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Vitamin D–Binding Protein and Vitamin D Status of Black Americans and White Americans


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Abstract

Background—Low levels of total 25-hydroxyvitamin D are common among black Americans. Vitamin D–binding protein has not been considered in the assessment of vitamin D deficiency.

Methods—In the Healthy Aging in Neighborhoods of Diversity across the Life Span cohort of blacks and whites (2085 participants), we measured levels of total 25-hydroxyvitamin D, vitamin D–binding protein, and parathyroid hormone as well as bone mineral density (BMD). We genotyped study participants for two common polymorphisms in the vitamin D–binding protein gene (rs7041 and rs4588). We estimated levels of bioavailable 25-hydroxyvitamin D in homozygous participants.

Results—Mean (±SE) levels of both total 25-hydroxyvitamin D and vitamin D–binding protein were lower in blacks than in whites (total 25-hydroxyvitamin D, 15.6±0.2 ng per milliliter vs. 25.8±0.4 ng per milliliter, P<0.001; vitamin D–binding protein, 168±3 μg per milliliter vs. 337±5 μg per milliliter, P<0.001). Genetic polymorphisms independently appeared to explain 79.4% and 9.9% of the variation in levels of vitamin D–binding protein and total 25-hydroxyvitamin D, respectively. BMD was higher in blacks than in whites (1.05±0.01 g per square centimeter vs. 0.94±0.01 g per square centimeter, P<0.001). Levels of parathyroid hormone increased with...
decreasing levels of total or bioavailable 25-hydroxyvitamin D (P<0.001 for both relationships), yet within each quintile of parathyroid hormone concentration, blacks had significantly lower levels of total 25-hydroxyvitamin D than whites. Among homozygous participants, blacks and whites had similar levels of bioavailable 25-hydroxy vitamin D overall (2.9±0.1 ng per milliliter and 3.1±0.1 ng per milliliter, respectively; P = 0.71) and within quintiles of parathyroid hormone concentration.

**Conclusions**—Community-dwelling black Americans, as compared with whites, had low levels of total 25-hydroxyvitamin D and vitamin D–binding protein, resulting in similar concentrations of estimated bioavailable 25-hydroxyvitamin D. Racial differences in the prevalence of common genetic polymorphisms provide a likely explanation for this observation. (Funded by the National Institute on Aging and others.)

Low levels of total 25-hydroxyvitamin D, which are more common in black Americans than in white Americans, are associated with negative health outcomes in epidemiologic studies.\(^1\)\(^-\)\(^4\) Such studies are responsible for the routine clinical practice of screening for vitamin D deficiency. Among the possible effects of vitamin D deficiency, the strongest evidence is for a role in skeletal disorders,\(^5\)\(^,\)\(^6\) but clinical investigations of vitamin D supplementation to decrease the risk of fracture have been inconclusive.\(^7\)\(^-\)\(^10\)

Because blacks consistently have lower levels of total 25-hydroxyvitamin D than whites, they are frequently given a diagnosis of vitamin D deficiency.\(^11\)\(^-\)\(^13\) Yet, as compared with whites, blacks have higher bone mineral density (BMD) and a lower risk of fragility fracture.\(^14\)\(^-\)\(^16\) Elevated levels of parathyroid hormone, often considered a sensitive marker of vitamin D deficiency, are more common in blacks than in whites.\(^17\) However, the relation between levels of parathyroid hormone and total 25-hydroxyvitamin D may differ in blacks and whites.\(^18\)

Vitamin D–binding protein is the primary vitamin D carrier protein, binding 85 to 90% of total circulating 25-hydroxyvitamin D.\(^19\) The non–vitamin D–binding protein fraction (bioavailable 25-hydroxyvitamin D) consists primarily of albumin-bound 25-hydroxyvitamin D (10 to 15% of total 25-hydroxyvitamin D), with less than 1% of total 25-hydroxyvitamin D in the free form. Vitamin D–binding protein appears to inhibit some actions of vitamin D, because the bound fraction may be unavailable to act on target cells.\(^20\)\(^,\)\(^21\) Common genetic polymorphisms in the vitamin D–binding protein gene produce variant proteins that differ in their affinity for vitamin D.\(^22\)\(^,\)\(^23\) The prevalence of these polymorphisms differs between racial groups.\(^24\)\(^,\)\(^25\) Clinical assays measure the level of total 25-hydroxyvitamin D without distinguishing fractions bound to carrier proteins.

We conducted a study to determine whether vitamin D–binding protein genotypes and concentrations of circulating vitamin D–binding protein differ between black Americans and white Americans, possibly accounting for observed racial differences in manifestations of vitamin D deficiency.
Methods

Study Population

Healthy Aging in Neighborhoods of Diversity across the Life Span (HANDLS) is a population-based cohort study involving 3720 participants that is supported by the Intramural Research Program of the National Institute on Aging.\textsuperscript{26} Study participants, who were 30 to 64 years of age and living in Baltimore at the time of enrollment, were recruited from 13 contiguous U.S. Census tracts. Participants were randomly selected within strata based on age, race, sex, and socio-economic status; those who did not identify themselves as black or white were excluded. The institutional review board of the National Institute of Environmental Health Sciences, National Institutes of Health, approved the protocol. The Partners HealthCare Human Research Committee exempted the present study from the requirement for review. The first and last authors vouch for the accuracy of the data and analyses.

Data Collection

We used cross-sectional data from the HANDLS study that were collected between 2004 and 2008. After providing written informed consent and being interviewed, participants underwent an examination on a mobile research vehicle in which blood was sampled, height and weight were measured, and bone densitometry was performed. Dietary intake of calcium and vitamin D were determined by means of the U.S. Department of Agriculture Automated Multiple-Pass Method. Only participants who completed the examination, including bone densitometry (performed with the use of the Lunar DPX-IQ densitometer [Lunar] and restricted to participants weighing <122.5 kg [270 lb]), and who had sufficient blood samples available were included in the present study (2085 participants) (Fig. S1 in the Supplementary Appendix, available with the full text of this article at NEJM.org). BMD at the femoral neck was used in this study, given its relevance as a risk factor for hip fracture.\textsuperscript{27}

Laboratory Analyses

Blood samples drawn at the examination were stored at −80°C. Levels of total 25-hydroxyvitamin D (D\textsubscript{2} and D\textsubscript{3}) were measured with the use of tandem mass spectrometry (interassay coefficient of variation, 8.6%). Levels of vitamin D– binding protein were measured by means of a commercial enzyme-linked immunosorbent assay (R&D Systems) that uses two monoclonal antibodies in a sandwich format (interassay coefficient of variation, 7.2%). Levels of intact parathyroid hormone were measured with the use of the Elecsys Parathyroid Hormone Immunoassay (Modular Analytics E170, Roche Diagnostics) (interassay coefficient of variation, 2.5%). Calcium levels were corrected for the participant's albumin level as follows: corrected calcium = (measured calcium in mg per deciliter) + [0.8 × (4.0 − serum albumin in g per deciliter)].

Genotyping

DNA samples from the participants were genotyped for two common single-nucleotide polymorphisms (SNPs) in the coding region of the vitamin D–binding protein gene (rs4588...
and rs7041) (see the Methods section in the Supplementary Appendix). These polymorphisms were chosen because of their high prevalence in the general population, their association with race, and their known link with vitamin D–binding protein function.\textsuperscript{22-25} We successfully genotyped samples from 1981 participants (95.0\%).

**Calculation Of Bioavailable 25-Hydroxyvitamin D**

Bioavailable 25-hydroxyvitamin D was defined as circulating 25-hydroxyvitamin D not bound to vitamin D–binding protein, which is analogous to the definition of bioavailable testosterone.\textsuperscript{28} Concentrations of bioavailable 25-hydroxyvitamin D were calculated in 1025 homozygotes, for whom we could use a single genotype-specific binding affinity constant on the basis of the presence of a single vitamin D–binding protein variant (see the Methods section in the Supplementary Appendix).\textsuperscript{22} Calculated concentrations of bioavailable 25-hydroxyvitamin D were validated by direct measurement in a subgroup of homozygous participants with the use of a competitive radioligand-binding assay (see the Methods section and Fig. S2 and S3 in the Supplementary Appendix). Measured and calculated 25-hydroxyvitamin D concentrations were correlated (Pearson's $r = 0.81$ in 32 Gc1F homozygotes and 0.90 in 13 Gc1S homozygotes; $P<0.001$ for both relationships) (Fig. S4 in the Supplementary Appendix).

**Statistical Analysis**

The characteristics of the study participants were compared according to race with the use of t-tests or chi-square tests and are presented as means ±SE or numbers and percentages. Non-normally distributed variables were natural log–transformed for parametric testing. Adjusted means were derived from multivariable linear regression models containing terms for age, sex, body-mass index (BMI), status with respect to poverty (defined as self-reported household income of <125\% of the federal poverty level in 2003), season, smoking status, and calcium intake. Microalbuminuria (defined as a urinary microalbumin-to-creatinine ratio $>30 \mu g$ of microalbumin per milligram of creatinine) was included as a covariate in models predicting levels of vitamin D–binding protein and total 25-hydroxyvitamin D. Squared semi-partial correlation coefficients (expressed as percentages) are presented for multivariable linear regression models exploring variation in total 25-hydroxyvitamin D and vitamin D–binding protein. Total $r^2$ values are presented for unadjusted models and for the overall variance explained in multivariable models.

Chi-square tests were used to compare allele frequencies according to race. Race-stratified linear regression models were used to summarize associations of levels of vitamin D–binding protein and total 25-hydroxyvitamin D with the two SNPs of interest (rs7041 and rs4588). For a subgroup of 774 samples from black participants with complete data from genomewide association studies, we created two models, one including 10 principal components from a discriminant analysis of racial groups and a second with only one covariate, percent African ancestry. Adjustment for population substructure had little effect on the model. Thus, these covariates are not included in reported results.
Participants were divided into quintiles to examine relationships between 25-hydroxyvitamin D measures and markers of vitamin D status (parathyroid hormone level, calcium level, and BMD).

Statistical analyses were conducted with the use of SAS software, version 9.2 (SAS Institute). Two-tailed P values of less than 0.05 were considered to indicate statistical significance, with the exception of the genotype analysis, in which the significance threshold was adjusted for the presence of two SNPs, with P values of less than 0.025 considered to indicate statistical significance.

Results

Characteristics of the Participants

Blacks (1181 participants) and whites (904 participants) were similar in terms of age, sex, BMI, and menopausal status (Table 1). Blacks were more likely than whites to be impoverished, to be active smokers, and to have microalbuminuria. Blacks were less likely than whites to have received a diagnosis of osteoporosis or to have been prescribed osteoporosis therapies. Use of hormone-replacement therapy and medications that affect vitamin D metabolism (e.g., antiepileptic agents and glucocorticoids) was uncommon (Table 1).

Total 25-Hydroxyvitamin D, Vitamin D–Binding Protein, and Markers of Vitamin D Status

Unadjusted levels of total 25-hydroxyvitamin D were lower in blacks than in whites (15.6±0.2 ng per milliliter vs. 25.8±0.4 ng per milliliter, P<0.001) (Fig. 1A). Racial differences in total 25-hydroxyvitamin D levels persisted after multivariable adjustment (17.3±0.3 ng per milliliter in blacks vs. 25.5±0.4 ng per milliliter in whites, P<0.001). There were seasonal differences in 25-hydroxyvitamin D levels (Table S1 in the Supplementary Appendix). Race explained 22.7% of the variation in total 25-hydroxyvitamin D levels in an unadjusted model.

Unadjusted levels of vitamin D–binding protein were lower in blacks than in whites (168±3 μg per milliliter vs. 337±5 μg per milliliter, P<0.001) (Fig. 1B). Racial differences in vitamin D–binding protein levels persisted after multivariable adjustment (169±5 μg per milliliter in blacks vs. 339±5 μg per milliliter in whites, P<0.001). There were seasonal differences in vitamin D–binding protein levels; they appeared to explain 0.5% of the variation in vitamin D–binding protein levels (Table S1 in the Supplementary Appendix). Race explained 30.5% of the variation in vitamin D–binding protein levels in an unadjusted model.

Adjusted mean BMD at the femoral neck was greater in blacks than in whites (1.05±0.01 g per square centimeter vs. 0.94±0.01 g per square centimeter, P<0.001), as were adjusted mean calcium levels (9.11±0.01 mg per deciliter vs. 8.99±0.01 mg per deciliter, P<0.001). Adjusted mean levels of parathyroid hormone were higher in blacks than in whites (39±1 pg per milliliter vs. 34±1 pg per milliliter, P<0.001). When we excluded participants with measurable 25-hydroxyvitamin D₂, our findings did not change appreciably (data not shown).
Genetic Polymorphisms, Vitamin D–Binding Protein, and Total 25-Hydroxyvitamin D

Blacks were more likely than whites to have the T allele at rs7041, whereas whites were more likely than blacks to have the G allele at this location (P<0.001 for both comparisons); blacks were less likely to have the A allele at rs4588 (P<0.001) (Table 2).

The T allele at rs7041 was associated with decreased levels of vitamin D–binding protein in both blacks and whites (Table 2). The A allele at rs4588 was associated with higher vitamin D–binding protein levels in both blacks and whites after we accounted for the allele at rs7041. The polymorphisms at rs7041 and rs4588 had additive effects on vitamin D–binding protein concentrations (Table 2). Genetic variants independently appeared to explain 79.4% of the variation in vitamin D–binding protein levels after we accounted for other factors. After genetic variants were taken into account, race appeared to explain less than 0.1% of the variation in vitamin D–binding protein levels.

The T allele at rs7041 was associated with decreased levels of total 25-hydroxyvitamin D among blacks. In whites, the A allele at rs4588 was associated with decreased levels of total 25-hydroxyvitamin D (Table 2). These genetic polymorphisms appeared to explain 9.9% of the variation in total 25-hydroxyvitamin D levels after other factors were taken into account. In the same model, season and race appeared to explain 10.5% and 7.3% of the variation in total 25-hydroxyvitamin D levels, respectively, whereas sex, age, smoking, calcium intake, BMI, poverty, and microalbuminuria each appeared to account for less than 2.0% of the variation. Overall, 31.2% of the variation in total 25-hydroxyvitamin D levels appeared to be explained in a model containing the aforementioned variables. The concentration of vitamin D–binding protein and the genotype of vitamin D–binding protein appeared to explain a similar amount of variation.

Findings In Homozygous Participants

Vitamin D–Binding Protein Phenotypes and Bioavailable 25-Hydroxyvitamin D

—Figure 2A shows the percentage of homozygous participants in each racial group with each variant vitamin D–binding protein (resulting from unique combinations of rs7041 and rs4588). Vitamin D–binding protein levels were lowest in Gc1F homozygous participants, highest in Gc1S homozygous participants, and intermediate in Gc2 homozygous participants (P<0.001 for all comparisons) (Fig. 2B). Among all 1025 homozygous participants, calculated levels of bioavailable 25-hydroxyvitamin D were similar in blacks and whites (2.9±0.1 ng per milliliter and 3.1±0.1 ng per milliliter, respectively; P = 0.71) (Fig. 2C).

Markers of Vitamin D Status and 25-Hydroxyvitamin D—BMD was not associated with levels of bioavailable or total 25-hydroxyvitamin D in black homozygous participants; however, in white homozygous participants, BMD generally increased with increasing levels of total or bioavailable 25-hydroxyvitamin D (Table S2 in the Supplementary Appendix). Calcium levels increased with increasing levels of total 25-hydroxyvitamin D in blacks only (Table S2 in the Supplementary Appendix). Lower levels of total or bioavailable 25-hydroxyvitamin D were associated with higher levels of parathyroid hormone in homozygotes of both races (P<0.001 for all relationships) (Table S2 in the Supplementary Appendix). As compared with white homozygotes with similar parathyroid hormone levels,
black homozygotes had significantly lower levels of total 25-hydroxyvitamin D (Fig. 3A). In contrast, homozygous blacks and whites with similar parathyroid hormone levels had similar levels of bioavailable 25-hydroxyvitamin D (Fig. 3B). Relationships between total 25-hydroxyvitamin D levels and markers of vitamin D status in homozygotes were similar to those in the overall study population.

Discussion

Because levels of total 25-hydroxyvitamin D are consistently lower in black Americans than in white Americans, blacks are frequently classified as being vitamin D–deficient. In our study involving community-dwelling adults, we found that levels of vitamin D–binding protein are also lower in blacks, probably because of the high prevalence of a common genetic variant. Lower levels of vitamin D–binding protein in blacks appear to result in levels of bioavailable 25-hydroxyvitamin D that are equivalent to those in whites. These data, combined with previous data from our group, suggest that low total 25-hydroxyvitamin D levels do not uniformly indicate vitamin D deficiency and call into question routine supplementation in persons with low levels of both total 25-hydroxyvitamin D and vitamin D–binding protein who lack other traditional manifestations of this condition.

Thresholds for vitamin D sufficiency have been based on total 25-hydroxyvitamin D levels at which calcium absorption declines or parathyroid hormone levels increase. Because experimental data are inconclusive, controversy surrounds the precise level of total 25-hydroxyvitamin D at which these changes occur. We studied a community-dwelling population, in which overt vitamin D deficiency was rare; in fact, few participants had parathyroid hormone levels outside the normal range. Still, on the basis of the current guidelines (suggesting a threshold for sufficiency of 20 or 30 ng per milliliter), 77 to 96% of our black participants would be classified as vitamin D–deficient. Labeling the majority of the black participants as vitamin D–deficient would be inconsistent with the observation that they had higher BMD, higher calcium levels, and only slightly higher parathyroid hormone levels than their white counterparts.

Low levels of vitamin D–binding protein in blacks may provide protection against the manifestations of vitamin D deficiency despite low levels of total 25-hydroxyvitamin D. The bio-availability of other lipophilic hormones, such as thyroid hormone, is known to be influenced by the concentration of carrier proteins. When the concentration of the thyroxine-binding globulin is low or undetectable, there is a lower total thyroid hormone requirement for sufficiency. Analogously, mice that lack vitamin D–binding protein have low levels of total 25-hydroxyvitamin D but do not show signs of vitamin D deficiency. Therefore, low levels of total 25-hydroxyvitamin D probably do not indicate true vitamin D deficiency when levels of vitamin D–binding protein are also low, as in many black Americans. Bioavailable 25-hydroxyvitamin D may be a more appropriate cross-racial marker of vitamin D sufficiency; however, investigations in populations with overt vitamin D deficiency are required before routine clinical use is warranted.
Levels of total 25-hydroxyvitamin D are, in part, genetically determined.\textsuperscript{24,35,36} In our study, genetic polymorphisms in vitamin D–binding protein appeared to account for a greater proportion of the variation in total 25-hydroxyvitamin D levels than most factors known to be associated with 25-hydroxyvitamin D levels.\textsuperscript{35-37} The effect of vitamin D–binding protein polymorphisms on total 25-hydroxyvitamin D concentrations appeared to be mediated by the concentration of vitamin D–binding protein, an observation that is consistent with the findings in a previous study.\textsuperscript{38} Data in genetically modified mice suggest that the function of vitamin D–binding protein is to prolong the half-life of 25-hydroxyvitamin D, supporting this hypothesis.\textsuperscript{20} Although mice lacking vitamin D–binding protein do not have manifestations of overt deficiency at baseline, they are more susceptible to deficiency than normal mice when deprived of vitamin D.\textsuperscript{20} Vitamin D–binding protein prolongs the half-life of 25-hydroxyvitamin D by serving as a reservoir and aiding in the reabsorption of filtered vitamin D through megalin in the kidney.\textsuperscript{20,39} We speculate that low levels of vitamin D–binding protein may confer a predisposition to inadequate 25-hydroxyvitamin D levels when vitamin D sources are scarce. Levels of vitamin D–binding protein only partially explained racial differences in levels of total 25-hydroxyvitamin D; other factors, including skin pigmentation and other polymorphisms, probably contribute to low levels of total 25-hydroxyvitamin D in blacks.\textsuperscript{1,40,41}

Our study has certain limitations. First, given the cross-sectional and observational nature of the study, we were unable to predict the effects of vitamin D–binding protein levels on the risk of fracture. Second, measurement of bone-turnover markers, levels of 1,25-dihydroxyvitamin D, and urinary calcium excretion might have provided additional insight into the effect of vitamin D–binding protein on mineral metabolism. Third, we did not have data on the use of vitamin D supplements. However, when we excluded participants with measurable 25-hydroxyvitamin D\textsubscript{2}, which suggests exogenous supplementation with vitamin D derived from plants or fungi, our findings did not change. Further investigation is needed to determine the effects of supplementation on total and bioavailable 25-hydroxyvitamin D levels in persons with different vitamin D–binding protein genotypes. Finally, we relied predominantly on calculation of bioavailable 25-hydroxyvitamin D rather than direct measurement. Among homozygous participants, however, direct measurement of bioavailable 25-hydroxyvitamin D was well correlated with calculated levels.

There is an alternative commercially available assay for measuring vitamin D–binding protein levels; the results of that assay are inconsistent with those of the assay used in this study (Fig. S5 and S6 in the Supplementary Appendix). The vitamin D–binding protein levels we report correlate inversely with the percentage of bioavailable 25-hydroxyvitamin D measured directly. Given the lack of genotype-specific standards in our direct assay format, we could not accurately report absolute concentrations. Our data should provide an impetus for the development of assays that directly measure bioavailable 25-hydroxyvitamin D.

Vitamin D deficiency is certainly present in persons with very low levels of total 25-hydroxyvitamin D accompanied by hyperparathyroidism, hypocalcemia, or low BMD. However, community-dwelling blacks with total 25-hydroxyvitamin D levels below the threshold used to define vitamin D deficiency typically lack the accompanying characteristic
alterations. The high prevalence among blacks of a polymorphism in the vitamin D-binding protein gene that is associated with low levels of vitamin D-binding protein results in levels of bioavailable 25-hydroxyvitamin D that are similar to those in whites, despite lower levels of total 25-hydroxyvitamin D. Alterations in vitamin D-binding protein levels may therefore be responsible for observed racial differences in total 25-hydroxyvitamin D levels and manifestations of vitamin D deficiency. To improve the determination of vitamin D status in diverse populations, the measurement of vitamin D-binding protein will most likely need to be incorporated into the assessment.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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References


Figure 1. Levels of Total 25-Hydroxyvitamin D and Vitamin D-Binding Protein in Community-Dwelling White and Black Study Participants

Histograms representing stacked distributions are shown. Mean (±SE) levels of total 25-hydroxyvitamin D were significantly lower in blacks than in whites (15.6±0.2 ng per milliliter vs. 25.8±0.4 ng per milliliter, P<0.001) (Panel A), as were levels of vitamin D-binding protein (168±3 μg per milliliter vs. 337±5 μg per milliliter, P<0.001) (Panel B).
Figure 2. Variant Vitamin D–Binding Proteins and Bioavailable 25-Hydroxyvitamin D

As shown in Panel A, unique combinations of the rs7041 and rs4588 polymorphisms produce amino acid changes resulting in variant vitamin D–binding proteins (left side of panel; Asp denotes aspartic acid, Glu glutamic acid, Lys lysine, and Thr threonine). The Gc1F phenotype was most common in black homozygotes, whereas the Gc1S phenotype was most common in white homozygotes (right side of panel). As shown in Panel B, levels of vitamin D–binding protein were lowest in Gc1F/Gc1F homozygotes (632 participants, 93±2 μg per milliliter), highest in Gc1S/Gc1S homozygotes (313 participants, 468±6 μg per milliliter), and intermediate in Gc2/Gc2 homozygotes (80 participants, 190±4 μg per milliliter). Plasma vitamin D–binding protein concentrations in Gc1F/Gc1S heterozygotes (413 participants, 285±4 μg per milliliter) were intermediate between those of Gc1F/Gc1F...
homozygotes and Gc1S/Gc1S homozygotes. These differences were significant (P<0.001 for all comparisons). Panel C shows a histogram representing stacked distributions. Among homozygous participants, levels of bioavailable 25-hydroxyvitamin D were similar in blacks and whites (2.9±0.1 ng per milliliter in blacks and 3.1±0.1 ng per milliliter in whites, P = 0.71).
Figure 3. Total and Bioavailable 25-Hydroxyvitamin D Levels among Homozygous Blacks and Whites with Similar Parathyroid Hormone Levels

Within quintiles of parathyroid hormone values, blacks generally had lower levels of total 25-hydroxyvitamin D levels than whites (Panel A) but similar levels of bioavailable 25-hydroxyvitamin D (Panel B). I bars indicate standard errors. One asterisk denotes P<0.01 for the comparisons between blacks and whites within the quintile, two asterisks P<0.01 for the comparison with the highest quintile among whites, and three asterisks P<0.01 for the comparison with the highest quintile among blacks.
Table 1
Characteristics of the Study Participants Overall and According to Race*

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>Overall (N = 2085)</th>
<th>Blacks (N = 1181)</th>
<th>Whites (N = 904)</th>
<th>P Value</th>
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<tr>
<td>Age — yr</td>
<td>48.3±0.2</td>
<td>48.3±0.3</td>
<td>48.3±0.3</td>
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<td>Male sex — no. (%)</td>
<td>921 (44.2)</td>
<td>523 (44.3)</td>
<td>398 (44.0)</td>
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<tr>
<td>BMI†</td>
<td>29.6±0.2</td>
<td>29.4±0.2</td>
<td>29.8±0.2</td>
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<td>Household income &lt;125% of poverty line — no. (%)</td>
<td>850 (40.8)</td>
<td>573 (48.5)</td>
<td>277 (30.6)</td>
<td>&lt;0.001</td>
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<tr>
<td>Score on Houston Activity Scale‡</td>
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<td>2.2±0.1</td>
<td>2.8±0.1</td>
<td>&lt;0.001</td>
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<tr>
<td>Current smoker — no./total no. (%)</td>
<td>930/1938 (48.0)</td>
<td>552/1091 (50.6)</td>
<td>378/847 (44.6)</td>
<td>0.01</td>
</tr>
<tr>
<td>Diagnosis of osteoporosis — no./total no. (%)</td>
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<td>19/946 (2.0)</td>
<td>32/784 (4.1)</td>
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<td>Prescribed osteoporosis therapies — no. (%)§</td>
<td>29 (1.4)</td>
<td>10 (0.8)</td>
<td>19 (2.1)</td>
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<td>Postmenopausal — no. of women/total no. (%)</td>
<td>623/1104 (56.4)</td>
<td>345/626 (55.1)</td>
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<td>Microalbuminuria — no./total no. (%)¶</td>
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<td>27/710 (3.8)</td>
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<td>Estimated GFR of &lt;60 ml/min/1.73 m² — no./total no. (%)║</td>
<td>114/2039 (5.6)</td>
<td>67/1141 (5.9)</td>
<td>47/898 (5.2)</td>
<td>0.53</td>
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<td>Prescribed antiepileptic agents — no. (%)**</td>
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<td>6 (0.5)</td>
<td>8 (0.9)</td>
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<td>Prescribed glucocorticoids — no. (%)††</td>
<td>28 (1.3)</td>
<td>20 (1.7)</td>
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<td>Dietary calcium intake — mg/day</td>
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</table>

* Race was self-reported. Plus–minus values are means ±SE. P values of less than 0.05 were considered to indicate statistical significance. HRT denotes hormone-replacement therapy.
† The body-mass index (BMI) is the weight in kilograms divided by the square of the height in meters.
‡ Scores on the Houston Activity Scale range from 0 to 7, with higher scores indicating greater habitual physical activity. Data were missing for 785 black participants (66.5%) and 672 white participants (74.3%).
§ Osteoporosis therapies included pamidronate, neridronic acid, olpadronate, alendronate, ibandronate, risedronate, zoledronate, denosumab, teriparatide, and raloxifene.
¶ Microalbuminuria was defined as a urinary microalbumin-to-creatinine ratio of more than 30 μg of microalbumin per milligram of creatinine.
║ The estimated glomerular filtration rate (GFR) was calculated with the use of the Chronic Kidney Disease Epidemiology Collaboration equation.
** Antiepileptic agents included phenobarbital, carbamazepine, phenytoin, and primidone.
†† Glucocorticoids included prednisone, hydrocortisone, methylprednisolone, prednisolone, and dexamethasone.
Table 2
Influence of Genetic Polymorphisms on Levels of Vitamin D–Binding Protein and Total 25-Hydroxyvitamin D

<table>
<thead>
<tr>
<th>SNP</th>
<th>Reference Allele</th>
<th>Variant Allele</th>
<th>Variant Allele Frequency</th>
<th>Change in Vitamin D–Binding Protein Level per Variant Allele Copy (95% CI)</th>
<th>P Value</th>
<th>Change in Total 25-Hydroxyvitamin D Level per Variant Allele Copy (95% CI)</th>
<th>P Value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>Vitamin D–Binding Protein Level (μg/ml)</td>
<td></td>
<td>Total 25-Hydroxyvitamin D Level (ng/ml)</td>
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<td></td>
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<td></td>
<td></td>
<td>Change in Vitamin D–Binding Protein Level per Variant Allele Copy (95% CI)</td>
<td>P Value</td>
<td>Change in Total 25-Hydroxyvitamin D Level per Variant Allele Copy (95% CI)</td>
<td>P Value</td>
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<tr>
<td>Blacks</td>
<td>rs7041</td>
<td>G</td>
<td>T</td>
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<td></td>
<td></td>
<td></td>
<td>−195.7 to −183.1</td>
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<td>−2.9 to −1.1</td>
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<td></td>
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<td></td>
<td>&lt;0.001</td>
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<tr>
<td></td>
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<td>C</td>
<td>A</td>
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<td>57.0</td>
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<td>49.2 to 64.7</td>
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<td>−1.7 to 0.6</td>
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<td>&lt;0.001</td>
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<tr>
<td>Whites</td>
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<td>T</td>
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<td>1.3 to 1.7</td>
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<td>−4.1 to −0.9</td>
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</table>

*Data on 1126 blacks and 855 whites were included in the analysis. CI denotes confidence interval, and SNP single-nucleotide polymorphism. P values of less than 0.025 were considered to indicate statistical significance (adjusted for two SNPs).