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Reference values for high attenuation areas on chest CT in a healthy, never-smoker, multi-ethnic sample: The MESA study

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Abstract

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Supplementary Information

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Data availability statement: Anonymized data from the MESA study have been made publicly available at BioLincc (https://biolincc.nhlbi.nih.gov/home/) and/or dbGAP (https://www.ncbi.nlm.nih.gov/gap/).

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Background and objective: Normative values for HAA—a quantitative, CT-based measure of subclinical ILD—in healthy adults are needed to improve interpretability in clinical and research settings.

Methods: HAA was measured on full-lung CT in 3110 participants in the MESA study. Clinical prediction models were developed using a healthy never-smoker subset with normal spirometry (n = 696). RMSE on cross-validation was used as the primary criterion for model selection. Parametric and non-parametric methods were considered. z-Scores were calculated for the entire study sample. Associations between z-scores and several ILD features were estimated.

Results: In the healthy never-smoker subset, the mean age was 69 years with a range of 54–93 years. The median HAA was 4.3% with a range of 2.7–17.8%. Linear regression had better predictive performance than other methods. The final model included race, height, weight, age and sex. The standard error of the estimate was 1.62 with a cross-validated RMSE of 1.64 and an adjusted R² of 0.139. z-Scores were associated with several ILD outcomes in adjusted models, including ILA (OR: 1.40 per z-unit; 95% CI: 1.30, 1.52), exertional dyspnoea (OR: 1.08 per z-unit; 95% CI: 1.02, 1.15) and FVC (expected increase per z-unit: –2.49; 95% CI: –2.95, – 2.03).

Conclusion: We present a reference equation and z-scores to define expected values of HAA on full-lung CT to aid HAA interpretation in middle-aged and older adults.

Keywords

high attenuation area; pulmonary fibrosis; quantitative computed tomography; reference equations; subclinical interstitial lung disease

INTRODUCTION

Interstitial lung disease (ILD) refers to a family of closely related respiratory disorders that cause progressive fibrosis and inflammation in the lung.¹ Idiopathic pulmonary fibrosis (IPF) is the most common ILD. It affects nearly 1 in 200 older U.S. adults over the age of 65 years and carries a poor prognosis.^{2,3} Two available therapies have been shown to slow disease progression in patients with mild to moderate IPF^{4,5}; however, effective management depends on early disease detection.⁶ There are no existing interventions that prevent, halt or reverse the development of fibrotic ILD.⁷ Improved methods of identifying early stage ILD could allow for earlier intervention and may help reduce the public health burden of the disease.

High attenuation areas (HAA) are a quantitative, computed tomography (CT)-based measure of subclinical ILD.^{8–12} HAA is associated with clinical respiratory outcomes, including ILD-specific hospitalization and death in community-dwelling adults.^{9,10} These data support HAA as a biomarker of the earliest biological changes in the lung parenchyma leading to ILD. A better understanding of the distribution of HAA in a healthy sample is needed to improve interpretability in clinical and research settings.

In contrast to pulmonary function testing and quantitative measures of emphysema on CT,¹³ a normative range for HAA has not been established. Here, we examine the natural variation of HAA in a healthy never-smoker sample of community-dwelling adults. We present an

age, sex and body size-specific prediction model that defines expected values of HAA and an upper limit of normal (ULN) on full-lung CT. We further demonstrate that adjusted zscores generated by this model are associated with visually identifiable interstitial lung abnormalities (ILA) antecedent to clinical ILD and with other ILD features and outcomes.

METHODS

Study design and participants

The Multi-Ethnic Study of Atherosclerosis (MESA) is a National Heart, Lung, Blood Institute (NHLBI) funded prospective cohort study of 6814 community-dwelling adults without cardiovascular disease sampled from six communities across the United States between 2000 and 2002.¹⁴ Of these, 3113 underwent full-lung CT imaging at the 10-year follow-up during 2010–2012. Three participants were excluded due to inadequate data. The present study is a cross-sectional analysis of 3110 MESA participants. All MESA participants provided informed consent, and the MESA study was approved by the institutional review boards at all centres.

The healthy never-smoker sample

In accordance with the NHANES III sampling criteria used to develop the spirometry reference equations¹⁵ and validated in the MESA cohort,¹⁶ healthy never-smokers were defined as all participants who denied all of the following: ever-smoking cigarettes, cigars or pipes; physician diagnosis of asthma, emphysema or lung cancer; and respiratory symptoms including chronic cough, chronic phlegm production, exertional dyspnoea and wheezing in chest. Further exclusions included abnormal spirometry, estimated glomerular filtration rate <30 mL/min/m² (eGFR), obesity (body mass index, BMI 35 kg/m²), self-report of bronchitis or pneumonia in the 2 weeks prior to CT scan, and inadequate data for analysis (Fig. 1). We chose to include overweight participants (BMI between 25.0 and 34.9 kg/m²) in the healthy never-smoker subset to make this cohort more representative of the general population in the United States where only one-third of adults over the age of 20 years have a BMI less than 25.¹⁷ In total, 696 participants were included in the healthy never-smoker sample.

Respiratory symptoms

Cough, phlegm production and wheezing were assessed at 10-year follow-up during the years 2010–2012 with the following questions administered verbally: 'Do you usually have a cough on most days for three or more months during the year?'; 'Do you usually bring up phlegm from your chest on most days for three or more months during the year?'; and 'In the last 12 months, have you had wheezing or whistling in your chest?'. Exertional dyspnoea was defined as affirmative response to any of the following questions administered verbally: 'When walking on level ground, do you get more breathless than people your own age?'; 'When walking up hills or stairs, do you get more breathless than people your own age?'; and 'Do you ever have to stop walking because of breathlessness?'.

Lung function

Lung function was assessed by spirometry in accordance with the American Thoracic Society (ATS)/European Respiratory Society guidelines as previously described.^{16,18}

HAA and interstitial lung abnormalities

Full-lung CT scans were obtained as previously described.¹⁶ Image attenuation was measured using a modified version of the Pulmonary Analysis Software Suite (University of Iowa, Iowa City, Iowa, USA), and HAA were defined as the percent of imaged lung volume having CT attenuation between –600 and –250 Hounsfield Units (HU), as previously described.^{8–10,12,19}

ILA on CT scans were defined as involvement of more than 5% of non-dependent lung by reticular abnormalities, ground-glass abnormalities, diffuse centrilobular nodularity, honeycombing, traction bronchiectasis and/or non-emphysematous cysts.^{9,20,21}

Statistical analysis

We developed prediction models for full-lung HAA at 10-year follow-up using a healthy never-smoker sample. To optimize predictive performance, parametric and non-parametric methods were considered, and root of mean square error (RMSE) on 10-fold crossvalidation was used as the primary criterion for model selection. Support vector machine (SVM) regression with a linear kernel, random forests, boosting, ordinary least squares (OLS) regression and elastic net regression were considered. Cross-validation to specify tuning parameters was nested inside of cross-validation to estimate model performance, and tuning parameter specification was performed on training sets as appropriate to the method.

To preserve comparability across models, cross-validation folds were defined prior to all analyses and the same folds were used for all models fitting (with the exception of tuning parameter specification).

Variable selection was performed by best subset selection for all methods. Candidate variables included race (as an indicator variable), waist circumference, hip circumference, height, weight, BMI, age, sex and a binary variable indicating if the subject received a low dose of radiation on CT scan due to having BMI below a certain threshold.

The presence of non-linear relationships and variable interactions were assessed on OLS models. For each continuous predictor, several non-linear terms, including a smoothing spline and polynomial terms of degree 2 through 6, were examined in the best subset setting. All possible subsets of all possible pairwise interaction terms were examined. Evidence of non-linearity or variable interactions was defined by reduction in RMSE of 0.1 when compared with the simplest model. Race/ethnicity- and sex-stratified OLS models were developed using the fitting process described above.

Predicted values of HAA were calculated for each study participant using the selected prediction model, and z-scores were calculated for each individual as described below:

 $z-Score_i = \frac{O_i - E_i}{SEE}$

where O_i is the observed value of HAA for subject *i*, E_i is the expected value of HAA for subject *i* and SEE is the standard error of the estimate calculated in the healthy never-smoker subset. The reader should note that SEE is equivalent to the SD of the errors of prediction. Here, we refer to the SD of the errors of prediction as the SEE when it is estimated using the entire study sample and as the RMSE when estimated as a mean over cross-validation samples. We define elevated HAA as the upper fifth percentile of the healthy normal distribution. As the distribution of z-scores is right-skewed, we use the empirical 95th quantile (1.634) as the cut-off value for elevated HAA instead of the 95th quantile of a standard normal distribution.

ULN were defined by one-tailed 95% prediction intervals as follows

 $ULN_i = O_i + 1.634 \times SEE$

where ULN_{*i*} is the ULN for subject *i*, and O_i and SEE are defined as described above. Elevated HAA is defined as O_i ULN_{*i*} or, equivalently, z-score_{*i*} 1.634.

Similar methods were used to develop a reference equation, predicted values, ULN and zscores for HAA measured from cardiac CT scans (Appendix S1, Table S1 in Supplementary Information).

z-Score associations with ILA, exertional dyspnoea and cough were estimated using logistic regression. z-Score associations with lung function measures and log-transformed pack-years were estimated using OLS regression. Associations were adjusted for study site, smoking status, pack-years, waist circumference, eGFR and educational attainment. The reader should note that adjusted z-scores function as a dimensionality reduction variable much like a principal component generated using principal component analysis. This means that the z-score contains *some* of the information present in the variables used to fit the prediction model, but not *all* of that information. Therefore, effect estimates were also examined in a model adjusted for all of the previously mentioned variables in addition to the predictor variables used to calculate the z-score (race, height, weight, age and sex). This model allows the reader to assess how effectively the dimensionality reduction variable (the z-score) is controlling for the confounding of the predictor variables used in the z-score.

The validity of all models used for inference was assessed by visual inspection of standard diagnostic plots and/or Hosmer–Lemeshow tests, as appropriate.

RESULTS

There were 3110 MESA participants with available data who underwent full-lung CT imaging. 52% of participants were women, 27% were Black, 39% were white, 21% were Hispanic and 13% were Chinese. About half (46%) of the participants were never-smokers,

47% were former smokers and 8% were current smokers. The mean \pm SD age was 69 \pm 6 years, the mean \pm SD weight was 70.3 \pm 13.8 kg and the mean \pm SD height was 165.4 \pm 9.9 cm. Mean \pm SD percent of predicted forced vital capacity (FVC) was 97.1 \pm 17.8%, mean \pm SD percent of predicted forced expiratory volume in 1 s (FEV₁) was 94.9 \pm 19.9% and mean \pm SD FEV₁/FVC ratio was 74.0 \pm 9.0% (Table 1).

Of these, 696 met criteria for inclusion in the healthy never-smoker subset (Fig. 1). The healthy never-smoker subset had a higher proportion of women (61.2%); a higher proportion of Chinese (26.1%); and lower proportions of Blacks (23.1%), whites (32.8%), and Hispanics (26.1%) when compared with the complete study sample. In the healthy never-smoker subset, the mean \pm SD age (69 \pm 9 years) and height (163.4 \pm 9.4 cm) were similar to that of the complete study sample, while the mean \pm SD weight (70.3 \pm 13.8 kg) was lower in the healthy never-smoker subset. The healthy never-smoker subset also had better lung function (mean \pm SD: FVC, 102.1 \pm 14.7; FEV₁, 103.4 \pm 15.1; FEV₁/FVC, 76.8 \pm 6.0) when compared with the complete study sample (Table 1).

The distribution of HAA was right-skewed in both the healthy never-smoker subset and in the complete study sample (Fig. 2A). The mean \pm SD value of HAA was 5.01 \pm 2.38% in the complete study sample and was lower (4.76 \pm 1.76%) in the healthy never-smoker subset (Table 1).

Prediction model for HAA using the healthy never-smoker subset

Linear regression with OLS estimation had improved predictive performance when compared with linear regression with an elastic net penalty implying good model stability despite covariate collinearity. OLS models also outperformed non-parametric methods, including random forests, boosting and support vector regression, suggesting good model specification with respect to parametric assumptions (Table 2).

The best-performing OLS model, given below, had an SEE of 1.62, an RMSE of 1.64 on 10-fold cross-validation and an adjusted coefficient of determination (adj. R²) of 13.9%:

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\begin{array}{l} {\rm HAA}_{\rm pred} = 10.953841 + 0.695687 \times I(\mbox{ race= Chinese }) \\ + 0.410185 \times I(\mbox{ race= Black }) + 0.964014 \\ \times I(\mbox{ race= Hispanic }) - 0.065887 \times \mbox{ height} \\ + 0.036438 \times \mbox{ weight } + 0.022561 \times \mbox{ age} \\ + 0.043759 \times I(\mbox{ sex= male }) \end{array}
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where HAA_{pred} is the predicted value of HAA, height is measured in cm, weight is measured in kg and age is measured in years. The distribution of predicted values of HAA was normally distributed and symmetric (Fig. 2B).

The weighted average of the RMSE of the best-performing gender-stratified models was 1.65, suggesting no benefit to gender stratification. The weighted average RMSE for the race/ethnicity-stratified models was 1.56. This reduction in RMSE was not considered substantial enough to justify the increased complexity of stratified prediction models. No evidence of non-linearity or variable interactions was observed in stratified or unstratified models.

HAA z-score distribution

The distribution of z-scores in healthy never-smoker sample (n = 696) was right skewed (skewness: 2.95) with a median of -0.23. The interquartile range was from -0.57 to 0.26 and the range was from -1.44 to 7.45. By definition, 4.8% had elevated HAA (Fig. 2C).

The distribution of z-scores in the complete study sample (n = 3110) had a heavier right tail (skewness: 5.02) when compared with the distribution of z-scores in the healthy neversmoker subset. In the complete study population, the median z-score was -0.21, the mean was 0.07, the SD was 1.39, the interquartile range was from -0.58 to 0.29 and the range was from -2.1 to 23.42. The proportion of those with elevated HAA was higher in the complete study sample (6.2%) when compared with the healthy never-smoker subset.

HAA z-scores and ILA

Among those with elevated HAA (n = 127), 38.6% had ILA compared to 11.0% among those with normal HAA (n = 2279). Associations between z-scores and ILA were observed in the unadjusted model (OR: 1.39 per z-unit; 95% CI: 1.29, 1.51), the minimally adjusted model (OR: 1.39 per z-unit; 95% CI: 1.29, 1.51) and the fully adjusted model (OR per z-unit: 1.40; 95% CI: 1.30, 1.52). Similar results were observed for the association between (binary) elevated HAA and ILA (Table 3). These results are consistent with our previously published work.⁹

The 95th percentile threshold for elevated HAA (z-score = 1.634) was associated with a sensitivity of 0.97 for the detection of ILA and a specificity of 0.06. Alternative thresholds for elevated HAA were examined using receiver operating curve (ROC) analysis. The area under the curve (AUC) was 0.67 (95% CI: 0.64, 0.70). The optimal threshold for ILA detection identified by Youden's J-statistic was -0.02 (sensitivity: 0.61; specificity: 0.66; Fig. 2D).

HAA z-scores and clinical features

Higher HAA z-score was associated with lower FVC in the unadjusted model (mean difference per z-unit: -2.61; 95% CI: -3.10, -2.13), the minimally adjusted model (mean difference per z-unit: -2.49; 95% CI: -2.96, -2.03) and the fully adjusted model (mean difference per z-unit: -2.49; 95% CI: -2.95, -2.03). Higher HAA z-score was also associated with lower FEV₁ in unadjusted (mean difference per z-unit: -2.00; 95% CI: -2.55, -1.45), minimally adjusted (mean difference per z-unit: -1.89; 95% CI: -2.42, -1.36) and fully adjusted (mean difference per z-unit: -1.89; 95% CI: -2.42, -1.36) and fully adjusted (mean difference per z-unit: -1.80; 95% CI: -2.33, -1.28) models. Higher FEV₁/FVC ratio was also observed to be associated with increasing HAA z-score in unadjusted (mean difference per z-unit: 0.57; 95% CI: 0.25, 0.89) and fully adjusted (mean difference per z-unit: 0.57; 95% CI: 0.25, 0.89) and fully adjusted (mean difference per z-unit: 0.57; 95% CI: 0.25, 0.89) and fully adjusted (mean difference per z-unit: 0.57; 95% CI: 0.25, 0.89) and fully adjusted (mean difference per z-unit: 0.57; 95% CI: 0.33, 0.97) models. Similar results were observed for the association between these lung function measures and binary elevated HAA (Table 4).

Higher HAA z-score was associated with dyspnoea in unadjusted (OR: 1.09 per z-unit; 95% CI: 1.03, 1.15), minimally adjusted (OR: 1.08 per z-unit; 95% CI: 1.02, 1.14) and fully adjusted (OR: 1.08 per z-unit; 95% CI: 1.02, 1.15) models (Table 3). An association was

also observed between binary elevated HAA and exertional dyspnoea in the unadjusted (OR: 1.75 per z-unit; 95% CI: 1.28, 2.37) and minimally adjusted models (OR: 1.52 per z-unit; 95% CI: 1.10, 2.09), but the estimate was unstable in the fully adjusted model (OR: 1.37 per z-unit; 95% CI: 0.98, 1.91) (Table 4).

No meaningful associations were observed between z-scores and cough (Table 3) or smoking pack-years (Table 4).

DISCUSSION

We examined several prediction modelling methods in order to develop optimized HAA reference equations and z-scores using a multi-ethnic, healthy, never-smoker sample of older adults. We demonstrate the validity of HAA z-scores as a measure of disease risk by showing that z-scores are associated with ILD features, such as smoking, and respiratory outcomes, including ILA, lung function and exertional dyspnoea.

Spirometric measures of lung function are known to have wide margins of variation in healthy samples.^{16,22,23} Clinical interpretations of spirometric measurements require reference to ranges standardized by height, weight, age, sex and race/ethnicity.^{16,23} More recently, CT-based measures of emphysematous lung have been shown to vary substantially by these same factors in a healthy non-smoking sample suggesting that demographic and body size characteristics play key roles in the natural variation of lung health measures.¹³ Importantly, models containing these same variables were shown to be optimal for predicting HAA out of several variables considered.

z-Scores measure the difference between an individual's observed measurement and what would be expected for that individual based on his or her demographic and anthropometric characteristics. This makes z-scores ideal for clinical decision-making because they adjust for variables that may confound interpretations of 'normal'.^{24,25} As z-scores also function as a dimensionality reduction variable, they may be used in place of observed values of HAA in small-sample research settings when the number of predictors is large relative to the sample size, impeding appropriate adjustment. However, we caution that as dimensionality reduction reduces the total information present in confounding variables, they may be underpowered to adjust for confounding.

Our methodology follows the latest recommendations for clinical prediction modelling^{26,27} and considers both parametric and non-parametric modelling approaches. In this sample, linear regression with OLS outperformed non-parametric machine learning approaches and the elastic net fit. Parametric methods may outperform non-parametric approaches for small data when the model is well specified with respect to parametric assumptions. OLS regression may also outperform an elastic net penalty when predictor collinearity does not affect coefficient stability and the model is at low risk for overfitting. OLS regression has the added benefit of providing unbiased coefficient estimates.

Some limitations of this study include the sample size of the healthy never-smoker subset used to develop the prediction models and the lack of an external validation set. It is also likely that additional variables not considered here play a role in the natural variation of

HAA. Furthermore, this cross-sectional analysis of the MESA cohort is left truncated on 10-year survivorship.

In conclusion, this study fills an important knowledge gap by establishing a normative range for HAA and by presenting reference equations to define expected values with respect to key variables associated with natural variation in HAA. These tools will aid in the interpretability of HAA in future studies and help to move HAA into the clinical sphere.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations:

AUC	area under the curve
СТ	computed tomography
eGFR	estimated glomerular filtration rate
FEV ₁	forced expiratory volume in 1 s
FVC	forced vital capacity
НАА	high attenuation area
ILA	interstitial lung abnormality
ILD	interstitial lung disease
IPF	idiopathic pulmonary fibrosis
MESA	Multi-Ethnic Study of Atherosclerosis
OLS	ordinary least square
RMSE	root of mean square error
ROC	receiver operating curve
ULN	upper limit of normal

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SUMMARY AT A GLANCE

To better understand the natural variation of HAA (a novel quantitative CT-based measure of subclinical ILD), we developed HAA reference equations and z-scores to define expected values of HAA with adjustment for key demographic and anthropometric variables, and we demonstrated that HAA z-scores correlate with several ILD features.

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Figure 1.

Flow diagram of MESA participants included in the healthy never-smoker subset. BMI, body mass index; CT, computed tomography; MESA, Multi-Ethnic Study of Atherosclerosis.

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Figure 2.

Density plots showing the distribution of: (A) observed HAA values, (B) predicted HAA values (solid) and ULN (dashed) and (C) z-scores (solid) and ULN (z-score = 1.634). Blue denotes the healthy never-smoker subset and orange denotes the complete study sample. (D) ROC curve showing sensitivity versus false positive rate (1 – specificity) for ILA detection for varying elevated HAA threshold values. AUC, area under the curve; HAA, high attenuation area; ILA, interstitial lung abnormality; ROC, receiver operating curve; ULN, upper limit of normal.

Table 1

Characteristics of study participants

	Complete study population	Healthy never-smoker subset
Participants, n	3110	696
HAA (%)	5.01 ± 2.38	4.76 ± 1.76
Age (years)	69 ± 9	69 ± 9
Sex		
Male	1489 (47.9%)	270 (38.8%)
Female	1621 (52.1%)	426 (61.2%)
Race/ethnicity		
Black	839 (27.0%)	161 (23.1%)
White	1202 (38.6%)	228 (32.8%)
Hispanic	658 (21.2%)	125 (18.0%)
Chinese	411 (13.2%)	182 (26.1%)
Weight (kg)	78.3 ± 17.5	70.3 ± 13.8
Height (cm)	165.4 ± 9.9	163.4 ± 9.4
Smoking history		
Never-smokers	1417 (45.6%)	696 (100%)
Former smokers	1459 (46.9%)	0 (0%)
Current smokers	234 (7.5%)	0 (0%)
Cigarette	20 ± 25	0 ± 0
pack-years [†]		
Lung function		
FVC (percent predicted)	97.1 ± 17.8	102.1 ± 14.7
FEV ₁ (percent predicted)	94.9 ± 19.9	103.4 ± 15.1
FEV ₁ /FVC ratio	74.0 ± 9.0	76.8 ± 6.0

Data are presented as mean ± SD or n (%). All variables were measured at MESA 10-year follow-up examinations during the years 2010-2012.

 † Among current and former smokers.

FEV1, forced expiratory volume in 1 s; FVC, forced vital capacity; HAA, high attenuation area; MESA, Multi-Ethnic Study of Atherosclerosis.

Table 2

Predictive performance of parametric and non-parametric modelling approaches for prediction of HAA measured from full lung CT scans

Modelling method	Variables included in the best subset	RMSE
Linear regression with least squares estimation	Race, height, weight, age, sex	1.64
Linear regression with elastic net penalty	Race, height, weight, age	1.72
Random forests	Race, weight, age, sex	1.68
Boosting	Race, sex	1.70
Support vector regression	Weight	1.81

RMSE is calculated on 10-fold cross-validation.

CT, computed tomography; HAA, high attenuation area; RMSE, root of mean square error.

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	u	Events	OR (95% CI)	P-value	u	Events	OR (95% CI)	<i>P</i> -value	u	Events	OR (95% CI)	P-value
HAA z-score												
Unadjusted	2406	299	1.39 (1.29, 1.51)	<0.001	3085	793	1.09 (1.03, 1.15)	0.002	3104	307	1.01 (0.92, 1.09)	0.81
Model 1	2352	285	1.39 (1.29, 1.51)	<0.001	3009	677	1.08 (1.02, 1.14)	0.01	3028	303	1.01 (0.93, 1.09)	0.77
Model 2	2352	285	$1.40\ (1.30,1.53)$	<0.001	3009	<i>611</i>	1.08 (1.02, 1.15)	0.01	3028	303	1.02 (0.93, 1.10)	0.68
Elevated HAA												
Unadjusted	2406	299	5.10 (3.47, 7.44)	<0.001	3085	793	1.75 (1.28, 2.37)	<0.001	3104	307	1.07 (0.64, 1.69)	0.78
Model 1	2352	285	5.42 (3.65, 8.00)	<0.001	3009	<i>6LT</i>	1.52 (1.10, 2.09)	0.01	3028	303	1.06 (0.64, 1.69)	0.80
Model 2	2352	285	5.68 (3.74, 8.60)	<0.001	3009	<i>6LT</i>	1.37 (0.98, 1.91)	0.06	3028	303	1.11 (0.66, 1.77)	0.68

as the upper fifth percentile of the ment. Model 2 is adjusted for all terms included in model 1 and race, height, BMI, age and sex.

BMI, body mass index; HAA, high attenuation area; ILA, interstitial lung abnormality; OR, odds ratio per z-unit increase or for elevated HAA compared to non-elevated HAA.

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		FVC			FEV ₁			FEV ₁ /FVC			$\mathbf{Pack-years}^{\hat{f}}$	
	u	Expected rate of change (95% CI)	<i>P</i> -value	u	Expected rate of change (95% CI)	<i>P</i> -value	u	Expected rate of change (95% CI)	<i>P</i> -value	u	Expected rate of change (95% CI)	<i>P</i> -value
HAA z-score												
Unadjusted	2740	-2.61(-3.10, -2.13)	<0.001	2735	-2.00(-2.55, -1.45)	<0.001	2789	0.58 (0.25, 0.90)	0.001	1425	$0.06\ (0.00,\ 0.11)$	0.06
Model 1	2674	- 2.49 (- 2.96, -2.03)	<0.001	2669	-1.89 (-2.42, -1.36)	<0.001	2669	$0.57\ (0.25,0.89)$	<0.001	1412	$0.06\ (0.00,\ 0.11)$	0.06
Model 2	2674	-2.49 (-2.95, -2.03)	<0.001	2669	-1.80 (-2.33, -1.28)	<0.001	2669	$0.65\ (0.33,\ 0.97)$	<0.001	1412	0.05 (-0.01, 0.10)	0.10
Elevated												
HAA												
Unadjusted	2740	-8.05 (-10.91, -5.18)	<0.001	2735	-5.12 (-8.36, -1.88)	0.002	2789	2.86 (0.95, 4.77)	0.003	1425	$0.18 \left(-0.14, 0.49\right)$	0.28
Model 1	2674	-6.61 (-9.39, - 3.83)	<0.001	2669	-3.83 (-6.97, -0.68)	0.02	2669	2.74 (0.86, 4.61)	0.004	1412	$0.15 \left(-0.16, 0.47\right)$	0.34
Model 2	2674	-7.57 (-10.31, -4.84)	<0.001	2669	-5.03 (-8.13, -1.93)	0.001	2669	2.32 (0.46, 4.18)	0.01	1412	0.21 (-0.10, 0.52)	0.19
Associations are fifth percentile of 2 is adjusted for	e estimated of the popu	I by ordinary least squares I lation (z-score 1.645). M included in model 1 and rec	regressions, lodel 1 is ad	P-values	are calculated from two- study site, smoking state	sided t-tests 1s, pack-yea	and RMS rs, waist c	E is calculated on 10-fold ircumference, estimated ξ	l cross-valida glomerular fi	ation. Elev Itration ra	vated HAA is defined as t ate and educational attain	he upper ment. Model

 \dot{r} Analysis is conducted among current and former smokers, and models are not adjusted for smoking status or pack-years.

BMI, body mass index; FEV1, forced expiratory volume in 1 s; FVC, forced vital capacity; HAA, high attenuation area; RMSE, root of mean square error.