ORIGINAL ARTICLE

Phenotypes of obstructive sleep apnea in the Hispanic Community Health Study/Study of Latinos

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Abstract

Study Objectives: Recent work on US Whites from clinical samples used obstructive sleep apnea (OSA) symptoms to generate phenotypes for individuals with moderate-severe OSA which suggested 3 to 5 symptom classes. However, it is unknown whether similar classes generalize to diverse Hispanics/Latino adults. Therefore, we sought to fill this gap by empirically deriving sleep phenotypes among a large sample of diverse Hispanics/Latinos.

Methods: We used data from The *Hispanic Community Health Study/Study of Latinos* (HCHS/SOL; 2008–2011), a prospective cohort study designed using a multisite multistage probability sample of adults 18–74 years old. The subpopulation of interest included participants with moderate-severe OSA symptoms (≥15 respiratory event index (REI) events per hour; $n = 1,605$). We performed latent class analysis for complex survey data using 15 common OSA symptoms (e.g. Epworth Sleepiness Scale) and 4 comorbidities to identify phenotype classes.

Results: Average age was 52.4 ± 13.9 years and 34.0% were female. Mean REI was 33.8 ± 22.5 events per hour. Fit statistics and clinical significance suggested that a three-class solution provided the best fit to the data. The three phenotypes were: (1) *Minimally Symptomatic* (47.7%), (2) *Excessive sleepiness* (37.1%), and (3) *Disturbed Sleep* (15.2%). Sensitivity models were consistent with the main proposed solution.

Conclusions: Derived sleep phenotypes among diverse Hispanic/Latinos were consistent with recent findings from the Sleep Apnea Global Interdisciplinary Consortium, but we found notable differences in class prevalence relative to Whites. Further research is needed to link derived sleep phenotypes to health comorbidities in diverse populations.

Statement of Significance

Sleep apnea subtypes of diverse Latinos are not well understood in sleep medicine. We used data from a large, diverse, and representative cohort study of US Latinos to identify sleep apnea phenotypes and link them to sociocultural characteristics and comorbid health conditions. Our study provides data-driven validated sleep apnea phenotypes like the Sleep Apnea Global Interdisciplinary Consortium, but with notable differences compared to non-Hispanic whites. We propose that the empirically derived sleep apnea phenotypes reported herein will help our understanding of sleep apnea among diverse Latinos and culturally tailor precise therapeutic approaches.

Introduction

The United States (US) population is becoming increasingly diverse. Hispanics/Latinos represent 18.0% of the US population and are projected to increase rapidly in coming decades [1]. Health disparities among Hispanics/Latinos are evident; including higher rates of diabetes and cognitive impairment compared to Whites [2, 3]. Sleep disorders may precede these health disorders and have been associated with a higher risk for heart failure, glucose intolerance, and decline in cognitive function [4–6]. In addition to its health sequelae, sleep disturbances also place a high economic burden. Recent estimates indicate that close to 450 billion dollars of economic output in the US are lost to sleep insufficiency alone [7]. Therefore, more research is needed to characterize sleep within the Hispanic/Latino population.

Demographic factors may influence sleep within the Hispanic/Latino population. For example, data from the Hispanic Community Health Study/Study of Latinos (HCHS/SOL) indicate that 25% of Puerto Ricans and 20% of Dominicans and South Americans sleep less than 7 h a day [8]. Another HCHS/SOL study found that Central Americans and Cubans had the most sleepiness symptoms, while Dominicans and Puerto Ricans had the least [9]. Gender/sex differences have also been observed: Mexican men had longer sleep duration compared to Mexican women [10], and are more likely to have obstructive sleep apnea (OSA) compared to women [9].

The prevalence of OSA in predominantly non-Hispanic whites ranges from 6% to 17% [11] and was up to 49% in older adults. Of importance, the home sleep apnea test, allow to measure the respiratory event index (REI), as the number of apnea and hypopnea events in an hour, and the % measures the oxygen desaturation during these events. The prevalence of OSA (REI3% ≥ 15) was 9.8% in middle-aged and older Latinos. The same study attributed increased hypertension and diabetes in Latinos to untreated OSA [9]. Indeed, OSA has also been linked to increased mortality and several cardiovascular risks such as heart failure and stroke [12, 13].

The presentation of OSA symptoms varies across individuals. Recent work has uncovered important heterogeneities in symptoms across patients with OSA and suggested that subgroups can be described using symptom profiles. For example, Ye et al. used latent class analysis (LCA) on 19 sleep symptoms and 4 comorbid conditions to derive 3 distinct derived sleep phenotypes using the *Icelandic Sleep Apnea Cohort* (ISAC) [14]. Their findings suggest the existence of three groups with select symptoms consistent with (1) minimally symptomatic, (2) disturbed sleep, and (3) excessive sleepiness. Follow-up studies from the ISAC cohort have shown that the effectiveness of treatment differs across OSA subtypes [15]. Kim et al. confirmed these three classes in a community-dwelling Korean population-based study [16]. More recently, data from the *Sleep Apnea Global Interdisciplinary Consortium* (SAGIC) uncovered five classes, including the three reported in Ye et al., in a primarily non-Hispanic white (58.8%) and Asian (21.1%) sample [17]. The two additional classes include individuals with (1) upper airway symptoms dominant and (2) sleepiness dominant. The Sleep Heart Health Study (SHHS) [18], in addition to the three main clusters, found a moderately sleepy group. The Canadian Study [19] found four clusters: the three main clusters and an "excessive sleepiness with disturbed sleep group." To date, there is a paucity of data on OSA symptoms in diverse Hispanic/Latino populations.

To address research gaps in characterizing sleep among diverse Hispanics/Latinos, we used data-driven approaches to extract OSA subtypes from a large community-dwelling sample of Hispanics/Latinos in the US. We hypothesized that the three groups found by Ye et al. [14] and other researchers will provide a good fit to the data. However, we expected that the prevalence of these phenotypes would vary relative to non-admixed non-Hispanic white populations, and that women and Hispanics/Latinos of Puerto Rican and Dominican heritage would have symptom profiles closer to the disturbed sleep group proposed by Ye and colleagues (e.g. difficulties falling asleep, waking up at night, and issues falling back asleep after waking up).

Methods

Data

We used baseline data from the HCHS/SOL, an ongoing population-based prospective cohort study of diverse Hispanics/Latinos. Data were collected from four major US metropolitan areas with substantial Hispanic/Latino concentrations (Bronx, NY; Chicago, IL; Miami, FL; and San Diego, CA) and include targeted area representation of the following Hispanic/Latino heritage: Central American, Cuban, Dominican, Mexican, Puerto Rican, and South American. Each site enrolled around 4,000 community-dwelling adults, for a total of 16,415 participants (18–74 years). Detailed methodological discussions of the HCHS/SOL sampling design and methods have been published elsewhere [20]. Institutional review boards at each site approved the study protocol and participants provided written informed consent.

Sleep questionnaires

The original aims of the HCHS/SOL sleep study were to evaluate the associations between sleep apnea and cardiovascular risks among Hispanics/Latinos. Questionnaires were administered in English or Spanish, depending on the participant's preferred language. A modification of the SHHS Sleep habits questionnaire, the International Restless Leg Syndrome questionnaire, [21] as well as the Epworth Sleepiness Scale questionnaire were administered to assess different health risk factors, including OSA. As described elsewhere, participants used an ARES Unicorder 5.2; B-Alert (Carlsbad, CA) to quantify the REI of 3% oxygen desaturation (REI3%) and 4% oxygen desaturation (REI4%) [9].

Sleep variables

In line with published work [14, 16–19], we accounted for 15 OSA symptoms and 4 comorbid health factors. A detailed list of these measures and their scales is included in Supplementary Table S1. We included covariates relating to sleep patterns (e.g. waking up during sleep, napping, having trouble while falling asleep), covariates relating to sleepiness (e.g. dozing off while watching television, dozing off while driving), insomnia (e.g. having trouble falling asleep), and four major comorbidities (i.e. hypertension, obstructive lung disease, cardiovascular disease, and diabetes).

All variables were dichotomized so that individuals reporting a symptom as present once a week. The Epworth Sleep Scale (ESS) was treated as a continuous measure. We classified individuals as meeting criteria for restless legs syndrome if they reported (1) experiencing a desire to move due to discomfort, (2) they felt they needed to move to relieve their discomfort, and (3) if symptoms were worse later in the day or at night. Participants who responded "don't know" were classified as not having the symptom and those who refused to answer as missing. Individuals with three or more missing covariates were excluded from the analysis. We did not perform imputation on missing covariates since LCA calculates estimates based on all available information.

Sociodemographic variables. Sociodemographic variables include age, sex, self-reported Hispanic/Latino heritage (Central American, Cuban, Dominican, Mexican, Puerto Rican, South American, other/mixed), language preference (English, Spanish), and acculturation (language and social acculturation). Previous research characterizes acculturation as "psychological, behavioral and attitudinal changes that occur when individuals and groups from different cultures come into prolonged contact with each other and whereby individuals adopt attitudes, values, customs, beliefs, and behaviors of another culture" [22, 23]. Acculturation, specifically negative acculturation, has been linked to accelerated decline in health and increase in morbidities [24, 25].

Cardiovascular risk variables. Cardiovascular risk variables include high-density lipoprotein (mg/dL), triglycerides (mg/dL), total cholesterol (mg/dL), body mass index (BMI; kg/m^2), and presence or absence (binary) of the following: current smoker, current alcohol use, hypertension (National Health and Nutrition Examination Survey definition [26]), type 2 diabetes (American Diabetes Association definition [27]; fasting glucose ≥ 126 mg/dL, post-OGTT ≥ 200 ng/DL, or A1C ≥ 6.5%), obstructive lung disease (Spirometry was performed using American Thoracic Society and European Respiratory Society guidelines [28]. HCHS/SOL reference equations for spirometry have also been published [29]; FEV_1/FVC < 0.70 or lower limit.), self-reported cardiovascular disease (CVD) including coronary heart disease (based on angina, heart attack, and coronary heart disease), self-

reported heart failure, Framingham CVD composite criterion [30], self-reported stroke or transient ischemic attack (TIA), and self-reported physical and mental health composite scores (12-item Short-Form Health Survey; SF-12 [31]).

Analytic sample

From the 16,415 participants in the cohort, n = 1,138 did not participate in the sleep study and an additional n = 808 had insufficient sleep data for analysis. We included n = 1,623 with moderate to severe OSA (REI3% ≥ 15 events per hour). We further excluded n = 18 participants who did not complete the sleep questionnaire. The final analytic sample was n = 1,605. Included participants were more likely to be male, less likely to have graduated high school, had higher BMI, and were on average 12-years older compared to the overall population, consistent with known risk factors for OSA.

Analysis

Mplus Automation software version 8 was used to fit a series of latent class models iteratively up to 10 potential class solutions. LCAs are probabilistic models and have been used to create classification in other clinical settings, which were used to generate the phenotypes in this study [32, 33]. The first model included the sleep variables as well as four comorbidities: hypertension, diabetes, presence of self-reported cardiovascular disease, and obstructive lung disease. All LCA models accounted for the HCHS/SOL survey weights to adjust for nonresponse bias and allow generalization to the target population. We used full information maximum likelihood to fit the LCAs and all models also accounted for data clustering and stratification to reflect the complex sampling design of the HCHS/SOL data.

In line with published recommendations on assessment on the number of classes derived from LCA, we used several statistical fit indices to adjudicate model fit across the iterated solutions [34]. We used the Bayesian information criterion, Akaike information criterion, entropy, and Vuong-Lo-Mendell Rubin to determine the optimal solution. Results from the LCA can be found in Table 1. For the chosen solution, we generated summary tables to characterize the sleep symptoms and sleep health sociodemographic and general health conditions, and

Table 1. Latent class analysis model fit statistics of sleep phenotypes in individuals with moderate to severe sleep apnea (REI ≥ 15). Unweighted n = 1,605

Solution	LL	Scaling correction factor	Free parameters	AIC	BIC	SSABIC	Entropy	VLMR P	LMR P	AICC
C2	−20,949.12	2.2891	44	41,986.24	42,223	42,083.22	0.766	0.000	0.000	41,988.78
C3	−20,453.12	2.2333	66	41,038.23	41,393.37	41,183.7	0.825	0.041	0.043	41,043.98
C4	−20,232.00	2.2434	88	40,639.99	41,113.51	40,833.95	0.789	0.500	0.502	40,650.32
C5	−20,092.13	2.0964	110	40,404.26	40,996.16	40,646.71	0.804	0.311	0.312	40,420.61
C6	−19,960.73	2.3792	132	40,185.47	40,895.74	40,476.4	0.797	0.773	0.773	40,209.32
C7	−19,872.23	2.6249	154	40,052.46	40,881.12	40,391.89	0.785	0.718	0.718	40,085.39
C8	−19,779.57	2.225	176	39,911.13	40,858.17	40,299.05	0.823	0.243	0.243	39,954.76
C9	−19,705.95	2.119	198	39,807.9	40,873.31	40,244.31	0.84	0.409	0.410	39,863.95
C10	−19,637.82	2.1967	220	39,715.63	40,899.43	40,200.53	0.814	0.730	0.730	39,785.89

*P values from non-survey adjusted LCA models.
C# indicates the number of classes estimated in the model.
REI, respiratory event index; LL, log likelihood; AIC, Akaike information criterion; BIC, Bayesian information criterion; SSABIC, sample size adjusted BIC; VLMR, Vuong-Lo-Mendell Rubin; LMR, Lo-Mendell-Rubin; AICC, sample corrected Akaike information criterion.

cardiovascular risk and disease profiles of the generated pheno-types. We tested overall differences between groups using survey adjusted chi-squared tests for categorical measures and *t*-tests for continuous measures. Additionally, we calculated and tested pairwise odds ratios to test for potential differences in characteristics between classes. To facilitate understanding of the distribution of the sleep symptoms across the chosen solu-tion, we generated heatmaps to visualize symptoms prevalence.

Sensitivity analysis

We conducted two sets of sensitivity models. First, we refit the LCAs to include only the sleep-related variables. We did so to examine the extent of change in the generated phenotypes as a result of excluding comorbid conditions from the classification process. As with the primary analyses, we characterize the sleep symptoms and sleep health, sociodemographic and general health conditions, and cardiovascular risk and disease profiles of the generated phenotypes.

Second, we validated our results using hierarchical clustering on principal components (HCPC) techniques. To start, multiple correspondence analysis (MCA) was used to generate the prin-cipal components for categorical variables. For missing data under MCA (around 1%), we replaced missing ESS scores with the median score and performed hot-deck imputation [35] using the VIM (version 6.1.0) package in R on missing categorical indi-cators. Details of hot-deck imputation are provided elsewhere [36, 37]. Briefly, this imputation technique sorts individuals by responses and replaces missing values based on the value of the closest matching pair. We used the *FactoMineR* package to per-form the MCA, and scree tests to determine the optimal number of extracted dimensions. Subsequently, we used HCPC with the ward method to determine the ideal number of classes. Results were generated using the *NbClust* package, which calculates up to 23 indices to determine the optimal solution. As with the pri-mary analyses, we generated descriptive tables and a heatmap to characterize the sleep symptoms of the adopted solution through this method. Lastly, we tabulated classification from the primary solution and MCA/HCPC model.

Results

Target population characteristics

The average age was 52.4 ± 13.9 years and 34.0% were female. Overall, 54.9% were hypertensive, and 8.4% had obstructive lung disease. The average respiratory event index (REI3%) was 33.8 ± 22.5. The most-reported sleep symptoms were snoring (76.8%), dozing off while watching television (72.4%), and waking up sev-eral times at night (62.3%).

Model selection

The statistical fit indices generated across the solutions of the full LCA models and the models not accounting for comorbidities are presented in Table 1 and Supplementary Table S3, respectively. Across models, the sample size corrected Akaike and Bayesian information criterion decreased consist-ently as the number of classes increased suggesting a better fit. The descent of both criteria narrowed and stabilized after the

three-class solution. Additionally, the Vuong-Lo-Mendell Rubin *p*-value became insignificant with four classes thus rejecting a better fit relative to the three-class solution. The entropy value for the three-class model was 0.82 suggesting good fit and dec-remented when larger classes were fit. The fit statistics were consistent in identifying the three-class solution as providing optimal fit for the LCA model that excluded comorbid conditions (Supplementary Table S3). The MCA/HCPC models also pointed to a three-class solution as providing the best fit.

Phenotypes characteristics

Based on the fit criteria, we selected the three-class solution as providing optimal fit to the data. Based on our examination of OSA symptoms prevalence within each class, we adopted a naming convention similar to Ye et al. [14] since the OSA pro-files were consistent with their findings: The most common de-rived sleep phenotype was (1) minimally symptomatic (47.7%), (2) disturbed sleep (37.1%), and (3) excessive sleepiness (15.2%). Representation of OSA symptoms for these classes can be seen in Figure 1. We found significant and consistent differences in OSA symptoms across the three derived sleep phenotypes (Table 2). For example, the excessive sleepiness group was more likely to report snoring, dozing while watching TV, or driving compared to the other derived sleep phenotypes. The disturbed sleep phenotype was associated with trouble falling asleep, awaking at night, and trouble going back to sleep. The minim-ally symptomatic phenotype had the lowest prevalence on every OSA symptom.

The minimally symptomatic class was more likely to be male (76%) and had the lowest BMI, REI3%, and REI4% aver-ages relative to the other phenotypes. Individuals in this group performed better (higher) on the mental and physical health measures and had an ESS score of 4.4 ± 3.7. This group was the least acculturated compared to the other classes; it also had the highest Spanish language preference of the three groups. The minimally symptomatic group had the lowest prevalence of OSA symptoms and cardiovascular comorbidities relative to the other phenotypes but had a prevalence of obstructive lung dis-ease that was like other groups (Tables 2–7).

The disturbed sleep group had the highest female compos-ition (53% male). Average BMI, REI3%, and REI4% were slightly above the minimally symptomatic group, but below the exces-sive sleepiness group. Individuals in this group had the highest prevalence of insomnia-related symptoms. For example, indi-viduals in this group had high rates of trouble falling asleep (78%), waking up several times at night (92%), and trouble going back to sleep (65.8%). The percentage of Dominicans and Puerto Ricans was higher compared to other classes. The average ESS score of individuals in this group was 6.0 ± 3.6. In addition to the higher levels of acculturation compared to the other groups, they were also the most likely to have insurance and had the lowest Spanish language preference of the three groups. The disturbed sleep group had higher odds of stroke/ TIA history compared to the minimally symptomatic group (Tables 2–7).

The excessive sleepiness group was composed of 67% males, had the highest prevalence of obstructive lung disease, highest average BMI, REI3%, REI4%. The average ESS score was 15.7 ± 4.6 with 61% of individuals reporting they nap for more than 5 min,

Sleep Derived Phenotype
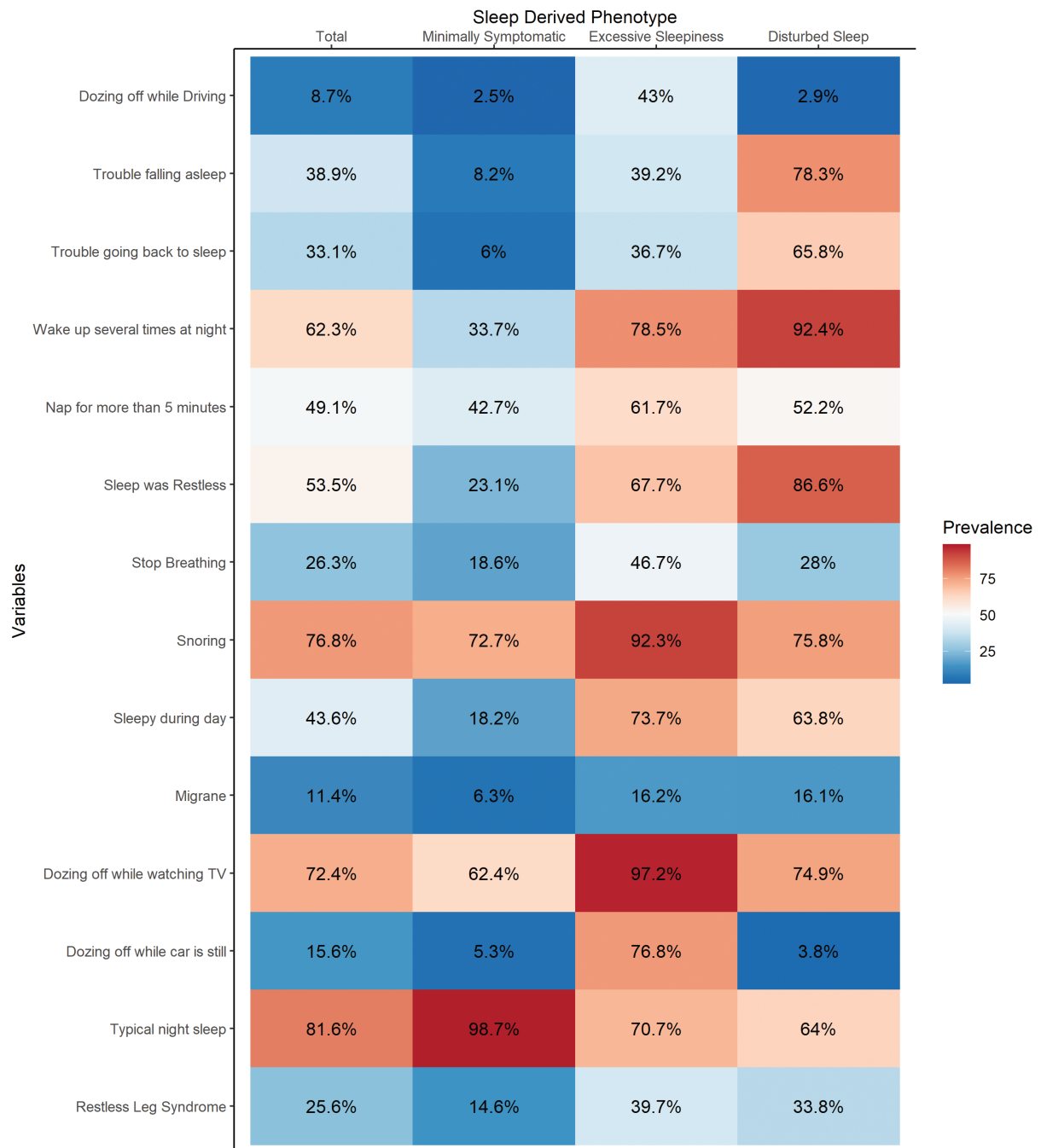
Figure 1. Symptom profile of primary latent class solution with HCHS/SOL individuals REI3% ≥ 15.

All variables are presented using the prevalence in % term.

TV, television.

73.7% feel sleepy during the day, and 97.2% may fall asleep or feel sleepy watching television. Insomnia-related symptoms were also higher for this group compared to the Minimally symptomatic group (Tables 2–7).

Similar trends in summary fit statistics were observed in the LCA solution without comorbidities, thus we also selected three-class solution. The three classes found in this solution with largely similar results to those with comorbidities (Supplementary Tables S4–S6).

Validation with MCA/HCPC

The three-class solution was validated by the MCA/HCPC models with 11 indices (e.g. silhouette) pointing to optimal fit and evidence for the highest relative loss of inertia through this solution. Cross-validation using different variables (e.g. no comorbidities) as well as different clustering techniques (hierarchical clustering, LCA) showed consistency of classes across models. The same three phenotypes with similar prevalence were extracted; however, we found differences in

Table 2. Sleep symptoms (prevalence or mean) by derived sleep phenotypes from LCA solution, HCHS/SOL (2008–2011)

	Total (n = 1,605)	Minimally symptomatic (n = 780)	Excessive sleepiness (n = 248)	Disturbed sleep (n = 577)	P
Sleep symptoms included in LCA†					
Dozing off while driving	8.7	2.5	43.0	2.9	0.001
Trouble falling asleep	38.9	8.2	39.2	78.3	0.001
Trouble going back to sleep	33.1	6.0	36.7	65.8	P < 0.001
Wake up several times at night	62.3	33.7	78.5	92.4	P < 0.001
Nap for more than 5 min	49.1	42.7	61.7	52.2	P < 0.001
Restless sleep	53.5	23.1	67.7	86.6	P < 0.001
Stop breathing	26.3	18.6	46.7	28.0	P < 0.001
Snoring	76.8	72.7	92.3	75.8	P < 0.001
Sleepy during day	43.6	18.2	73.7	63.8	P < 0.001
Migraine	11.4	6.3	16.2	16.1	P < 0.001
Dozing off while watching TV	72.4	62.4	97.2	74.9	P < 0.001
Dozing off while car is still	15.6	5.3	76.8	3.8	P < 0.001
Sound or restful sleep	81.6	98.7	70.7	64.0	P < 0.001
Restless legs	25.6	14.6	39.7	33.8	P < 0.001
ESS score*	6.73 (5.65)	4.39 (3.67)	15.71 (4.53)	6.03 (3.56)	P < 0.001
Sleep variables not included in LCA*					
REI3%	33.76 (22.53)	31.40 (18.19)	42.50 (30.28)	33.40 (23.02)	P < 0.001
REI4%	26.42 (21.95)	24.16 (17.62)	36.05 (30.23)	25.51 (21.93)	P < 0.001
Minimum oxygen saturation (%)	77.64 (8.43)	77.84 (8.09)	75.42 (9.94)	78.24 (8.07)	0.006

*Means and standard deviations are presented.
†% are presented.
P value: Pearson's chi square test for continuous variables; Regression-based F test for categorical variables.
ESS, Epworth Sleepiness Scale; LCA, latent class analysis; REI, respiratory event index.

Table 3. Baseline sociodemographic and health characteristics by primary solution of three derived sleep phenotypes, HCHS/SOL (2008–2011)

	Total	Minimally symptomatic	Excessive sleepiness	Disturbed sleep	P
Unweighted n	1,605	780	248	577	
% of total	100.0	47.7	15.2	37.1	
Males†	934 (66.0)	520 (75.5)	150 (67.3)	264 (53.3)	P < 0.001
Background†					
Central American	154 (6.2)	77 (6.2)	31 (9.3)	46 (5.1)	P < 0.001
Cuban	281 (26.9)	145 (31.1)	42 (26.3)	94 (21.8)	
Dominican	117 (8.5)	49 (6.8)	15 (5.2)	53 (12.1)	
Mexican	632 (34.9)	326 (36.0)	88 (34.8)	218 (33.4)	
Puerto Rican	287 (16.7)	104 (11.4)	55 (19.6)	128 (22.4)	
South American	90 (3.8)	52 (4.5)	12 (2.9)	26 (3.4)	
Other	42 (2.9)	26 (4.1)	5 (1.9)	11 (1.8)	
Language preference†					
Spanish	1,361 (81.8)	682 (86.2)	202 (80.0)	477 (76.7)	P = 0.017
English	244 (18.2)	98 (13.8)	46 (20.0)	100 (23.3)	
Age years*	52.41 (13.93)	52.13 (13.98)	52.48 (13.19)	52.95 (14.10)	P = 0.812
SF-12 physical component score*	46.75 (11.13)	49.33 (9.37)	44.33 (12.79)	44.57 (11.56)	P < 0.001
SF-12 mental component score*	49.83 (12.11)	53.40 (10.87)	47.93 (12.19)	46.00 (12.07)	P < 0.001
SASH acculturation language*	1.92 (1.14)	1.78 (1.04)	2.03 (1.20)	2.03 (1.21)	P = 0.007
SASH acculturation social*	2.19 (0.62)	2.16 (0.59)	2.19 (0.60)	2.22 (0.65)	P = 0.460
Insured†	791 (57.5)	349 (53.3)	111 (53.0)	331 (64.8)	P = 0.006

*Means and standard deviations are presented.
†Counts and % are presented.
P value: Pearson's chi square test for continuous variables; Regression-based F test for categorical variables.
SF-12, 12-Item short-form survey.

comorbidities prevalence through this extraction method. For a visual representation of these classes, see Supplementary Figure S1. While there were some differences between these two solutions, the OSA symptoms, as well as sociodemographics of each group, were largely equivalent (Tables 2 and Supplementary Tables S7–S9).

We tested the consistency of classification between the LCA and MCA/HCPC models. The minimally symptomatic group was the most consistent; 96% of individuals were classified as minimally symptomatic under both methods. The disturbed sleep phenotype was also stable at 83%. The excessive sleepiness was the least stable, with only a 67% overlap. These results were

Table 4. Baseline cardiovascular characteristics by primary solution derived sleep phenotype

	Total	Minimally symptomatic	Excessive sleepiness	Disturbed sleep	P
Major comorbidities included in LCA†					
Hypertension	54.9	52.6	57.4	56.8	P = 0.450
Diabetes	34.1	30.4	37.7	37.5	P = 0.119
Framingham CVD	37.6	27.7	41.6	48.8	P < 0.001
Obstructive lung disease	8.4	9.2	12.7	5.6	P = 0.036
Major comorbidities not included in LCA					
HDL* (mg/dL)	44.38 (11.56)	44.76 (12.18)	43.93 (11.34)	44.17 (10.83)	P = 0.648
Total cholesterol* (mg/dL)	206.21 (46.36)	206.14 (44.82)	203.87 (45.39)	207.46 (48.98)	P = 0.758
Triglycerides* (mg/dL)	173.00 (146.37)	166.51 (115.59)	165.96 (94.23)	184.23 (191.10)	P = 0.272
BMI* (kg/m^2)	33.81 (6.83)	32.97 (6.43)	35.04 (7.70)	34.21 (6.66)	P = 0.003
Current smoker†	18.9	19.1	18.2	18.8	P = 0.974
Alcohol use†	47.4	49.8	46.8	44.5	P = 0.387
CHD†	7.7	6.3	11.5	7.9	P = 0.167
Heart failure†	3.1	3.1	3.5	2.8	P = 0.899
Stroke/TIA†	4.4	3.0	3.1	6.6	P = 0.040

*Means and standard deviations are presented.
†% are presented.
P value: Pearson's chi square test for continuous variables; Regression-based F test for categorical variables.
HDL, high-density lipoproteins; BMI, body mass index; CHD, coronary heart disease; TIA, transient ischemic attack; LCA, latent class analysis; CVD, cardiovascular disease. This is a 0/1 indicator variable that defines a composite CVD definition based on the Framingham Study criterion (http://www.framinghamheartstudy.org/risk/gencardio.html).

Table 5. Odds ratio of derived sleep phenotypes by sleep symptoms

	Disturbed sleep (ref) vs excessive sleepiness		Disturbed sleep (ref) vs minimally symptomatic		Minimally symptomatic (ref) vs excessive sleepiness	
	Odds ratio	P	Odds ratio	P	Odds ratio	P
Sleep symptoms included in LCA†						
Dozing off while driving	25.074	0.001	0.845	P = 0.691	29.690	P < 0.001
Trouble falling asleep	0.179	P < 0.001	0.025	P < 0.001	7.256	P < 0.001
Trouble going back to sleep	0.302	P < 0.001	0.033	P<0.001	9.148	P < 0.001
Wake up several times at night	0.300	P < 0.001	0.042	P < 0.001	7.187	P < 0.001
Nap for more than 5 minutes	1.476	P = 0.083	0.685	P = 0.018	2.156	P < 0.001
Sleep was restless	0.322	P < 0.001	0.046	P < 0.001	6.960	P < 0.001
Stop breathing	2.255	P < 0.001	0.587	P = 0.004	3.840	P < 0.001
Snoring	3.830	P < 0.001	0.849	P = 0.377	4.154	P < 0.001
sleepy during day	1.592	P = 0.046	0.126	P < 0.001	12.585	P < 0.001
Migraine	1.011	P = 0.969	0.352	P < 0.001	2.872	P < 0.001
Dozing off while watching television	11.666	P < 0.001	0.556	P = 0.001	20.963	P < 0.001
Dozing off while car is still	84.280	P < 0.001	1.424	P = 0.349	59.188	P < 0.001
Sound or restful sleep	1.361	P = 0.153	42.367	P < 0.001	0.032	P < 0.001
Restless legs	1.288	P = 0.238	0.333	P < 0.001	3.863	P < 0.001

†% are presented.
Odds ratios were calculated using survey weighted logic regression models.
LCA, latent class analysis.

consistent in the models without comorbidities (Supplementary Table S10). Lastly, we tested whether recoding "don't know" answers could affect models by recoding them to missing. While the subpopulation was smaller, cross-tabulation with the original LCA model shows that results are consistent with the original model (Supplementary Table S11).

Discussion

Using three clustering algorithms and covariates, we showed evidence for three sleep phenotypes: (1) minimally symptomatic, (2) disturbed sleep, and (3) excessive sleepiness among diverse Hispanics/Latinos. We also found noteworthy differences in comorbidity profiles across the three phenotypes. The excessive sleepiness group had an increased prevalence of obstructive lung disease, while the disturbed sleep group had higher odds of self-reported cerebrovascular disease and/or TIA. Hispanics/Latinos overall had higher cardiovascular risk (diabetes and hypertension) and cardiovascular disease compared to the populations represented in the ISAC and SAGIC cohorts [14, 17], and our results underscore the importance of considering these factors [38–40] in the context of sleep in this population. The prevalence of obstructive lung disease in our cohort was lower compared to the ISAC cohort (8.4% HCHS/SOL vs. 18.7% ISAC).

Our study is partly consistent with previous research of predominantly non-Hispanic White and Korean samples [14,

Table 6. Mean comparisons across sleep health and cardiovascular risk indicators for primary LCA solution

	Minimally symptomatic	Excessive sleepiness	Disturbed sleep
	β [95% CI]	β [95% CI]	β [95% CI]
Age years	52.13 [50.77;53.49]	52.48 [50.16;54.80]	52.95 [50.89;55.01]
REI3%	31.40 [29.89;32.92][B]	42.50 [37.88;47.11][C]	33.40 [30.73;36.07]
REI4 %	24.16 [22.67;25.64][B]	36.05 [31.45;40.64][C]	25.51 [23.12;27.90]
Minimum oxygen saturation	77.84 [77.20;78.48][B]	75.42 [73.93;76.92][C]	78.24 [77.32;79.16]
SF-12 physical component score	49.33 [48.50;50.16][B,C]	44.33 [42.17;46.49]	44.57 [43.28;45.87]
SF-12 mental component score	53.40 [52.46;54.34][B,C]	47.93 [46.08;49.78]	46.00 [44.37;47.63]
HDL (mg/dL)	44.76 [43.67;45.84]	43.93 [42.24;45.62]	44.17 [42.97;45.37]
Total cholesterol (mg/dL)	206.14 [202.25;210.02]	203.87 [196.62;211.12]	207.46 [201.99;212.93]
Triglycerides (mg/dL)	166.51 [157.11;175.91]	165.96 [152.67;179.25]	184.23 [164.07;204.38]
BMI (kg/m²)	32.97 [32.43;33.51][B,C]	35.04 [33.80;36.27]	34.21 [33.31;35.10]
Language acculturation	1.78 [1.68;1.88][B,C]	2.03 [1.84;2.22]	2.03 [1.87;2.19]
Social acculturation	2.16 [2.10;2.21]	2.19 [2.10;2.28]	2.22 [2.13;2.31]

Group differences testing for continuous variables calculated through survey adjusted linear regression of the clustering variable on latent class membership.
B: Group differences significant at *P* < 0.05 relative to the daytime sleepiness.
C: Group differences significant at *P* < 0.05 relative to the disturbed sleep.
LCA, latent class analysis; REI, respiratory event index; HDL, high-density lipoproteins; BMI, body mass index; SF-12, 12-Item short-form survey.

Table 7. Odds ratio comparisons across cardiovascular risk indicators for primary LCA solution

	Disturbed sleep (ref) vs excessive sleepiness		Disturbed sleep (ref) vs minimally symptomatic		Minimally symptomatic (ref) vs excessive sleepiness	
	OR	P	OR	P	OR	P
Included in the LCA model						
Hypertension	1.027	P = 0.900	0.844	P = 0.297	1.217	P = 0.318
Diabetes	1.008	P = 0.972	0.729	P = 0.073	1.383	P = 0.119
Framingham CVD	0.748	P = 0.175	**0.401**	P < 0.001	**1.862**	P = 0.002
Obstructive lung disease	**2.436**	P = 0.015	1.696	P = 0.058	1.436	P = 0.259
Not included in the LCA model						
Male	**1.804**	P = 0.008	**2.706**	P < 0.001	**0.667**	P = 0.050
Cigarettes	0.963	P = 0.879	1.018	P = 0.923	0.945	P = 0.816
Alcohol	1.100	P = 0.658	1.238	P = 0.192	0.889	P = 0.547
CHD	1.515	P = 0.193	0.788	P = 0.406	1.923	P = 0.065
Heart failure	1.276	P = 0.635	1.125	P = 0.778	1.135	P = 0.817
Stroke/TIA	0.462	P = 0.105	**0.444**	P = 0.032	1.042	P = 0.929

Odds ratios were calculated using survey weighted logic regression models.
OR, odds ratios; LCA, latent class analysis; CHD, coronary heart disease; TIA, transient ischemic attack; CVD, cardiovascular disease.

16, 17]. We provide added evidence on the stability of OSA symptom classes across populations, and our findings help generalize symptom heterogeneity to diverse community-dwelling Hispanics/Latinos. Our study is unique in that it points to differences in the prevalence of symptoms classes across populations. Our representative target populations had a higher prevalence of minimally symptomatic individuals compared to the ISAC and SAGIC samples and a smaller excessive sleepiness group compared to ISAC, SAGIC, and Korean samples. Furthermore, the Central/South Americans in SAGIC (6.6% of cohort) were less prevalent in the disturbed sleep group, but we did not see the same relationship in our cohort. These results have implications for targeted treatment strategies in Hispanics/Latinos. For example, some evidence [15] suggests that OSA patients in the excessive sleepiness group have better positive airway pressure (PAP) adherence compared to the other two groups. Future work should examine the implications of different treatment modalities across sleep-derived phenotypes among Hispanics/Latinos.

We posit several explanations for differences in prevalence by cohorts. First, our findings are based on community-dwelling

adults, whereas ISAC and SAGIC focused on clinical samples, suggesting a referral bias in the latter studies. Second, despite consistent selection on moderate and severe apnea (REI3% ≥ 15), our target population is relatively young and had a lower OSA symptom load, most likely a consequence of our study design. Indeed, a recent study using a population-based cohort, found that sleep measures did not significantly differ between OSA and non-OSA individuals [41]. Third, the sociodemographic, cultural, and health profiles of our target population are distinct from those of non-Hispanic whites and Koreans. Our findings point to distinct socio-demographic (e.g. sex and Hispanic/Latino background) and cardiovascular and pulmonary profiles that could guide research and clinical work on Hispanics/Latinos as well as the non-Latino population. Socioeconomic characteristics [42, 43], cultural practices (e.g. acculturation [44, 45]), and health conditions (e.g. BMI [46]) influence sleep health, sleep management, and clinical management and treatment of symptoms. Kim et al.'s [16] community-dwelling Korean population-based study also found a high prevalence of minimally symptomatic patients (55.7%) in a healthier population. However, they reported a much lower prevalence of disturbed

sleep relative to our cohort (38.1% in our cohort vs. 14.5%). The Sleep Heart Health study [18], a population-based study, reported around 12% disturbed sleep, but the Canadian Study [19] reported around 30% disturbed sleep and found another cluster with disturbed sleep symptoms. We found a disproportionate representation of Puerto Ricans, Dominicans, and women in the disturbed sleep group compared to the minimally symptomatic and excessive sleepiness groups. We posit the following explanations for these differences: Previous studies on HCHS/SOL cohort have found female, but not male, Puerto Ricans had the highest risk for three or more cardiovascular risk factors [40]. The majority of the Dominican and Puerto Ricans in our cohort came from the field center in the Bronx and thus may experience more environmental noise from living in a dense urban environment which could have affected sleep [47]. Results from HCHS/SOL have shown Dominicans and Puerto Ricans had the highest depressive symptoms of all groups [48], and sleep disturbances are common in individuals with depression [49, 50]. Similar to other studies [14, 16, 17], we found that despite greater numbers of men to women in each group, women were overrepresented in the disturbed sleep group, underscoring the importance of characterizing individuals into OSA subgroups. Women usually report more insomnia-like symptoms compared to men [51]. Women also have shorter apnea/hypopnea events [52] and greater reporting of somatic symptoms among women compared to men [53].

Our findings suggest that, despite evidence for moderate to severe OSA (REI3% ≥ 15), a large portion of individuals meeting clinical OSA criteria could go undiagnosed and untreated due to minimum symptoms/complaints present. Even compared to other population-based studies like the Sleep Heart Health Study [18] and the Canadian Study [19], there was an overrepresentation of minimally symptomatic individuals in our cohort. On the one hand, fewer OSA symptoms are potential markers of a better quality of life. Excessive sleepiness and insomnia have been linked with decrements in health quality of life [54], while many studies have shown no relation between AHI and quality of life [54–56]. On the other hand, fewer symptoms presentation could lead to longer periods of undiagnosed disorders and portend more severe downstream development of adverse health risks (e.g. hypertension [57]) and negative health outcomes. Underdiagnosis of OSA can lead to substantial adverse downstream health outcomes including cognitive impairment and dementia [58]. OSA may impact overall health through amplification of oxidative stress, endothelial dysfunction, and increased inflammation [59, 60].

Our data suggest that minimally symptomatic Hispanics/Latinos may face unique barriers to achieving diagnosis and treatment. For example, over half of the individuals in the minimally symptomatic group did not have health insurance (53%), had the lowest language acculturation of the three groups, and highest Spanish language preference. Spanish language preference is associated with lower rates of insurance and access health care [61], and this is in addition to lower access to healthcare among US Hispanics/Latinos, even after adjusting for income and health insurance [62]. Overall, these factors may present as barriers for both diagnosis and treatment of OSA. US nativity could also influence OSA classification. US-born Mexican Americans have higher rates of short sleep (both <7 and <6 h) and report more insomnia and excessive sleepiness symptoms

compared to immigrants (also known as the "Hispanic Paradox") [10]. Future studies should account for acculturation and other sociocultural factors (years in the US, health literacy, nativity, etc.). Prospective studies are needed to determine if derived sleep phenotypes have a differential incidence of health outcomes/chronic illness.

Strengths and limitations

To our knowledge, this is the first study that provides estimates for empirical classifications of OSA symptoms among diverse community-dwelling Hispanics/Latino adults in the US. Our study benefits from the design of the HCHS/SOL cohort, the large sample size relative to existing studies, and the potential generalizability to a wider range of the OSA spectrum compared to clinical samples. Our findings provide additional evidence for the validity and the generalizability of classes currently reported in the literature.

Our study has some limitations. First, our selected symptom items did not exactly match the ones reported in previous studies. However, the criteria in this study include largely similar array of OSA symptom comparable to previously published work [14, 16, 17]. This potentially limits comparisons of specific derived sleep phenotypes and their correlates across studies. Second, community-based population studies may have confounding conditions that lead to systematic errors (e.g. through differential measurement). HCHS/SOL minimized the influence of these confounders by creating strict regulations and guidelines across testing centers with centrally trained bilingual staff and technicians administering the tests in the individual's preferred language. As with other studies, using subjective data can also introduce classification error which can have implications for interpretation in clinical settings. Even though we provided cardiovascular information for each cluster, given the scope of the paper, we did not perform any multivariable modeling to link our extracted phenotypes to the prevalence and incidence of cardiovascular disease and risk factors. Furthermore, the adherence in classification between LCA and MCA models was mixed for the excessive sleepiness group. Specifically, we found that 33% of individuals were classified out of the excessive sleepiness group by the MCA algorithm and classified as minimally or disturbed sleep. However, this reclassification was primarily driven by the MCA algorithm's selection relative to the "sleepy while driving" item. Otherwise, our findings show that the symptoms were consistent with what the LCA classified as excessive sleepiness. Further explorations are required to validate which of the two classification methods is more precise. Finally, given the scope of the study, we did not test for invariance in phenotype composition across socio-demographically interesting groups. Future studies should explore whether the LCA classifications generated through this work are invariant to language. This can potentially provide supportive evidence that language differences in questionnaire compositions (e.g. through translation) did not lead to differential classifications. We believe that the differences in phenotype characteristics published in this work provide a starting point for future work focused on testing the stability of phenotypes across demographic and other health characteristics.

Conclusion

We reported three OSA phenotypes among diverse Hispanics/Latinos; a large US population that has been understudied in sleep research. These derived phenotypes were comparable to previous studies of non-Latinos. However, we found notable differences in the prevalence of these classes relative to non-Hispanic white, suggesting that other biopsychosocial and lifestyle factors such as diet, environment, and physical activity may contribute to derived sleep phenotypes among Hispanics/Latinos. Our reported empirically derived OSA phenotypes in Hispanics/Latinos may guide research focused on health outcomes (e.g. dementia) and could inform clinical diagnoses, development of targeted therapeutics, and allocation of public health resources. There is value in relating these symptom-based classes to physiological endotypes (arousal threshold, neuromuscular collapsibility, loop gain, etc.) which would provide specific therapeutic targets as well as risk prediction.

Supplementary Material

Supplementary material is available at *SLEEP* online.

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