

The predictive role of lung clearance index on FEV₁ decline in cystic fibrosis

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ABSTRACT

Background. The lung clearance index (LCI) is a sensitive lung function index that is used to detect early lung disease changes in children with cystic fibrosis (CF). This study aimed to define the predictive role of baseline LCI, along with other potential factors on the change in forced expiratory volume in one second (FEV₁) during one-year follow-up in CF patients who had a percent predicted (pp) FEV₁ ≥80.

Methods. LCI was concurrently performed on 57 CF patients who had ppFEV₁ ≥80 at month zero. The ppFEV₁ decline was evaluated prospectively during the one year follow up. The primary outcome of ppFEV₁ decline in the study group in one year was dichotomized according to the median value for the decline in ppFEV₁, which was 3.7. The LCI value predicting ppFEV₁ decline at the end of one year was calculated with receiver operating characteristic curve analysis. Regression analysis was performed. Furthermore, a decision tree was constructed using classification and regression tree methods to better define the potential effect of confounders on the ppFEV₁ decline.

Results. The LCI value for predicting ppFEV₁ decline >3.7% at the end of one year was 8.2 (area under the curve: 0.80) Multivariable regression analysis showed that the absence of the F508del mutation in at least one allele, LCI >8.2 and initial FEV₁ z-score were predictors of a ppFEV₁ decline >3.7 (p<0.001). Factors altering ppFEV₁ decline >3.7% at the end of one-year evaluated by decision trees were as follows: initial FEV₁ z-score, type of CFTR mutation, LCI value and initial weight-for-age z-score.

Conclusions. LCI is sensitive for predicting ppFEV₁ decline in patients with ppFEV₁ ≥80 along with the initial FEV₁-z-score and type of CFTR mutation.

Key words: cystic fibrosis, multiple breath wash-out, lung clearance index, spirometry, FEV₁ decline.

Cystic fibrosis (CF) is one of the most common life-limiting genetic diseases that mainly affects the respiratory system. In recent years, the early diagnosis of the disease, close monitoring of lung health and on-site therapy interventions have prolonged the lives of patients with CF (pwCF). Pulmonary involvement is of early onset and progresses insidiously.¹

Conventional spirometry has been the standard test used for evaluating lung function in patients with CF. Failure to perform spirometry effectively in young children and the failure to capture changes in spirometry in the early stages of the disease have led to a search for new tests. The lung clearance index (LCI) derived from the multiple breath washout (MBW) test, is the most commonly used index for measuring ventilation distribution inhomogeneity.^{2,3}

The rate of forced expiratory volume in one second (FEV₁) decline is related to disease progression and has been shown to predict

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mortality. To date risk factors for FEV₁ decline have been identified as non-modifiable factors such as sex, age, CF transmembrane regulator (CFTR) genotype, history of meconium ileus, pancreatic insufficiency, and modifiable factors such as pulmonary exacerbations, airway colonization high initial FEV₁ and CF-related diabetes.⁴ However, the predictive role of LCI along with these parameters on FEV₁ decline has not been fully established.

The primary aim of this study was to define the predictive role of baseline LCI derived from the MBW technique, along with other potential factors by decision trees on FEV₁ change in a one year follow up in pwCF who had a percent predicted (pp) FEV₁ ≥80 when LCI was performed. Our hypothesis in this study was that a higher baseline LCI would be related to a higher FEV₁ decline in patients with accompanying factors and we also aimed to investigate these factors.

Materials and Methods

Study design

Children between 5-18 years with a confirmed diagnosis of CF, attending a tertiary pediatric CF clinic, who were able to cooperate to spirometry and LCI and had ppFEV₁ ≥80 were enrolled.⁵ This was a single-center, prospective cohort study. The study was approved by Hacettepe University Ethics Committee for Non-Interventional Studies (GO 20-637). Patients and caregivers gave written consent to participate in the study. The study was conducted between April 2018- April 2019.

The sample size was calculated using the formula below:

$$n = \frac{Z_{1-\alpha/2}^2 \times S_P \times (1 - S_p)}{L^2 \times (1 - Prevalans)}$$

(S_p = anticipated specificity, α = size of the critical region [1 - α is the confidence level], Z_{1-α/2} = standard normal deviation corresponding to the specified size of the critical region [α] L = absolute precision desired on either side

[half-width of the confidence interval, CI] of sensitivity or specificity)⁶

According to this formula, when Z_{1-α/2}=1.96, SP=0.85, L=0.15 and prevalence=0.39 (based on the Turkish cystic fibrosis registry data, where 39% of the patients have low FEV₁% predicted⁷), the minimum sample size was calculated as 56.

The MBW test was performed at the initial visit at the time of inclusion, along with spirometry. The spirometry was then performed at 3, 6 and 12 months during routine outpatient visits. The study was terminated by performing the last spirometry at the twelfth month after the MBW was performed. The patients' demographic data including weight, height, and body mass index (BMI), transcutaneous oxygen saturation, physical examination findings, microorganism growth and colonization status in sputum culture, were recorded. The radiological examinations findings which were performed according to the decision of the clinician, considering the clinical condition of the patients were recorded. At each visit, patients were questioned for acute pulmonary exacerbation (aPEX) signs and compliance with their standard therapies (medications and chest physiotherapy). All aPEX events and intravenous / oral antibiotic therapies that occurred between visits were recorded.

aPEX was defined according to the study conducted by Fuchs et al.⁸ Chronic *P. aeruginosa* infection was defined according to the study by Lee et al.⁹ Mucus samples for microbiologic analysis were collected at all test occasions.

Measurements

Patients performed the MBW test using a Nitrogen Analyzer connected to a Sensormedics Vmax Spectra 22 Device. Nitrogen (N₂) gas was used as the inert gas. The MBW tests were performed in alliance with the segregation rules according to the ATS/ERS consensus statement.¹⁰⁻¹³ MBW measurements were performed by the same staff using the same equipment. MBW test was performed with

the patient in the sitting position, breathing through a mouthpiece, wearing a nose clip. After a relaxed tidal breath, the patient was switched to 100% oxygen breathing until the end-tidal N_2 concentration declined below $1/40^{\text{th}}$ of the starting end-tidal concentration for at least three consecutive breaths. The LCI was calculated as the cumulative expired volume divided by the functional residual capacity (FRC) at $1/40$ of the initial N_2 concentration. MBW test completion was established when three technically acceptable tests were achieved. All MBW traces were controlled for technical quality and convenient breathing pattern. Calibration was performed prior to testing on each test day. Quality control of MBW trials were conducted according to the 2013 ATS/ERS MBW consensus guideline.¹⁰⁻¹²

Spirometry was performed according to the ATS/ERS statement.¹⁴ Spirometry completion was established when three technically acceptable tests were achieved. The percent predicted values and z-scores were calculated using all age prediction equations for spirometry from the Global Lung Function Initiative.¹⁵ The decline in ppFEV₁ at the 12th month was calculated according to the initial ppFEV₁.

Statistical analyses were performed using the SPSS version 25.0 software package (IBM, SPSS, Chicago, IL, USA). Normal distribution of data was tested analytically (Kolmogorov-Smirnov/ShapiroWilk tests) and visually (histogram, probability plots). Categorical variables were described as relative and absolute frequencies. Continuous variables which are normally distributed were summarized as mean \pm standard deviation (SD) and analyzed using Student's t-test. Repeated measure ANOVA was used for dependent variables for repeated measures that are normally distributed. For significant ANOVA results, Bonferroni test was used for binary comparisons between groups.

Correlations between parameters were assessed using Pearson correlation coefficient (r). The primary outcome ppFEV₁ decline in one year was dichotomized according to the median

value for the decline in ppFEV₁ in one-year follow-up which was 3.7. To define LCI value to predict ppFEV₁ decline at the end of one year the patients were grouped into two: patients who had a decline in ppFEV₁ ≤ 3.7 and patients who had a decline in ppFEV₁ > 3.7 and it was calculated using receiver operating characteristic (ROC) curve analysis. The cut-off point was determined based on Youden index. Regression analysis was performed to adjust the effect of potential confounders on the ppFEV₁ decline. The SMOTE approach was used to rebalance the data. Independent variables that had a relation with dependent variables with a p-value of ≤ 0.25 in univariable regression analysis were further analyzed in a multivariable regression model. We used backward logistic regression analysis using ppFEV₁ decline > 3.7 (dichotomous variable) as the dependent variable and presence/absence of F508del mutation in at least one allele, baseline LCI, initial FEV₁ z-score, initial weight for age (WFA) as independent variables.

A decision tree was constructed to better define the potential effect of confounders on the ppFEV₁ decline. In the decision tree patients were divided into two groups according to the median ppFEV₁ decline > 3.7 . The decision trees were created using the R programming language version 4.0.0. "Rpart" library was used to construct decision trees. The data was divided into training and test data. During the learning phase, a training dataset was used to develop the classification model. In the second step, the test set was used to evaluate the classification model's accuracy after it had been trained using the training set. Accuracy rate, sensitivity and specificity were measures used to assess the model's performance. The "Performance Estimation" library was used to rebalance data using the SMOTE technique.¹⁶⁻¹⁸

Results

Over the one-year recruitment period among 360 pwCF aged 0-18 years, 57 subjects with ppFEV₁ ≥ 80 who were able to perform the MBW and

spirometry were enrolled. The demographic, microbiologic, radiologic, treatment data and the latest chest X-rays and thorax CT of the patients are summarized in Table I. During the one-year follow-up, 23 patients (40.3%) had aPEX; 14 (24.6%) patients had one, 8 (14%) patients had two, and one (1.8%) patient had 4 exacerbations. The median pulmonary exacerbation number was 1 (interquartile range [IQR]: 1-2) per patient per year. A total of 54 courses of non-prophylactic antibiotics (oral 84.4%, intravenous 15.6%) were used during the study.

The initial, 3, 6, and 12-month spirometry values and BMI-z-score are summarized in Table II. A significant decline in FEV₁ (pp-z-score) and FEF₂₅₋₇₅ (pp-z-score) in the third, sixth, and twelfth months was shown by using repeated measure ANOVA. There was no significant decline in BMI z-score at the end of one year.

The mean LCI value was 7.39 (\pm 2.00). The mean LCI was higher in girls, patients without F508del mutation in at least one allele, patients with pancreatic insufficiency, and patients with bronchiectasis/atelectasis. However, they were not statistically significant (p=0.7, p=0.6, p=0.6, p=0.1 respectively) (Table III). According to the Pearson correlation test, there was no correlation between LCI and initial, and twelfth month ppFEV₁ and z-scores (r=-0.1 p=0.46, r=-0.08 p=0.53 for initial ppFEV₁ and z-scores; r=-0.18 p=0.20, r=-0.13 p=0.3 for twelfth month ppFEV₁ and z-scores). No correlation was found between BMI-z-score and FEV₁, FEF₂₅₋₇₅ z-scores and LCI.

The median ppFEV₁ decline for the study group was 3.7 (IQR: 0.5-6.5). The LCI value for predicting ppFEV₁ decline >3.7 at the end of one year was calculated using the ROC curve analysis and was found to be 8.2 (area under the ROC curve: 0.80, sensitivity 46.2% [95% CI: 24.1-68.3], specificity 86.4% [95% CI: 75.7-96.3], p<0.001) (Fig. 1).

Regression analysis was performed to evaluate the effect of covariates on ppFEV₁ decline. Age, gender, age at diagnosis, presence/absence of F508del mutation in at least one allele, baseline LCI, initial FEV₁ z-score, total acute pulmonary exacerbations, *Pseudomonas aeruginosa* and *Staphylococcus aureus* colonization, presence of bronchiectasis \pm atelectasis, initial WFA and BMI z-score were selected as covariates for predicting a ppFEV₁ decline >3.7 in univariable analysis. Multivariable backward logistic regression analysis with presence/absence of F508del mutation in at least one allele, baseline LCI, initial FEV₁ z-score, initial WFA z-score showed that absence of F508del mutation in at least one allele, LCI >8.2 and initial FEV₁ z-score were predictors of ppFEV₁ decline >3.7 (p<0.001, Table IV, accuracy: 0.84, sensitivity: 0.81, specificity: 0.88).

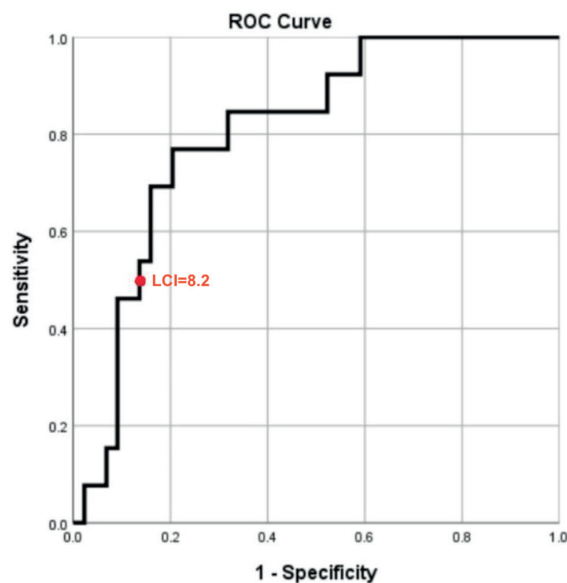


Fig. 1. Receiver operating characteristic (ROC) curve of lung clearance index for predicting ability of FEV₁ decline by > 3.7% at the end of one year. The red dot represents the cut of value lung clearance index=8.2 which is defined by the ROC curve analysis [area under the ROC curve: 0.80, sensitivity 46.2% (95% CI: 24.1-68.3), specificity 86.4% (95% CI: 75.7-96.3), p<0.001]

Table I. Demographic, microbiologic, radiologic, and treatment data of subjects at time of the initial visit (N=57).

Male/female	31/26
Age, yr, mean (SD)	11.6 (3.5)
Age at diagnosis, yr, median (IQR)	0.3 (0.2-1.5)
Genotype, n (%)	
F508del/F508del	10 (17.5)
F508del/other	13 (22.8)
Other/other	34 (59.7)
Exocrine pancreatic insufficiency	50 (87.7)
Chronic liver disease, n (%)	15 (26.3)
Cystic fibrosis related diabetes, n (%)	4 (7)
Weight for age z-score, mean (SD)	-0.31 (1.10)
Height for age z-score, mean (SD)	-0.47 (1.08)
BMI z-score, mean (SD)	0.01 (1.09)
Transcutaneous oxygen saturation (%), mean (SD)	96.9 (1.5)
Microbiological growth (sputum or deep oropharyngeal), n (%)	
Methicillin sensitive <i>Staphylococcus aureus</i>	32 (56.1)
Methicillin resistant <i>Staphylococcus aureus</i>	9 (15.8)
<i>Pseudomonas aeruginosa</i>	16 (28.1)
<i>Haemophilus influenzae</i>	2 (3.5)
<i>Burkholderia cepacia</i>	2 (3.5)
<i>Stenotrophomonas maltophilia</i>	2 (3.5)
<i>Achromobacter</i>	1 (1.8)
Candida species	9 (15.8)
<i>Aspergillus fumigatus</i>	2 (3.5)
Microbiological colonization, n (%)	
<i>Staphylococcus aureus</i>	42 (71.9)
<i>Pseudomonas aeruginosa</i>	15 (26.3)
Chest X-ray, n (%)	57 (100)
Normal	3 (5.3)
Pulmonary infiltration	3 (5.3)
Chronic changes (air trapping, mosaic pattern, bronchiectasis and/ or atelectasis)	51 (86.4)
Thorax CT, n (%)	10 (17.5)
Atelectasis	6 (10.5)
Bronchiectasis	8 (14)
Mucus plugs	8 (14)
Lymphadenopathy	4 (7)
Medications	
Inhaled dornase alpha, n (%)	
Adherent to treatment	52 (91.2)
Not adherent to treatment	3 (5.3)
Not recommended by the physician	2 (3.5)
Inhaled mannitol or hypertonic saline, n (%)*	
At month zero	0 (0)
Chest physiotherapy, n(%)	
Regular	44 (77.2)
Irregular	13 (22.8)

BMI: body mass index, CT: computed tomography, IQR: interquartile range, SD: standard deviation, yr: year.

* Only one patient started inhaled mannitol treatment at the third month, and five patients started inhaled hypertonic saline treatment at the sixth month visit.

Table II. Percent predicted and z-scores of FEV₁ and FEF₂₅₋₇₅ and BMI z-score at initial, third, sixth and twelfth month.

Spirometry parameters	Clinical visit time (month)				p value
	0	3	6	12	
FEV ₁ mean (SD)					
% predicted	107.0 (15.7)	102.6 (17.7)	102.6 (17.7)	101.4 (18.9)	0.001*
z-score	0.55 (1.59)	0.12 (1.90)	0.12 (1.90)	-0.09 (1.98)	0.001**
FEF ₂₅₋₇₅ mean (SD)					
% predicted	107.2 (35.9)	98.8 (37.4)	98.8 (37.4)	94.1 (37.3)	<0.001+
z-score	0.14 (1.94)	-0.29 (2.11)	-0.29 (2.11)	-0.53 (2.17)	<0.001**
BMI z-score, mean (SD)	0.01 (1.09)	-0.09 (1.06)	-0.09 (1.06)	-0.08 (1.03)	0.5

* FEV₁ %predicted month 0 is statistically significant with FEV₁ %predicted month 6 (sig: 0.01) and FEV₁ %predicted month 12 (sig: 0.01), but not with FEV₁ %predicted month 3 (sig: 0.06). FEV₁ %predicted month 3 is not significant with FEV₁ %predicted month 6 (sig: 0.9) and FEV₁ %predicted month 12 (sig: 0.9). FEV₁ %predicted month 6 is not significant with FEV₁ %predicted month 12 (sig: 0.9).

** FEV₁ z score month 0 is statistically significant with FEV₁ z score month 3 (sig: 0.03) FEV₁ z score month 6 (sig: 0.01), FEV₁ z score month 12 (sig: 0.005). FEV₁ z score month 3 is not significant with FEV₁ z score month 6 (sig: 0.9) and FEV₁ z score month 12 (sig: 0.9). FEV₁ z score month 6 is not significant with FEV₁ z score month 12 (sig: 0.9).

+ FEF₂₅₋₇₅ %predicted month 0 is statistically significant with FEF₂₅₋₇₅ %predicted month 3 (sig: 0.01), FEF₂₅₋₇₅ %predicted month 6 (sig: 0.01) and FEF₂₅₋₇₅ %predicted month 12 (sig: 0.001). FEF₂₅₋₇₅ %predicted month 3 is not significant with FEF₂₅₋₇₅ %predicted month 6 (sig: 0.9) and FEF₂₅₋₇₅ %predicted month 12 (sig: 0.6). FEF₂₅₋₇₅ %predicted month 6 is not significant with FEF₂₅₋₇₅ %predicted month 12 (sig: 0.7).

** FEF₂₅₋₇₅ z score month 0 is statistically significant with FEF₂₅₋₇₅ z score month 3 (sig: 0.02) FEF₂₅₋₇₅ z score month 6 (sig: 0.01), FEF₂₅₋₇₅ z score month 12 (sig: 0.001). FEF₂₅₋₇₅ z score month 3 is not significant with FEF₂₅₋₇₅ z score month 6 (sig: 0.9) and FEF₂₅₋₇₅ z score month 12 (sig: 0.9). FEF₂₅₋₇₅ z score month 6 is not significant with FEF₂₅₋₇₅ z score month 12 (sig: 0.9).

BMI: body mass index, FEF₂₅₋₇₅: forced expiratory flow between 25% and 75% of forced vital capacity, FEV₁: forced expiratory volume in the first second, SD: standard deviation.

Table III. The mean lung clearance index value of patients according to different categories.

	Lung Clearance Index Mean (SD)	p value
Gender (n)		0.7
Female (26)	7.52 (1.40)	
Male (31)	7.28 (1.56)	
CFTR mutation (n)		0.6
F508del in at least one allele present (23)	7.21 (1.74)	
F508del absent (34)	7.51 (2.18)	
Pancreatic insufficiency (n)		0.6
Present (50)	7.46 (1.10)	
Absent (7)	6.88 (0.94)	
Bronchiectasis / atelectasis (n)		0.1
Present (33)	7.76 (2.10)	
Absent (24)	6.90 (1.57)	
Pulmonary exacerbation during one year interval (n)		0.3
Present (23)	7.10 (1.50)	
Absent (34)	7.60 (1.30)	

* CFTR: cystic fibrosis transmembrane conductance regulator, SD: standard deviation.

Table IV. Backward logistic regression analysis for variables predicting ppFEV₁ decline >3.7.

Model	Odds ratio	95% CI	Sig
Step 1			
Constant	1.34	0.65-2.75	0.4
Absence of F508del mutation	0.014	0.002-0.093	<0.001
LCI >8.2	7.15	2.1-24.0	0.001
Initial FEV ₁ z- score	0.19	0.09-0.41	<0.001
Initial WFA z -score	1.22	0.62-2.42	0.6
Step 2			
Constant	1.32	0.65-2.70	0.4
Absence of F508del mutation	0.013	0.002-0.082	<0.001
LCI >8.2	7.65	2.31-25.31	<0.001
Initial FEV ₁ z- score	0.19	0.09-0.42	<0.001

CI: confidence interval, LCI: lung clearance index, ppFEV₁: percent predicted forced expiratory volume in one second, sig: significance, WFA: weight for age.

The independent variables that had a relation with >3.7 ppFEV₁ decline in univariate analysis were also analyzed by decision trees. Patients who had ppFEV₁ decline >3.7 were grouped as "high", whereas the second group consisted of patients who had ppFEV₁ decline ≤3.7 and were grouped as "low" in the decision trees. Only 22.8% of the patients (n:13) had ppFEV₁ decline >3.7. To overcome the class imbalance problem in the first step 10% of the dataset was reserved for validation. The SMOTE approach was used to resample and balance the remaining dataset. The data set which became balanced in terms of the class distributions of outcome then were then split into parts: 70% for train data set and 30% for test data set. Using the train data set, a decision tree was created. The first splitting variable is initial FEV₁ z-score followed by CFTR mutation analysis, LCI ≤8.2 defined by ROC curve, and initial WFA z-score. The predicted class is shown by colors, with pink representing a decrease of ≤3.7% in FEV₁ and blue representing a decrease of >3.7% in FEV₁.

The classification rules of the decision tree generated by the CART algorithm are given as follows:

- 1) If a patient's initial FEV₁-z-score is ≥0.62, a decline in ppFEV₁ in one year follow up will be ≤3.7 with a 100% probability.
- 2) If a patient's initial FEV₁-z-score is <0.62 and has a F508del mutation in at least one allele, a decline in ppFEV₁ in one year follow up will be ≤3.7 with a 91% probability.
- 3) If a patient's initial FEV₁-z-score is between -0.60 and 0.62, the patient does not have a F508del mutation, LCI is ≤8.2 and initial WFA-z-score is ≥0.27, a decline in ppFEV₁ in one year follow up will be ≤3.7 with an 80% probability.
- 4) If a patient's initial FEV₁-z-score is between -0.60 and 0.62, the patient doesn't have a F508del mutation, LCI is ≤8.2 and initial WFA-z-score is less than 0.27, a decline in ppFEV₁ in one year follow up will be >3.7 with an 88% probability.
- 5) If a patient's initial FEV₁-z-score is between -0.60 and 0.62, the patient doesn't have a F508del mutation and LCI >8.2, a decline in ppFEV₁ in one year follow up will be >3.7 with a 93% probability.
- 6) If a patient's initial FEV₁-z-score is less than -0.60, a decline in ppFEV₁ in one year follow up will be >3.7 (Fig. 2).

The results of the decision tree show that factors altering ppFEV₁ decline >3.7% at the end of one year are the initial FEV₁ z-score, type of CFTR mutation, LCI value and initial weight-for-age z-score. According to the results of performance

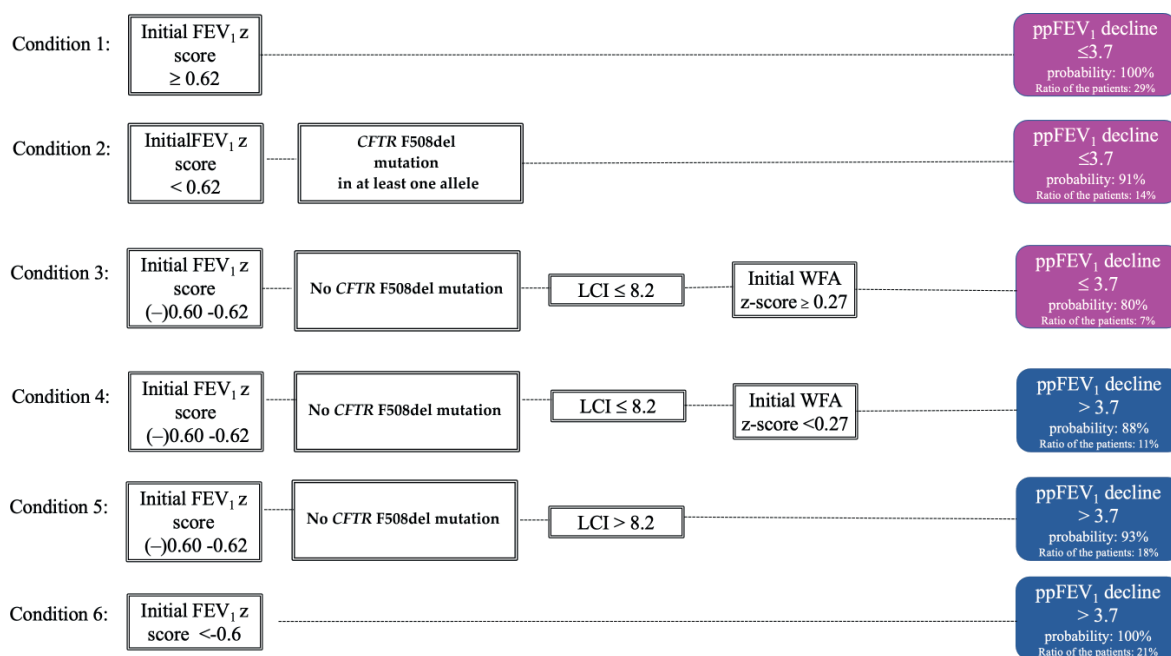


Fig. 2. Decision tree for determining FEV1 decline >3.7.

The first splitting variable is initial FEV₁ z-score followed by CFTR mutation analysis, LCI ≤ 8.2 defined by ROC curve, and initial WFA z-score. The predicted class is shown by colors, with pink representing a decrease of ≤3.7% in FEV₁ and blue representing a decrease of >3.7% in FEV₁.

The classification rules of the decision tree generated by the CART algorithm are given in the figure and in the text.

*CFTR: cystic fibrosis transmembrane regulator, LCI: lung clearance index, ppFEV₁: percent predicted forced expiratory volume in one second, WFA: weight for age.

metrics the model’s ability to predict whether a decrease in ppFEV₁ in one-year follow-up was lower than or equal to greater than 3.7% which is quite strong (Train dataset accuracy: 94.7%, sensitivity: 95%, specificity 95%; Test dataset accuracy: 87.5%, sensitivity: 81%, specificity: 94%; validation dataset accuracy: 84.6%, sensitivity: 100%, specificity 80%). The detailed results of the decision tree model including the accuracy, sensitivity and specificity and confusion matrix for each dataset (train, test and validation) are shown in Table V and VI.

Discussion

In our tertiary pediatric CF clinic, spirometry was performed on 250 patients during 2019, among these patients 23% had ppFEV₁ ≥ 80. Therefore, this wide cohort of patients required a more sensitive method for monitoring lung health which would enable us to apply the right interventions when necessary. To achieve this

goal, we aimed to determine an LCI value that could be a warning for a near-future ppFEV₁ decline along with other potential predictors. In this first report of LCI results in pwCF from Turkey designed in a mild CF population including school-age children and adolescents we found that ppFEV₁ significantly declined during a one-year follow-up in patients who initially had a ppFEV₁ ≥ 80. This decline in ppFEV₁ was >3.7 in almost one quarter of the patients. For this reason, LCI values were calculated using ROC curve analysis; the value that could predict FEV₁ decline >3.7% was found to be 8.2. The other predictors for ppFEV₁ decline along with LCI were the initial FEV₁-z-score, type of CFTR mutation and initial WFA z-score.

The annual ppFEV₁ decline in different studies ranges between 0.06-3.64%.⁴ The ppFEV₁ decline in our study, which was 3.7%, is significantly higher compared to other studies. The higher

Table V. The accuracy, sensitivity and specificity of the decision tree model.

	Train dataset	Test dataset	Validation dataset
ppFEV ₁ decline <3.7	38 (50%)	16 (50%)	10 (76.9%)
ppFEV ₁ decline ≥ 3.7	38 (50%)	16 (50%)	3 (23.1%)
Accuracy	94.74 %	87.5 %	84.62 %
Sensitivity	0.947	0.812	1.000
Specificity	0.947	0.937	0.800
Positive predictive value	0.947	0.928	0.600
Negative predictive value	0.947	0.833	1.000

ppFEV₁: percent predicted forced expiratory volume in one second

Table VI. The confusion matrix for train, test and validation dataset for the decision tree model.

Prediction for train dataset, n	Reference for train dataset, n	
	Low	High
Low	36	2
High	2	36
Prediction for test dataset, n	Reference for test dataset, n	
	Low	High
Low	15	3
High	1	13
Prediction for validation dataset, n	Reference for validation dataset, n	
	Low	High
Low	8	0
High	2	3

ppFEV₁ decline in our cohort can be explained by the following: First of all, our cohort consisted of children with high baseline ppFEV₁ which has been reported as an independent risk factor for FEV₁ decline in several studies.¹⁹⁻²¹ The reason for ppFEV₁ decline in patients with high baseline FEV₁ can be explained by several reasons. First of all, physicians may underestimate the early decline in FEV₁ which is still within normal limits and not intervene as quickly as possible in pwCF with moderate/severe disease. Secondly, patients might think that being in a state of wellness means there is no need for any treatment and they may not take their standard therapies/physiotherapy either involuntarily (simply forgetting their therapies) or voluntarily (nonadherence). In our study, a quarter of the patients were non-adherent to physiotherapy which we believe

is related to the high number of adolescents (n=44) who are prone to being non-adherent to treatments. We believe the second reason for the higher FEV₁ decline when compared with the literature is associated with newborn screening (NBS). It's well known that patients diagnosed with NBS have improved respiratory, nutritional outcomes and survival.²² NBS for CF was implemented in Türkiye on 01.01.2015 by the Ministry of Health, so none of the patients enrolled in the study were diagnosed through NBS. The third reason for higher FEV₁ decline can be explained by the low rate of inhaled hypertonic saline (HS) or inhaled mannitol which is mainly due to financial obstacles as the drugs were only reimbursed after 2018. All of the factors mentioned above put our mild CF patients at greater risk compared to pwCF with moderate/severe disease.

The mean LCI for healthy populations ranges between 6.2-7.2, and the upper normal limit for LCI, which is defined as mean LCI+2SD, is reported as 7.9-8.2.²³⁻²⁵ Differences in normal values between studies can be explained by age differences, methods of analysis, software used, devices and set-up, and the tracer gas used. The upper limit of normal (ULN) of LCI for healthy school-age children and adolescents was defined as 7.91.²⁵ When this upper limit was taken into consideration, almost one quarter of the patients had an elevated LCI. In the study of Ellemunter et al.²⁶, they found an elevated LCI in almost 80% of the mild pwCF. Similarly, in the 2014 study of Fuchs et al.²⁴, they showed that 83% had an increased LCI. Also, Fuchs et al.²⁴ reported a mean baseline LCI of 8. One of the reasons for the lower LCI in the current study may be the difference in FEV₁-z-scores of patients who were enrolled. In our study, the mean FEV₁-z-score was 0.55, whereas it was -0.26 in the study of Fuchs et al. In addition, in the current study, a more homogeneous group by age was included, whereas adults were also involved in Fuchs et al. study.

With the understanding that early lung disease arises in the peripheral airways, LCI has become a tool that can predict future respiratory functions when the patient has preserved spirometry values.^{27,28} In our study, the LCI value, predicting near future FEV₁ decline was 8.2 which is slightly higher than the ULN in published reference equations.²⁵ LCI measurement can be performed in functional lung tissue, therefore it is mostly preferred in mild CF patients in order to detect early lung changes and monitor disease progression. LCI can also be used to detect pulmonary exacerbations and first PA growth.^{29,30} Besides, there are several studies evaluating medical treatment (dornase alpha, hypertonic saline, CFTR modulators) responses by LCI measurements.^{31,32} Recently, Kurz et al. showed that increased LCI is associated with a greater risk of death or lung transplantation.³³ Future studies are needed to define how to implement LCI monitoring in CF patients in daily practices

in terms of LCI monitorization frequency and the change in LCI value. It is also necessary to clearly state whether follow-up with LCI improves management and survival in patients with CF.

Our study has some limitations. This was a single-center study designed without a control group. Compliance with medical therapies was evaluated according to the patients' declaration which may cause information/recall bias. The reported compliance with medical therapies has been shown to be less sensitive and correct by overestimating the real number of performed treatments compared to electronic measures such as inhalers counting the inhalation cycles. Besides the low sensitivity of the LCI cut off defined by the ROC curve to predict FEV₁ decline can be related to the sample size which is within the acceptable limits but still within the lower limit of normal. Also, due to low sensitivity, the false negative rate was high and negative diagnoses were not very reliable. Because of this limitation, our results cannot predict the FEV₁ decline by only LCI, but can predict the FEV₁ decline along with initial FEV₁-z-score, initial WFA z-score and type of CFTR mutation. To overcome this limitation larger prospective, longitudinal, multicenter studies are needed. In addition, the relatively small sample size is a problem in terms of applying logistic regression and decision tree models. This problem affects the validity of the results derived from regression and classification models. In addition, since the class distributions are imbalanced, the use of the SMOTE method and the low number of observations in the validation set can reduce the reliability of the results. Last of all, the variability in LCI ranging between 15-25%, which has been shown to be highest in pwCF with more severe lung disease could not be interpreted because we only measured LCI values at the beginning of the study.³⁴ However, the study conducted by Oude Engberink et al.³⁵ also showed that the variability is lower in LCI measurements taken 24 hours apart, which suggests that our findings can reflect close to normal LCI values.

Conclusions

Our study confirms that LCI is a helpful tool for predicting FEV₁ decline in patients who have ppFEV₁ ≥80 along with several parameters such as the initial FEV₁ z-score, type of CFTR mutation. Given the feasibility of LCI in infants and preschool children, we should encourage its use in children with mild CF, alongside spirometry, to enhance medical treatment strategies and improve survival.

Ethical approval

Informed consent was obtained for participation in the study and the study was approved by Hacettepe University Ethics Committee for Non-Interventional Studies (GO 20-637 Date: February 2018).

Author contribution

The authors confirm contribution to the paper as follows: study conception and design: BÖ, EY, NE, HKÜ, DD, UÖ, NK; data collection: BÖ, DAT, BS, CC; analysis and interpretation of results: BÖ, HKÜ, EY, DD, UÖ, NK; draft manuscript preparation: BÖ, EY, HKÜ, NE. All authors reviewed the results and approved the final version of the manuscript.

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Conflict of interest

The authors declare that there is no conflict of interest.

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