations of a reference DNA sample spanning a 75-fold range of DNA concentrations were prepared by serial dilution and analyzed in triplicate; these reactions provided the data for the generation of the standard curves used for relative quantification. All experimental DNA samples were assayed in triplicate with an average coefficient of variation of 16.4%. The intrabatch and interbatch control had a coefficient of variation of 4.3% and 4.8%, respectively. All primers were manufactured by Integrated DNA Technologies (Coralville, IA).

The thermal cycling profile was as follows: Stage 1, 15 minutes at 95°C; Stage 2, 2 cycles of 15 seconds at 94°C and 15 seconds at 49°C; and Stage 3, 32 cycles of 15 seconds at 94°C, 10 seconds at 62°C, 15 seconds at 73°C with signal acquisition, 10 seconds at 84°C, and 15 seconds at 87°C with signal acquisition. The 73°C reads provided the Ct values for the amplification of the telomere template (in early cycles where the hbg signal is still at baseline); the 87°C reads the Ct values for the amplification of the hbg template (at this temperature, there is no signal from the telomere PCR product because it is fully melted). After thermal cycling and raw data collection were complete, Ct values generated by CFX Manager Software Version 1.6 (Bio-Rad) were exported to REST software (Relative Expression Software Tool), version REST-384. PCR efficiency for each primer set was calculated by serial dilutions of reference DNA using the REST software tool. The relative telomere length per cell was also calculated using the REST software as the ratio (T/S) between relative content of telomere PCR product (T) and hbg PCR product (S). The T/S ratio was used the measure of leukocyte telomere length in all analysis presented here and was calculated for each individual participant relative to the average telomere length of the total analyzed group (344 participants).

All PCRs were performed with CFX96 Touch Real-Time PCR Detection System (Bio-Rad, Hercules, CA) in a 25-μl volume in the biochemistry laboratory, North-West University (accredited by the Health Professional Council of South Africa). PCR reactions comprised 20 μl master mix (including 5 μl of 5× HOT FIREPol EvaGreen qPCR Mix Plus (no ROX; Solis BioDyne, Tartu, Estonia) and 5 μl of each experimental DNA sample. For multiplex Q-PCR, the telomere primer pair telg: ACACCTAAGGTGGTGGTGGTTGGGTTGGITGTAAGTGT and telc: TTGTAAAGTACCATCCCTATCCCTATCCCTATCCCTATCC (final concentrations 900 nM each) was combined with primers for the single copy gene human β-globin (hbg); hbg: CGCGCGCGGGCAGCGGGCCGGCTGGCGGCGGTCCACACGTTACCTCCCTG and hbgd: GCCGCCGCCGGCGCCGGGCCGTCCCGCGGAGGAGAAGTCTGCCGTT (final concentrations 500 nM each), to give PCR products of 79 and 106 base pairs, respectively. All primers were manufactured by Integrated DNA Technologies (Coralville, IA).

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**Statistical Analysis**

Data were analyzed using SPSS version 21.0 statistical software for Windows (SPSS Inc, Chicago, IL). The level of significance was set at p < .05 (two tailed). To obtain more normally distributed values, we log10 transformed values for telomere length, age, years having stayed in the area, systolic BP, diastolic BP, T-C, HDL-C, BMI, HbA1c, CRP, γ-GT, total energy expenditure, depressive symptoms, and symptoms of psychological distress and work stress before the analyses. We applied independent t test and Pearson χ² test for group comparisons on continuous and categorical data, respectively. Pearson correlation and partial correlation analyses (controlling for age and sex) were used to estimate the bivariate association between telomere length and covariates. We used univariate analysis of covariance to compute the relationship between ethnic identity and telomere length. We only covaried for those variables that were significantly associated with telomere length in the bivariate correlation analysis, after controlling for age and sex as standard confounders of telomere length. Covariates were selected a priori based on previously described and/or hypothesized relationships with telomere length (2–11,14–17,21): a) sociodemographic factors (age, sex, number of years having lived in the area) and HIV infection status, b) metabolic factors (hypertension, BMI, T-C/HDL-C ratio, HbA1c, CRP),

### TABLE 1. Participant Characteristics.

<table>
<thead>
<tr>
<th>Variables</th>
<th>Black South Africans (n = 161)</th>
<th>White South Africans (n = 180)</th>
<th>p</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age, y</td>
<td>47.0 (42.0–54.0)</td>
<td>50.0 (44.0–58.0)</td>
<td>.269</td>
</tr>
<tr>
<td>Female sex, n (%)</td>
<td>76 (47.2)</td>
<td>92 (51.1)</td>
<td>.471</td>
</tr>
<tr>
<td>Having lived in the area, y</td>
<td>38.0 (21.5–48.0)</td>
<td>23.0 (13.0–32.0)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>HIV infection, n (%)</td>
<td>21 (13.0)</td>
<td>0 (0)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Systolic blood pressure, mm Hg</td>
<td>133.0 (123.5–145.0)</td>
<td>122.0 (116–130)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Diastolic blood pressure, mm Hg</td>
<td>83.0 (75.5–91.0)</td>
<td>75.0 (69.0–80.0)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Ambulatory hypertension, n (%)</td>
<td>109 (67.7)</td>
<td>63 (35.0)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Body mass index, kg/m²</td>
<td>30.1 (25.7–34.8)</td>
<td>27.9 (24.4–30.7)</td>
<td>.005</td>
</tr>
<tr>
<td>Total cholesterol, mmol/l</td>
<td>4.41 (3.81–5.27)</td>
<td>4.24 (3.68–4.88)</td>
<td>.008</td>
</tr>
<tr>
<td>HDL cholesterol, mmol/l</td>
<td>0.96 (0.79–1.13)</td>
<td>0.97 (0.80–1.23)</td>
<td>.144</td>
</tr>
<tr>
<td>Total cholesterol/HDL cholesterol</td>
<td>4.68 (3.83–5.90)</td>
<td>4.31 (3.23–5.32)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Hemoglobin A1c, %</td>
<td>5.84 (5.59–6.13)</td>
<td>5.48 (5.29–5.64)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>C-reactive protein, mg/L</td>
<td>4.63 (1.83–8.51)</td>
<td>1.19 (0.58–2.36)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Current smoker, n (%)</td>
<td>23 (14.3)</td>
<td>25 (13.9)</td>
<td>.466</td>
</tr>
<tr>
<td>γ-Glutamyltransferase, U/L</td>
<td>38.4 (22.9–69.8)</td>
<td>18.2 (11.9–31.6)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Alcohol abuse, n (%)</td>
<td>61 (37.9%)</td>
<td>16 (8.9%)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Total energy expenditure, kcal/d</td>
<td>2896 (2416–3639)</td>
<td>3045 (2480–3892)</td>
<td>.284</td>
</tr>
<tr>
<td>Depressive symptoms (score)</td>
<td>7.88 (5.00–11.00)</td>
<td>5.00 (3.00–8.00)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Moderate depression, n (%)</td>
<td>58 (36.0)</td>
<td>34 (18.9)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Distress symptoms (score)</td>
<td>5.00 (1.00–10.0)</td>
<td>2.00 (0.00–7.00)</td>
<td>&lt;.001</td>
</tr>
<tr>
<td>Work stress (score)</td>
<td>71.0 (58.0–82.0)</td>
<td>63.0 (49.0–73.0)</td>
<td>&lt;.001</td>
</tr>
</tbody>
</table>

HIV = human immunodeficiency virus; HDL = high-density lipoprotein.

Data are given as number of observations (n) and median of total (%) or median with interquartile range (25th–75th percentile).
c) life-style factors (current smoking, total energy expenditure, alcohol abuse), and d) psychosocial factors (depressive symptoms, moderate depression, psychological distress, work stress).

We further undertook a moderator analysis for all covariates that were significantly different between black and white participants and thus might differently affect telomere length depending on ethnicity. For this purpose, we probed for significant two-way interactions between ethnicity and each such covariate controlling for age and sex, and in the case of psychosocial characteristics, additionally for other significant covariates. Effect sizes are expressed as partial $\eta^2$.

**RESULTS**

**Study Participants**

Table 1 summarizes the characteristics of the 341 study participants per ethnic group. Compared with white participants, black participants had been living longer in the Potchefstroom area. All of the HIV-positive individuals were black. As compared with their white counterparts, black participants had a worse metabolic profile showing more ambulatory hypertension; higher levels of BMI, HbA1c, and CRP; and higher T-C/HDL-C ratio. Regarding life-style factors, there was evidence for significantly more alcohol abuse in black participants than in white participants. Psychosocial stress in the form of depressive symptoms, distress, and perceived work stress was significantly higher in black participants relative to white participants.

**Associations Between Telomere Length and Covariates**

**Sociodemographic Factors**

Telomeres (median, interquartile range) were shorter in men compared with women (0.89, 0.74–1.04 versus 0.94, 0.76–1.17; $p = .030$) and in relation to greater age ($r = -0.16$, $p = .003$). All of the subsequent bivariate correlation analyses were adjusted for age and sex as standard confounders of telomere length. There emerged significant correlations between shorter telomeres and longer residency in the area ($r = -0.16$, $p = .004$) and HIV infection ($r = -0.23$, $p < .001$).

**Cardiometabolic Factors**

Shorter telomeres were associated with higher systolic ($r = -0.12$, $p = .029$) and diastolic ($r = -0.13$, $p = .016$) BP as well as with hypertension status ($r = -0.13$, $p = .014$). Shorter telomeres were also found in those with higher BMI ($r = -0.13$, $p = .018$) and CRP ($r = -0.26$, $p < .001$) levels. The relationships between telomere length and HbA1c ($r = -0.10$, $p = .074$) and the T-C/HDL-C ratio was not significant ($r = -0.04$, $p = .46$).

**Life-Style Factors**

Shorter telomeres correlated with higher $\gamma$-GT levels ($r = -0.34$, $p < .001$) and also with alcohol abuse ($r = -0.22$, $p < .001$). In contrast, telomere length did not relate to smoking status ($r = 0.04$, $p = .48$) and total energy expenditure ($r = 0.02$, $p = .69$).

**Psychosocial Factors**

All of the relationships of telomere length with measures of psychosocial stress pointed in the expected inverse direction, but failed to reach statistical significance: depressive symptoms ($r = -0.11$, $p = .051$), moderate depression ($r = -0.06$, $p = .30$), symptoms of distress ($r = -0.08$, $p = .13$), and perceived work stress ($r = -0.07$, $p = .21$).

**Association Between Ethnicity and Telomere Length**

The median (interquartile range) of the relative telomere length (T/S ratio) was 0.91 (0.75–1.09) in the total sample, whereby black South Africans had shorter telomeres than white South Africans (0.79, 0.70–0.95 versus 1.06, 0.87–1.21; $p < .001$). Figure 1 shows the average relative telomere length in original units for black participants in comparison with white participants with adjustment for those covariates, which were significantly associated with telomere length in the above correlation analysis (i.e., age, sex, years having lived in the area, HIV infection, hypertension, BMI, CRP, and alcohol abuse). Across five models with subsequent adjustment for sets of covariates, ethnicity was significant for all group comparisons ($p < .001$), with black participants consistently showing shorter average relative telomere length than their white counterparts. Moreover, regardless of any covariate adjustments, the absolute difference in telomere length changed very little between black and white participants.

Table 2 shows the final model for the relation of ethnicity with the log-transformed telomere length with adjustment for those covariates, which were significantly related to telomere length in the above correlation analyses (i.e., age, sex, years having lived in the area, HIV infection, hypertension, BMI, CRP, and alcohol abuse). In addition to shorter telomeres in black...
TABLE 2. Univariate Analysis of Covariance for Leukocyte Telomere Length  

<table>
<thead>
<tr>
<th>Entered Variables</th>
<th>Unstandardized Coefficient</th>
<th>Standard Error</th>
<th>p</th>
<th>Partial $\eta^2$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Intercept</td>
<td>0.582</td>
<td>0.146</td>
<td>&lt;.001</td>
<td>0.046</td>
</tr>
<tr>
<td>Black South African ethnicity</td>
<td>−0.098</td>
<td>0.014</td>
<td>&lt;.001</td>
<td>0.131</td>
</tr>
<tr>
<td>Age</td>
<td>−0.280</td>
<td>0.065</td>
<td>&lt;.001</td>
<td>0.053</td>
</tr>
<tr>
<td>Female sex</td>
<td>0.030</td>
<td>0.012</td>
<td>.011</td>
<td>0.020</td>
</tr>
<tr>
<td>Years having lived in the area</td>
<td>0.006</td>
<td>0.020</td>
<td>.77</td>
<td>0.000</td>
</tr>
<tr>
<td>Human immunodeficiency virus infection</td>
<td>−0.054</td>
<td>0.024</td>
<td>.023</td>
<td>0.016</td>
</tr>
<tr>
<td>Ambulatory hypertension</td>
<td>0.016</td>
<td>0.013</td>
<td>.20</td>
<td>0.005</td>
</tr>
<tr>
<td>Body mass index</td>
<td>−0.085</td>
<td>0.068</td>
<td>.21</td>
<td>0.005</td>
</tr>
<tr>
<td>C-reactive protein</td>
<td>−0.006</td>
<td>0.012</td>
<td>.59</td>
<td>0.001</td>
</tr>
<tr>
<td>Alcohol abuse</td>
<td>−0.016</td>
<td>0.014</td>
<td>.25</td>
<td>0.004</td>
</tr>
</tbody>
</table>

All continuously scaled data were entered as log10-transformed values. Effect sizes are expressed as partial $\eta^2$.

participants than white participants (corresponding to model 5 in Fig. 1), significantly shorter telomeres were also found in those with advanced age, in men compared with women, and in HIV-positive versus HIV-negative participants.

**Moderator Analysis**

Most of the covariates of telomere length, including psychosocial characteristics, were significantly different between black and white participants (Table 1) and thus might differ in their association with telomere length depending on ethnicity. However, adjusting for age and sex, ethnicity did not significantly interact with any covariate to determine telomere length: years having lived in the area ($p = .63$), hypertension ($p = 1.00$), BMI ($p = .40$), T-C/HDL-C ratio ($p = .46$), HbA1c ($p = .17$), CRP ($p = .072$), and alcohol abuse ($p = .59$).

In terms of psychosocial characteristics, there were also no significant interactions of ethnicity with depressive symptoms ($p = .48$), moderate depression ($p = .23$), psychological distress ($p = .21$), and work stress ($p = .12$) in determining telomere length, controlling for age and sex. Moreover, these results did not change with additional adjustment for years having lived in the area, HIV infection, hypertension, BMI, CRP, and alcohol abuse (all $p$ values $\geq .11$ for interactions between ethnicity and psychosocial characteristics).

**DISCUSSION**

The main finding from this study is that black South African teachers had significantly shorter telomeres compared with their white counterparts, even after controlling for a range of socio-demographic, HIV-infection–related, cardiometabolic, life-style, and psychosocial factors which have previously been identified as important correlates of telomere length; this suggests that the observation of shortened telomeres in black South Africans versus white South Africans is a robust one. Among the significant correlates of telomere length, ethnicity showed the greatest effect size for an association with telomere length in absolute terms, explaining 13.1% of the variance in the fully adjusted model; to compare, the well-known correlates age and sex explained substantially less of the variance in telomere length.

We found that sociodemographic, HIV-infection–related, cardiometabolic, life-style, and psychosocial factors which significantly differentiated black and white participants did not emerge as significant moderators of the relation between ethnicity and telomere length after adjustment for all other covariates. This might indicate that genetic factors more than those investigated in our study contributed to shorter telomeres in black South Africans than white South Africans (2,3). Previous studies reported heritability estimates ranging from 34% to 82% (3); for instance, there might be gene polymorphisms that contributed toward telomere length (38), but we did not explore this. However, it should be emphasized that demographic and health-related variables are not irrelevant for telomere length maintenance, but rather of equal importance in black and white South Africans.

We confirmed several previously reported independent associations of telomere length with age, sex, and HIV infection. Women had longer telomeres than did men, whereas older and HIV-positive individuals had shorter telomeres than did younger participants and those HIV negative (4,21). Although showing the expected inverse relationship, we did not find a significant association between psychosocial factors and telomere length, namely, depression, psychological distress, and work stress; this is contrary to some previous studies (8–10), but similar to others (39,40). Psychosocial risk factors of CVD cover different domains of social adversities, personality traits, and negative affect (41). Further studies are needed to answer the question whether cellular aging as reflected by telomere length is a pathway that critically links psychosocial stress to CVD risk and the molecular mechanisms underlying these relationships in South Africans. The underlying mechanism might be complex and, as was revealed for various physical stressors disrupting telomere homeostasis (42), will likely differ in their effects on telomere attrition. For instance, a recent study found shorter telomeres but higher telomerase activity in high-hostile versus low-hostile men, whereby the heightened telomerase activity was interpreted as a compensatory response in men with high levels of cynical hostility (43).

To our knowledge, this is the first study to examine telomere length in black South Africans compared with white South Africans. Interestingly, except from one study also showing shorter telomeres in African Americans compared with whites and also Hispanics (28), the available literature suggests that African Americans have longer (age- and sex-adjusted) telomeres than do white (5,24–27). Similarly, African American adolescents aged 14 to 18 years also had longer telomeres than did their white counterparts (44). Altogether, these population-based studies cover the whole age distribution from adolescence to old age, suggesting that telomeres are consistently longer in African Americans compared with whites up to the age of 80 years (24).

We mention two explanations that might apply to the discrepancy with our study showing that black South Africans had markedly shorter telomeres than white South Africans. First, South Africans have quite a different genetic makeup compared with African Americans. It has been revealed that most African
Americans have mixed ancestry from different regions of western Africa, whereas black South Africans show the most private alleles within Africa (29). Second, although the aforementioned studies, including ours, were all cross-sectional, one of the few longitudinal studies on telomere length found that African Americans had longer telomeres than did whites, but that telomere shortening occurred relatively faster in African Americans over a follow-up of 6 to 10 years (45). Moreover, metabolic factors (46), but also depressive symptoms (8), result in more rapid telomere length attrition, and our black South African participants were clearly worse off in these variables than their white South African counterparts. Our cross-sectional analysis could only capture a snapshot of telomere length across the life span. Theoretically, our black participants could have begun life with longer telomeres (similar to African Americans) than their white counterparts, but experienced exaggerated telomere length attrition due to their progressively increasing CVD risk factors. This idea concurs with the notion that urbanization is a key stress factor in black South Africans in terms of both physical and psychological health sequelae (20), and indeed, an increased cardio-metabolic risk was demonstrated in South Africans attempting to cope with an urban life-style compared with their rural counterparts (47). In addition, this hypothesis is consistent with recent evidence indicating a differential role of telomere shortening at various stages of atherosclerosis, with preferential involvement in advanced vessel pathology and acute vascular syndromes (48–50). However, we acknowledge that our study lacked a comparison group of rural South Africans, so the assumed trajectory in telomere attrition due to the sequel of urbanization must remain speculative.

As shortened telomeres have been implied in CVD risk (51), telomere biology might partly explain the rapidly increasing risk of incident atherothrombotic disease and CVD risk factors in the African population (12–20). The increased CVD risk in South Africans as related to urbanization might be a function of time an individual has been living in an urban environment. We found that the longer participants had been living in the Potchefstroom area, the shorter were their telomeres. However, this relationship disappeared after adjustment for covariates, and it also did not differ by ethnicity. It should be emphasized that “years participants had been living in the area” is a very crude proxy measure of urbanization stress, as this variable does not take into account how participants dealt with the stress. Therefore, more sophisticated measures of urbanization should be used to probe for an association with telomere length in future studies. Additional psychosocial variables, such as cultural differences, might also be important to investigate in this context (52).

The comprehensive assessment of several important correlates of telomere length and the comparable socioeconomic background of participants to equate major health disparities due to differences in the SES were strengths of our study. However, the study also had its limitations. Our results may not transfer to more rural populations in South Africa, those with a lower SES, elderly South African individuals, and those with clinically apparent atherothrombotic disease. Also, we did not assess nutrition status and telomerase activity, both of which might account for differences in telomere length between ethnicities (19,38,43).

Because circulating white cells reproduce rapidly and thus differ in telomere kinetics from, for instance, heart or nerve cells, leukocyte telomeres only can act as a marker, not an indicator of telomere length in other tissues (53).

Taken together, we found strong evidence for shorter telomeres in black South Africans compared with their white counterparts independent of important correlates of telomere length. Genetic factors rather than sociodemographic, HIV-infection-related, cardiometabolic, life-style, and psychosocial ones might have accounted for this ethnic difference. The extent to which relatively shorter telomere length in black South Africans contributes to their increased risk of CVD needs to be explored further.

The authors are grateful to Péter Szabolcs, Chrissie Lessing, and Tina Scholz for their technical assistance and support and to Dr. Oksana Levanes for performing PCR analysis.

Source of Funding and Conflicts of Interest: The study was partly funded by The Metabolic Syndrome Institute, France; the Medical Research Council; National Research Foundation; ROCHE Diagnostics; North-West University; and North-West Department of Education, South Africa. Any opinion, findings, and conclusions or recommendations expressed in this material are those of the authors, and therefore, the National Research Foundation does not accept any liability in regard thereto. The authors declare no conflicts of interest.

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