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# Nocturnal Asthma and the Importance of Race/Ethnicity and Genetic Ancestry

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## Abstract

**Rationale:** Nocturnal asthma is a common presentation and is associated with a more severe form of the disease. However, there are few epidemiologic studies of nocturnal asthma, particularly in minority populations.

**Objectives:** To identify factors associated with nocturnal asthma, including the contribution of self-identified race/ethnicity and genetic ancestry.

**Methods:** The analysis included individuals from the Study for Asthma Phenotypes and Pharmacogenomic Interactions by Race-ethnicity (SAPPHIRE) cohort. Nocturnal asthma symptoms were assessed by questionnaire. Genome-wide genotype data were used to estimate genetic ancestry in a subset of African American participants. Logistic regression was used to evaluate the association of various factors with nocturnal asthma, such as self-identified race/ethnicity and genetic ancestry.

**Measurement and Main Results:** The study comprised 3,380 African American and 1,818 European Americans individuals with

asthma. After adjusting for other potential explanatory variables, including controller medication use, African Americans were more than twice as likely (odds ratio, 2.56; 95% confidence interval, 2.24–2.93) to report nocturnal asthma when compared with European American individuals. Among the subset of African American participants with genome-wide genotype data ( $n = 1,040$ ), estimated proportion of African ancestry was also associated with an increased risk of nocturnal asthma ( $P = 0.007$ ). Differences in lung function explained a small, but statistically significant ( $P = 0.02$ ), proportion of the relationship between genetic ancestry and nocturnal asthma symptoms.

**Conclusions:** Both self-identified race/ethnicity and African ancestry appear to be independent predictors of nocturnal asthma. The mechanism by which genetic ancestry contributes to population-level differences in nocturnal asthma appears to be largely independent of lung function.

**Keywords:** asthma; nocturnal symptoms; race/ethnicity; lung function; genetic ancestry

Asthma is a common respiratory disorder affecting more than 30 million people in the United States (1) and 300 million people worldwide (2–4). Nocturnal asthma is a common disease presentation, with studies

of asthma reporting up to 75% of patients with nocturnal symptoms (5). Nocturnal asthma has also been associated with disease severity and is accompanied by higher rates of morbidity (6) and mortality (7, 8).

Furthermore, although asthma mortality is relatively low, there is a higher chance of dying at night relative to during the daytime (9), suggesting that nocturnal asthma is a major medical and public health concern.

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## At a Glance Commentary

### Scientific Knowledge on the Subject:

Nocturnal asthma is a common presentation and is associated with a more severe form of the disease. However, there are few epidemiologic studies of nocturnal asthma in minority populations, and none have evaluated the genetic component of ancestry in relationship to the risk of developing nocturnal asthma.

### What This Study Adds to the

**Field:** In a large, ethnically diverse cohort of individuals from southeastern Michigan with asthma, African American individuals were more likely to report nocturnal asthma when compared with European American individuals. Within the subset of African Americans, the estimated proportion of African ancestry was associated with higher likelihood of symptoms, although this relationship was only partially explained by differences in lung function.

Nocturnal asthma is diagnosed in patients based on reported sleep disturbance due to asthma-related symptoms. However, lung function studies have shown that affected individuals may demonstrate reduced FEV<sub>1</sub> both at night and throughout the day (10, 11). The relative decrease appears to be greatest during the nighttime hours (12) and may be accompanied by increased eosinophil inflammation in the lung (13). Individuals with nocturnal asthma may also have higher levels of blood eosinophils and neutrophils relative to individuals with asthma without nocturnal symptoms (14). These findings suggest that nocturnal symptoms may constitute a specific subset of individuals with asthma. Nevertheless, little work has been done to characterize the epidemiology of nocturnal asthma, particularly among African American individuals, who as a group suffer disproportionately from asthma-related complications (15–17).

The current study reports results from an epidemiologic investigation of nocturnal asthma in a multiethnic cohort, the Study for Asthma Phenotypes and Pharmacogenomic Interactions by Race-Ethnicity (SAPPHIRE) cohort. Using this large cohort, we were able to

identify race/ethnicity differences in both the frequency of nocturnal symptoms and the characteristics associated with nocturnal asthma.

## Methods

### Study Subjects

This study was approved by the Institutional Review Board of Henry Ford Hospital and was in compliance with its Health Insurance Portability and Accountability Act policy. The SAPPHIRE cohort is an ongoing study to identify the genetic predictors of asthma controller medication response among a population-based sample of individuals with asthma. Specifically, the cohort includes members of a large health system, which serves southeast Michigan and the greater Detroit metropolitan area. Participants are identified from health care claims and recorded asthma diagnoses in the electronic medical record. The eligibility criteria for SAPPHIRE participants are as follows: age 12 to 56 years, a prior clinical diagnosis of asthma, and no recorded diagnosis of chronic obstructive pulmonary disease or congestive heart failure. Individuals, who consent to participation undergo a detailed enrollment evaluation.

### Measurement of Nocturnal Asthma Symptoms and Accompanying Covariates

All participants completed a detailed staff-administered survey at the time of enrollment. The Asthma Control Test (ACT) (QualityMetric Inc., Lincoln, RI) was included on the survey. The ACT included the following question about nocturnal asthma symptoms: “During the past 4 weeks, how often did your asthma symptoms (wheezing, coughing, shortness of breath, chest tightness or pain) wake you up at night or earlier than usual in the morning?” (18) The possible responses to this question were not at all (= 0), once or twice (= 1), once a week (= 2), two to three nights a week (= 3), or four or more times a week (= 4). We collapsed these responses into a single dichotomous variable for nocturnal symptoms to compare no symptoms (= 0) with any nocturnal symptoms (= 1).

The survey also asked participants to identify all race/ethnicity categories that applied to themselves. Patients were also queried about other demographic

characteristics and environmental exposures, including age, smoking history, and age of asthma onset. Field staff obtained the participants’ height and weight, and these measures were used to calculate body mass index (BMI) in kilograms divided by meters squared (kg/m<sup>2</sup>). Patients had their lung function assessed, and this spirometry was performed according to 2005 American Thoracic Society/European Respiratory Society recommendations using a Fleisch-type pneumotachometer (19). Bronchodilator reversibility (BDR) was quantified as the percentage change in FEV<sub>1</sub> after administration of 360 μg of inhaled albuterol hydroxyfluoroalkane delivered by metered dose inhaler using an AeroChamber Plus Flow-Vu spacer (Monahan Medical Corp., Plattsburgh, NY).

We assessed West African ancestry (heretofore, referred to as African ancestry) in individuals whose self-reported race-ethnicity was African American. To measure genetic ancestry, genomic DNA was first isolated from a blood specimen taken at the time of the initial study visit. Genome-wide genotyping was performed using the Axiom genome-wide AFR array (Affymetrix, Santa Clara, CA) on 1,040 SAPPHIRE subjects with asthma (20). We assessed local ancestry (i.e., African and European ancestry) at each autosomal single nucleotide polymorphism location (21, 22), and we estimated the total percentage of African ancestry (i.e., global ancestry) in African American individuals as the proportion of African alleles.

### Statistical Analysis

We restricted the analysis to African American and European American individuals. The primary outcome variable was nocturnal asthma (i.e., any nocturnal symptoms vs. no symptoms). To test for heterogeneity by race/ethnicity, we compared models with and without a covariate-by-race interaction term—a 1 degree of freedom likelihood ratio test comparing a model with the interaction term to a reduced model with just the individual covariates of interest and a term for self-reported race/ethnicity (i.e., two-variable models). We used forward-stepwise selection to separately build multivariable models of nocturnal asthma for African American and European American individuals. Variables were entered into the model one at a time based

on their likelihood ratio *P* value. A threshold *P* value of 0.05 was used to determine whether variables entered into the model were retained; however, sex, age, and height were included in the model regardless of statistical significance.

Causal mediation analysis (23, 24) was used to test whether the relationship between genetic African ancestry and nocturnal asthma was mediated by lung function (i.e., FEV<sub>1</sub> or FVC). For these models, percent African ancestry was considered the treatment, lung function was considered the mediator, and dichotomous nocturnal asthma status was used as the outcome. As the treatment must be dichotomous for this model, the continuous variable of percent African ancestry was divided into quintiles, and mediation effects were estimated using the subjects in the lowest quintile of percent African ancestry ( $\leq 73.7\%$ ) as the “control” group. Each successively higher quintile was a “treatment” group. Two equations were used to model each mediator and outcome. The mediator equation was estimated using a linear model of lung function with the primary predictor being African ancestry. The outcome equation was estimated using a logistic model for the binary outcome of nocturnal asthma. These models were also adjusted for additional variables associated with nocturnal asthma in African Americans identified from the forward stepwise selection procedure. The confidence intervals (CIs) were based on the nonparametric bootstrap with 1,500 samples.

As a sensitivity analysis, we assessed whether adjusting for inhaled corticosteroid (ICS) and long-acting  $\beta$ -agonist (LABA) medication use at baseline affected the relationships between nocturnal asthma and both self-reported race/ethnicity and African ancestry. We also assessed the effect of adjusting for asthma controller medication adherence in the 6-month period preceding symptom assessment. Medication adherence was quantified using pharmacy claims data in the manner described by us previously (25). We also reassessed the relationship of race/ethnicity and nocturnal asthma symptoms after stratifying by BMI and adjusting for different measures of tobacco smoke exposure.

A flowchart of the analyses performed and the numbers available for each analysis are shown in Figure E1 of the online supplement. The causal mediation analysis was conducted using the “mediation”

package (26) implemented in the R statistical programming language (version 3.0.2) (27). We used SAS version 9.2 (SAS institute Inc., Cary, NC) for all other analyses.

## Results

The demographic and clinical characteristics of the study participants are presented in Table 1. Self-identified African American individuals composed 65% of the total study population. The African American and European American groups were similar in age and sex. Relative to European American participants, African Americans had a higher average BMI (31.8 vs. 28.9), were more likely to have smoked (31% vs. 25%), were more likely to have had asthma before the age of 12 years (64 vs. 56%), and were nearly twice as likely to report a prior hospitalization for asthma (46% vs. 24%). Regarding lung function, mean percent of predicted FEV<sub>1</sub> was lower in African American individuals as compared with European American individuals (87.4% vs. 90.5%), whereas having a BDR value greater than 12% was approximately twice as high in the former group (29.4% vs. 16.9%).

Table 2 presents the unadjusted univariable associations with nocturnal

asthma. Higher measures of lung function (i.e., higher FEV<sub>1</sub>, FVC, and FEV<sub>1</sub>/FVC values) were associated with a lower likelihood of nocturnal asthma symptoms. Conversely, increasing age, female sex, higher BMI, older age at asthma diagnosis, higher BDR, a positive smoking history, and self-reported African American race-ethnicity were all associated with a higher likelihood of reporting nocturnal asthma symptoms. African American individuals were approximately three times more likely to report nocturnal asthma symptoms when compared with European American individuals (odds ratio [OR], 2.95; 95% CI, 2.61–3.34).

As shown in Figure 1, African American subjects also reported greater nights with disrupted sleep from asthma when compared with European American participants (*P* < 0.001). For example, relative to those not experiencing nocturnal symptoms, African American individuals were approximately four times as likely to report having nocturnal symptoms four or more times per week (OR, 4.05; 95% CI, 3.04–5.45) when compared with European American individuals.

To determine if the effect of race-ethnicity could be explained by other factors associated with nocturnal asthma, we

**Table 1.** Characteristics of Study Participants at the Time of Study Enrollment (n = 5,198)

	Combined (n = 5,198)	African American (n = 3,380)	European American (n = 1,818)	<i>P</i> Value*
Age, yr	33.4 ± 14.4	32.8 ± 14.0	34.5 ± 15.1	<0.001
Female,	3,324 (64)	2,183 (65)	1,151 (63)	0.202
Height, m	1.7 ± 0.1	1.67 ± 0.1	1.68 ± 0.1	<0.001
African American race/ethnicity	3,380 (65)	—	—	—
BMI, kg/m <sup>2</sup>	30.8 ± 9.3	31.8 ± 9.5	28.9 ± 8.4	<0.001
Smoking, ever smoker	1,518 (29)	1,061 (31)	457 (25)	<0.001
Age at diagnosis, yr	13.8 ± 13.7	12.9 ± 13.5	15.5 ± 13.8	<0.001
Age at diagnosis > 12 yr	2,027 (39)	1,223 (36.2)	804 (44.2)	<0.001
Ever hospitalized	1,995 (38)	1,556 (46)	439 (24)	<0.001
FEV <sub>1</sub> , L	2.8 ± 0.9	2.6 ± 0.8	3.1 ± 0.8	<0.001
FVC, L	3.6 ± 1.0	3.3 ± 0.9	4.0 ± 1.0	<0.001
FEV <sub>1</sub> /FVC ratio	76.8 ± 9.5	76.2 ± 9.8	77.8 ± 8.8	<0.001
Percent of predicted FEV <sub>1</sub> <sup>†</sup>	88.5 ± 18.9	87.4 ± 19.5	90.5 ± 17.6	<0.001
Percent of predicted FVC <sup>†</sup>	96.2 ± 17.2	96.3 ± 17.7	95.9 ± 16.2	0.382
BDR <sup>‡</sup>	8.3 ± 12.6	9.2 ± 13.6	6.7 ± 10.1	<0.001

*Definition of abbreviations:* BDR = bronchodilator reversibility; BMI = body mass index.

Data are presented as mean ± SD or n (%).

\**P* values for tests of differences between African American and European American individuals.

<sup>†</sup>Based on standardized equations found in Hankinson *et al.* (48).

<sup>‡</sup>BDR was measured as the change in FEV<sub>1</sub> after albuterol administration according to the following formula: BDR = [(FEV<sub>1</sub>(post-bronchodilator) - FEV<sub>1</sub>(pre-bronchodilator)]/FEV<sub>1</sub>(pre-bronchodilator) × 100.

**Table 2.** Assessment of Factors for Association with Nocturnal Asthma and Potential Differences by Race/Ethnicity

	All Subjects		African American		European American		$P_{int}^{\dagger}$
	OR*	95% CI	OR*	95% CI	OR*	95% CI	
Age at enrollment, yr	1.02	1.02–1.03	1.03	1.03–1.04	1.02	1.01–1.03	1.000
Race/ethnicity, African American	2.95	2.61–3.34	—	—	—	—	—
Sex, female	1.61	1.43–1.82	1.64	1.43–1.89	1.91	1.30–2.00	0.940
Height, m	0.22	0.13–0.40	0.32	0.16–0.64	0.23	0.08–0.68	0.620
BMI, kg/m <sup>2</sup>	1.04	1.03–1.05	1.03	1.02–1.03	1.05	1.04–1.07	<0.001
Smoking, ever smoker	2.51	2.22–2.83	2.71	2.32–3.16	2.00	1.60–2.51	0.028
Age at diagnosis, yr	1.01	1.00–1.01	1.01	1.01–1.02	1.01	1.00–1.02	0.230
Age at diagnosis > 12 yr	1.28	1.15–1.44	1.49	1.29–1.71	1.31	1.06–1.61	0.378
Ever hospitalized, yes	2.22	1.98–2.49	1.78	1.55–2.04	2.17	1.73–2.73	0.142
FEV <sub>1</sub> , L	0.47	0.43–0.50	0.52	0.48–0.57	0.54	0.47–0.62	0.668
FVC, L	0.60	0.57–0.64	0.68	0.63–0.73	0.67	0.60–0.75	0.917
FEV <sub>1</sub> /FVC ratio	0.96	0.95–0.96	0.96	0.95–0.97	0.97	0.96–0.98	0.164
Bronchodilator reversibility	1.03	1.03–1.04	1.03	1.02–1.03	1.05	1.04–1.06	0.002

*Definition of abbreviations:* BMI = body mass index; CI = confidence interval; OR = odds ratio;  $P_{int}$  =  $P$  value from a 1 degree of freedom likelihood ratio test of the multiplicative interaction between self-reported race/ethnicity and each variable.

The outcome variable for the association analyses was nocturnal asthma defined as having any night with disrupted sleep due to asthma in the preceding 4 weeks.

\*ORs represent the univariable relationship with nocturnal asthma. These estimates reflect the effect of a one-unit increase in the variable of interest.

<sup>†</sup>The  $P$  value for the likelihood ratio test of interaction between race/ethnicity and each variable in a bivariable model, which includes a covariate for self-reported race/ethnicity.

constructed two multivariable models. The first model included all factors, and the second was constructed using a forward-stepwise variable selection approach (see Table E2). In both models, African American race/ethnicity was associated with nocturnal symptoms (OR, 2.48; 95% CI, 2.12–2.89; and OR, 2.56; 95% CI, 2.24–2.93, respectively). Together, these results suggested that race-ethnicity had an independent association with nocturnal asthma and could not be fully explained by the other factors.

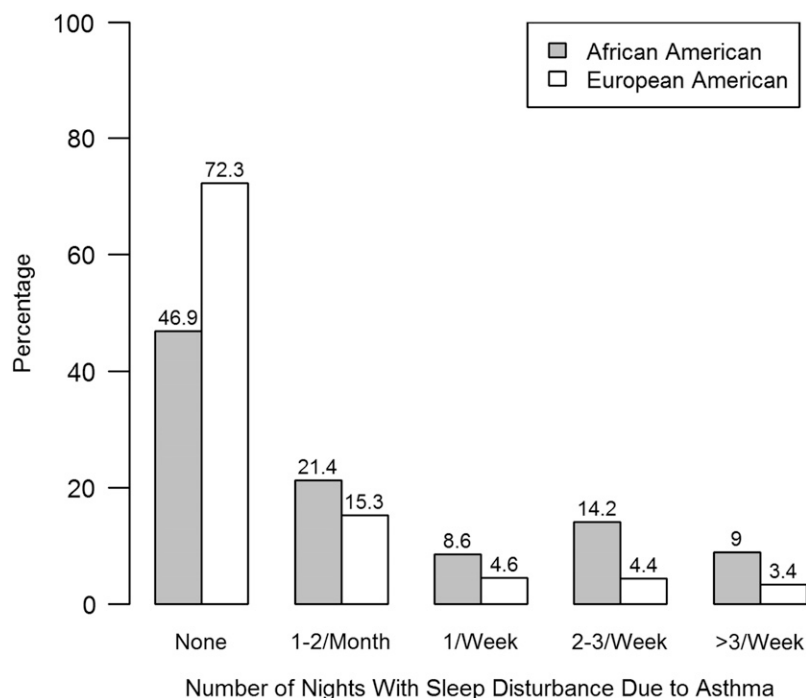
In light of the aforementioned differences by race/ethnicity, we used forward-stepwise variable selection to determine the factors associated with nocturnal asthma for African American patients and European American patients, separately (Table 3). Factors associated with nocturnal symptoms in African American individuals with asthma included current age, sex, BMI, smoking status, history of asthma hospitalization, BDR, FEV<sub>1</sub>, and FVC. With the exceptions of FVC and age, the same factors were associated with nocturnal asthma in European American individuals (i.e., sex, BMI, smoking status, history of asthma hospitalization, BDR, and FEV<sub>1</sub>), and the magnitude of the effect estimates was also similar.

Given the differences in nocturnal asthma by race/ethnicity, we also assessed whether African ancestry was associated with nocturnal symptoms among African

American study participants. For this analysis, the sample was restricted to the subset of African American participants with genome-wide genotype data ( $n = 1,040$ ). Association analysis revealed an increasing trend in proportion of African ancestry with increasing frequency of nocturnal asthma symptoms ( $P = 0.007$ ). The average proportion of African Ancestry was 0.79 ( $\pm 0.10$  SD), 0.80 ( $\pm 0.09$  SD), and 0.81 ( $\pm 0.10$  SD) for patients reporting no symptoms, symptoms less than one night per week, and symptoms one or more nights per week, respectively. After adjusting for all variables retained in the stepwise model for African American individuals (Table 3), increasing African ancestry was positively associated with nocturnal symptoms (OR, 3.47; 95% CI, 0.90–13.39;  $P = 0.069$ ), although the statistical significance was reduced. This reduction in statistical significance may have been due to the effect of outlier individuals with a lower proportion of African ancestry (Figure E2). Mitigating the effect of outliers by grouping individuals into quintiles based on their proportion of African ancestry (i.e., 20th percentile = 73.7%; 40th percentile = 79.2%; 60th percentile = 83.2%; 80th percentile = 87.0%) resulted in a statistically significant association between African ancestry and nocturnal asthma ( $P = 0.045$ ), even after accounting for potential confounders.

We used causal mediation analysis to examine whether the observed relationship between African ancestry and nocturnal symptoms was likely to be a direct effect or mediated by changes in lung function (as measured by FEV<sub>1</sub> or FVC). Due to the high correlation between FEV<sub>1</sub> and FVC, these measures were examined in separate models. The results of these analyses are presented in Table 4. For both FEV<sub>1</sub> and FVC, the findings support a small, but statistically significant, mediating effect of lung function on the relationship between African ancestry and nocturnal asthma. The strongest support for a mediating effect was achieved for FVC, with statistically significant evidence ( $P \leq 0.02$ ) across all four quintiles (i.e., the “treatment” groups) relative to the lowest quintile (i.e., the “control” group). For the group with the highest proportion of African ancestry, 13% of the total effect of ancestry on nocturnal asthma appeared to be mediated by FVC ( $P = 0.04$ ) (Table 4). The mediating effect of lung function appeared to be consistent across all of the other quintile groups defined by African ancestry (Table 4). These findings support an independent effect of African ancestry on nocturnal asthma.

We performed a number of sensitivity analyses to test whether our findings were robust. First, we assessed whether use of ICS or LABAs may have confounded the



**Figure 1.** Percentage of study participants reporting nocturnal asthma symptoms by race/ethnicity. Strata represent the number of nights with asthma-related sleep disturbance in the preceding month. The reported relationships are unadjusted for other variables.

association that we found between race-ethnicity and nocturnal asthma symptoms. For this analysis, we analyzed the subset of individuals for whom we had longitudinal

clinical records of medication use (992 African American patients and 637 European American patients). Among the African American patients, 59 (6.0%) were

using an ICS alone and 112 (11.3%) were using an ICS/LABA combination inhaler. Among the European American patients, 54 (8.5%) were using an ICS alone and 90 (14.1%) were using an ICS/LABA combination inhaler. Accounting for ICS and LABA use had almost no effect on the association between self-reported African American race/ethnicity and nocturnal asthma symptoms (OR, 1.96 for African American vs. European American race-ethnicity; 95% CI, 1.50–2.56;  $P < 0.001$ , before adjusting for ICS and LABA use; and OR, 1.96; 95% CI, 1.50–2.56;  $P < 0.001$ , after adjusting for ICS and LABA use). Similarly, African ancestry was still significantly associated with nocturnal symptoms among African American individuals after adjusting for ICS and LABA use ( $P = 0.027$ ). Among individuals who filled an ICS prescription, accounting for the level of medication adherence did not substantively affect the relationship between race/ethnicity and nocturnal symptoms (OR, 2.24 for African American vs. European American race/ethnicity; 95% CI, 1.22–4.13;  $P = 0.009$ ).

Using different measures of tobacco smoke exposure (i.e., cigarette pack-years and separate variables for past and current cigarette smoking) also did not substantively change the relationship between race-ethnicity and nocturnal symptoms (data not shown). Moreover, restricting the sample to normal-weight individuals (i.e., BMI, 18.5–24.9;  $n = 1,440$ ), we found that race-ethnicity was even more strongly associated with nocturnal symptoms (OR, 2.95 for African American vs. European American race/ethnicity; 95% CI, 2.20–3.94;  $P < 0.001$ ); however, among overweight individuals (BMI  $\geq 25$ ), race/ethnicity was still associated with nocturnal asthma (OR, 1.94; 95% CI, 1.63–2.30;  $P < 0.001$ ).

**Table 3.** Factors Associated with Nocturnal Asthma after Stepwise Selection among Race/Ethnic Groups

Covariate	African American*			European American*		
	OR <sup>†</sup>	95% CI	P Value	OR	95% CI	P Value
Age at enrollment, yr	1.01	1.01–1.02	<0.001	1.00	1.00–1.01	0.620
Sex, female	1.32	1.09–1.61	0.006	1.32	0.97–1.79	0.075
Height, m	1.30	0.42–4.03	0.644	1.52	0.27–8.46	0.633
BMI, kg/m <sup>2</sup>	1.01	1.01–1.02	<0.001	1.04	1.03–1.05	<0.001
Smoking, ever smoker	2.13	1.80–2.52	<0.001	1.55	1.22–1.97	<0.001
Age at diagnosis, yr	—	—	—	—	—	—
Age at diagnosis > 12 yr	—	—	—	—	—	—
Ever hospitalized, yes	1.56	1.35–1.81	<0.001	1.75	1.37–2.23	<0.001
FEV <sub>1</sub> , L	0.55	0.41–0.74	<0.001	0.75	0.62–0.93	0.009
FVC, L	1.29	1.09–1.65	0.043	—	—	—
FEV <sub>1</sub> /FVC ratio	—	—	—	—	—	—
Bronchodilator reversibility	1.01	1.01–1.02	<0.001	1.03	1.02–1.05	<0.001

*Definition of abbreviations:* BMI = body mass index; CI = confidence interval; OR = odds ratio.

The outcome variable for the association analyses was nocturnal asthma defined as having any night with disrupted sleep due to asthma in the preceding 4 weeks.  $P$  value from a 1 degree of freedom likelihood ratio test for each variable.

\*Forward stepwise used for variable selection for the race/ethnicity specific models. Covariates without values (i.e., —) were those not retained in the final model.

<sup>†</sup>ORs reflect the estimated effect of a one-unit increase in the variable of interest, accounting for the other variables retained in the model.

## Discussion

The current study has demonstrated large and statistically significant differences in nocturnal asthma symptoms between African American and European American individuals enrolled in the SAPHIRE cohort. Moreover, we show that genetic ancestry may independently contribute to the risk of having nocturnal asthma.

To our knowledge, there has been only one previous study that has compared

**Table 4.** Causal Mediation Analysis Results Evaluating FEV<sub>1</sub> and FVC as Potential Mediating Traits between African Ancestry and Nocturnal Asthma among African American Participants (n = 1,040)

Mediator	% African Ancestry*	Average Mediation Effect			Average Direct Effect			% Mediated	
		Estimate	95% CI	P Value	Estimate	95% CI	P Value	%	P Value
FEV <sub>1</sub>	>20th to 40th	0.002	6.1 × 10 <sup>-5</sup> to 0.01	0.04	0.021	-4.5 × 10 <sup>-4</sup> to 0.04	0.06	9.0	0.08
	>40th to 60th	0.004	5.4 × 10 <sup>-5</sup> to 0.01	0.05	0.042	2.1 × 10 <sup>-3</sup> to 0.04	0.04	8.9	0.07
	>60th to 80th	0.007	-1.1 × 10 <sup>-4</sup> to 0.02	0.05	0.063	4.0 × 10 <sup>-3</sup> to 0.12	0.03	8.8	0.07
	>80th	0.009	-1.9 × 10 <sup>-4</sup> to 0.20	0.06	0.081	5.3 × 10 <sup>-3</sup> to 0.16	0.04	8.7	0.08
FVC	>20th to 40th	0.003	5.0 × 10 <sup>-4</sup> to 0.01	0.01	0.020	-1.2 × 10 <sup>-4</sup> to 0.04	0.05	12.4	0.04
	>40th to 60th	0.006	9.8 × 10 <sup>-4</sup> to 0.01	0.02	0.040	-2.5 × 10 <sup>-3</sup> to 0.08	0.06	12.7	0.05
	>60th to 80th	0.009	1.5 × 10 <sup>-3</sup> to 0.02	0.01	0.060	3.4 × 10 <sup>-4</sup> to 0.12	0.05	12.7	0.03
	>80th	0.012	2.1 × 10 <sup>-3</sup> to 0.01	0.02	0.079	-9.8 × 10 <sup>-4</sup> to 0.16	0.05	13.0	0.04

Definition of abbreviation: CI = confidence interval.

Nocturnal asthma was defined as having any night with disrupted sleep due to asthma in the preceding 4 weeks. Mediation effects were estimated using the subjects in the lowest quintile of percent African ancestry (i.e., ≤73.7%) as the “control” group and each successive higher quintile as the “treatment” group. The CIs for both the mediation and direct effect estimates were based on the nonparametric bootstrap with 1,500 samples. The mediation effect was estimated using a linear model of lung function (i.e., FEV<sub>1</sub> or FVC) and a logistic model for the binary outcome of nocturnal asthma.

\*The quintiles of % African ancestry were defined as follows: 20th percentile = 73.7%; 40th percentile = 79.2%; 60th percentile = 83.2%; and 80th percentile = 87.0%.

nocturnal asthma rates between African Americans and European Americans. In contrast to our present findings, Trochtenberg and colleagues studied 27 individuals and reported a higher overall prevalence of nocturnal symptoms (82%), with European American patients having a higher occurrence when compared with African American patients (100% vs. 67%, respectively) (28). However, the average age of these study participants was higher (53 and 44 yr for African American and European Americans, respectively) when compared with those enrolled in the SAPPHERE cohort. As we have found increasing age to be positively associated with the risk of reporting nocturnal asthma symptoms, demographic factors may explain a portion of the between study differences. Furthermore, the estimates of this earlier study are inherently less precise given the smaller number of participants.

The current study is also the first to investigate the possible role of genetic ancestry as a risk factor for nocturnal asthma. To date, genetic studies of nocturnal asthma have focused primarily on candidate genes, such as the β<sub>2</sub>-adrenergic receptor (29). Our findings suggest genetic differences between nocturnal and nonnocturnal asthma that may be spread throughout the genome and that this genetic variation is associated with ancestral genetic background. Genome-wide admixture mapping and association studies of nocturnal asthma have not been published to date but appear warranted.

In addition, we show that the effect of genetic ancestry on nocturnal symptoms appears to be mediated in part, but not completely, through its association with lung function (i.e., FVC and FEV<sub>1</sub>). These results are consistent with our earlier work, which showed African ancestry to be inversely associated with measured FVC and FEV<sub>1</sub> (30) and with results showing nighttime lung function to be particularly decreased in individuals with nocturnal asthma relative to those without nocturnal symptoms (10). Although the causal mediation analysis demonstrated a marginally more significant result for FVC as compared with FEV<sub>1</sub>, both FVC and FEV<sub>1</sub> are highly correlated. Therefore, we are unable to conclude which of these lung function measures is the primary mediator of nocturnal symptoms. However, the overall consistency of these findings suggests that the genetic predictors of lung function may be partially shared with the phenotype of nocturnal asthma. As such, a future genome-wide study might benefit from a multitrait mapping approach that includes lung function, given the potential mediating effect of lung function identified and power increases associated with the multitrait over single-trait approaches (31, 32).

According to current U.S. guidelines, individuals in the age range of SAPPHERE participants (i.e., ≥12 yr of age) should consider step-up therapy for uncontrolled or moderately severe persistent asthma when nighttime asthma awakenings occur more than once per week (33). The

guidelines also indicate that combination ICS and LABA therapy be considered for individuals with this level of asthma severity. However, these recommendations are tempered by conflicting findings in the literature, which suggest both a beneficial (34, 35) and detrimental effect (36) of supplemental LABA use, particularly among African American individuals. As a sensitivity analysis in a subset of SAPPHERE participants with longitudinal prescription information, we found that self-reported African American race-ethnicity was still a risk factor for nocturnal symptoms even after accounting for LABA and ICS use. Therefore, it is unlikely that differences in asthma controller medication use alone could explain the observed differences in nocturnal symptoms.

Current asthma treatment guidelines use a symptom-based approach for assessing asthma severity and directing therapeutic decisions (33), and it is not clear that these treatment algorithms would be improved by incorporating information on patient race/ethnicity or genetic ancestry. Nevertheless, our observations provide the impetus for investigating the mechanisms through which race/ethnicity and genetic ancestry contribute to nocturnal asthma. The knowledge gained by these additional studies and this one may, in turn, identify unique and actionable disease mechanisms.

This study is not without other limitations. First, the study population comprised individuals from a single, large health system serving southeast Michigan and metropolitan Detroit. However, we have

previously shown that our patient population is representative of the census population of the region (37), and the proportion of African and European ancestry in our African American participants is similar to that which has been described for African American communities in other parts of the United States (38, 39). Second, it is possible that some cases of nocturnal asthma were incorrectly diagnosed. We would expect that this misclassification would decrease our power to detect associations. However, differences in symptom frequency by race-ethnicity could still be attributable to other factors contributing to nocturnal symptoms, such as obstructive sleep apnea and gastroesophageal reflux disease (40, 41). Given the relationship of these conditions with body weight (42, 43), we stratified our analyses by BMI. However, the association of race/ethnicity with nocturnal symptoms was still present

among individuals with BMI less than 25. Population-level differences in allergic sensitization by race/ethnicity are well described (44, 45) and also could contribute to asthma severity (46, 47). Therefore, future analyses could investigate these and other potential mechanisms (e.g., sleep studies, allergic sensitivity testing, and nocturnal spirometry). Last, our definition of nocturnal asthma was based on responses to an ACT question assessing nighttime or early morning awakenings over the course of 4 weeks (18). It is uncertain whether this is the correct time course for labeling one as having nocturnal asthma, because undoubtedly some patients may have had previous symptoms. Nevertheless, we have no reason to believe that the time course for assessment would have influential differences by race-ethnicity, especially because our findings appeared to be independent of patient age and asthma duration.

In conclusion, the current study provides further insight into the epidemiological factors associated with nocturnal asthma among African American and European American individuals. In particular, this study found that self-identified race/ethnicity appears to be an independent risk factor for nocturnal asthma, even after accounting for likely confounders, such as patient age, sex, BMI, age of asthma onset, lung function, and controller treatment. We also found African ancestry to be associated with nocturnal symptoms among African American individuals, further supporting our findings. Therefore, further studies are needed to dissect the genetic architecture of nocturnal asthma and the mechanisms contributing to the observed disparity. ■

**Author disclosures** are available with the text of this article at [www.atsjournals.org](http://www.atsjournals.org).

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