Subclinical Left Ventricular Dysfunction is Influenced by Genotype Severity in Patients with Cystic Fibrosis

Michael Pallin¹, Dominic Keating¹, David M Kaye², Tom Kotsimbos¹ and John W Wilson¹

¹Cystic Fibrosis Service, Alfred Health and Central Clinical School, Monash University, Melbourne, VIC, Australia. ²Cardiovascular Medicine, Alfred Health and Central Clinical School, Monash University, Melbourne, VIC, Australia.

Clinical Medicine Insights: Circulatory, Respiratory and Pulmonary Medicine Volume 12: 1-5 © The Author(s) 2018 Article reuse guidelines: sagepub.com/journals-permissions DOI: 10.1177/1179548418794154



ABSTRACT

BACKGROUND AND OBJECTIVE: Over 2000 genotypes in the cystic fibrosis (CF) gene have been described. These genotypic differences result in variable clinical manifestations of CF, with severity of disease dependent on CF transmembrane conductance (CFTR) protein function. CFTR is widely distributed in nucleated cells, including cardiac myocytes, but the effect of genotype on cardiac function is not known.

METHODS: This retrospective review of echocardiographic data is from a single adult CF centre between 2000 and 2015. Patients were cohorted based on the functional classification of genotype. 'Severe' patients had both CF genes from functional classification groups 1-3; 'mild' patients had one or no gene from these groups, or in the event of the second gene being unknown were pancreatic sufficient.

RESULTS: Genotype and echocardiography were recorded during the inclusion period in 100 patients, 79 of whom were classified as having severe genotypes. Although the severe group were younger they had a lower fractional shortening (33.66 ± 6.6 vs 36.9 ± 6.3 , P < .05), left atrial area (14.9 \pm 3.6 versus 18.0 \pm 4.2 cm²; P < .01) and volume (39.9 \pm 18.7 versus 51.0 \pm 18.7 mL; P < .05) and showed a trend to lower left ventricular ejection fraction.

CONCLUSIONS: This study is the first to show that in CF, severity of genotype (functional classification) is associated with cardiac impairment. Patients with severe CF genotype and cardiac dysfunction should be identified to evaluate cardiac response to gene-modifying treatments prior to consideration for lung transplantation.

KEYWORDS: Cystic fibrosis, echocardiography, genetics, heart failure

RECEIVED: January 7, 2018. ACCEPTED: July 20, 2018.

TYPE: Original Research

FUNDING: The author(s) received no financial support for the research, authorship, and/or publication of this article

DECLARATION OF CONFLICTING INTERESTS: The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

CORRESPONDING AUTHOR: John W Wilson, Cystic Fibrosis Service, Alfred Health and Central Clinical School, Monash University, Commercial Rd, Melbourne, VIC 3004, Australia. Email: john.wilson@monash.edu

Introduction

Cystic fibrosis (CF) is an autosomal, recessive disorder resulting from mutation of the cystic fibrosis trans-membrane conductance regulator (CFTR) gene, which causes a dysfunctional CFTR protein and apical cell-surface chloride ion channel. The consequences of this are multi-systemic and eventuate in well-characterised disease of the lung, exocrine pancreas, liver and sweat ducts.¹ Exercise capacity is impaired in CF, which is not solely explained by the reduced lung function and remains to be fully elucidated.^{2,3} Optimal exercise proficiency will be impaired by changes in lung function, cardiac function, intravascular volume or cell energetics.⁴ Exercise tolerance can be significantly improved by treatments that potentiate the function of CFTR.5

The identification of CFTR on human myocytes⁶ has prompted investigation of a possible CF-related cardiomyopathy. Using the echocardiographic technique of longitudinal strain measurement, subclinical impairment in left ventricular function has been reported in a CF population.^{7,8} This finding is supported by earlier studies, which demonstrated exercise

limitation related to a failure to augment cardiac output in patients with CF.9-11 It is unclear, however, whether abnormal cardiac function relates directly to pulmonary resistance, systemic sepsis, autonomic dysfunction or whether there is a direct effect of CFTR dysfunction on the cardiac myocyte.12

There are over 2000 identified mutations of the CFTR gene, of which 276 are known to cause CF.13 Mutations are classified based on the mechanism by which they disrupt CFTR function. Class 1, 2 and 3 mutations result from defective synthesis of full-length CFTR protein, defective CFTR protein processing and trafficking, and defective CFTR channel-gating, respectively. Classes 4 and 5 are a consequence of defective CFTR channel conductance or reduced synthesis/ stability of CFTR protein, respectively. Whereas class 1-3 mutations typically show almost complete loss of CFTR function and severe CF disease, classes 4 and 5 mutations retain some CFTR activity and tend to result in milder phenotypes with longer survival.^{14,15} There is therefore a wellestablished precedence for categorising CF severity and prognosis on the basis of 'low-risk' or 'high-risk' genotype.¹⁴ In this



Creative Commons Non Commercial CC BY-NC: This article is distributed under the terms of the Creative Commons Attribution-NonCommercial 4.0 License (http://www.creativecommons.org/licenses/by-nc/4.0/) which permits non-commercial use, reproduction and distribution of the work without further permission provided the original work is attributed as specified on the SAGE and Open Access pages (https://us.sagepub.com/en-us/nam/open-access-at-sage).

retrospective study in patients with CF, we hypothesised that patients with a high-risk genotype had greater cardiac dysfunction than those with a low-risk genotype based on echocardiographic findings.

Methods

Study design and patient population

This is a retrospective review of cardiac indices measured by transthoracic echocardiogram (TTE). Ethics approval was obtained from the institutional ethics committee at The Alfred Hospital (IEC46/18), and specific patient consent was not required due to the low risk nature of the study and collated de-identified data. Patients attended the Cystic Fibrosis Service, Alfred Health, Melbourne, Australia, and had a diagnosis of CF based on genotype and a positive sweat chloride. Patients with TTE performed as part of clinical care between 2000 and 2015 were included. Selection criteria were age≥18 years, no known history of cardiac disease or recent exacerbation at the time of echocardiography. The clinical record was reviewed from the time of TTE, and subjects were excluded if there was any clinical change or if physiologic support was required (oxygen supplementation, ventilation, fluid resuscitation or inotropic/vasopressor agents).

Collected data included demographic details, genotype, FEV_1 , laboratory data and echocardiographic indices of cardiac function. Laboratory results performed within 60 days and closest temporally to the date of TTE were recorded. The most recent FEV_1 to the date of echocardiography was also recorded.

Subjects were classified into 2 genotype groups on the basis of functional class of identified CFTR allele mutations. Patients were considered to have a severe genotype if they had 2 CFTR mutations from classes 1-3. Patients with only one or no class 1-3 mutations were considered to have a mild phenotype. When a second allele mutation was not identified, patients were only included in the analysis if they were documented to be pancreatic sufficient (suggestive of a milder clinical phenotype) at transition to adult CF care.

Echocardiography measurements

Echocardiographic studies were performed (iE33; Philips Healthcare, North Ryde, New South Wales, Australia) with subjects examined in a semi-supine, left-lateral position. The electrocardiogram was recorded continuously. Global left ventricular function was assessed from an apical four-chamber view by measuring left ventricular ejection fraction using the modified monoplane Simpson's rule.

Left ventricular end-diastolic diameter (LVDD), left ventricular end-systolic diameter (LVSD), right ventricular enddiastolic diameter, and right ventricular anterior wall thickness were measured from the M mode from a left parasternal view. Left ventricular fractional shortening (FS) was calculated from the formula (LVDD – LVSD/LVDD) X 100. Left atrial area (LA area) and right atrial area were measured from the apical four-chamber view. LA volume (LA volume) was calculated using the biplane area-length method and normalised to body surface area. The mitral Doppler signal was recorded in the apical four-chamber view, with the Doppler sample volume placed at the tip of the mitral valve.

The peak velocities of early (E) and late (A) filling waves and early/late filling ratios of peak velocities (E/A) were measured on the basis of trans-mitral flow velocities. Pulmonary systolic artery pressure was estimated from continuous-wave Doppler of tricuspid regurgitation using the Bernoulli equation. Tricuspid regurgitation velocity was recorded from the apical view and the parasternal short-axis view. Right atrial pressure (RAP) was estimated by assessing the diameter of the inferior vena cava and the percentage decrease in its diameter during inspiration. Tricuspid Annular Plane Systolic Excursion (TAPSE) was recorded on M-mode using the apical four-chamber view with the cursor is placed at the junction of the tricuspid valve plan with the free wall of the right ventricle.

Statistical analysis

Statistical analysis was performed using GraphPad InStat version 3.10 for Windows (GraphPad Software, San Diego, California, USA). Mean and standard deviation were calculated for comparison of demographic and laboratory data. Comparisons of echocardiographic indices of cardiac function were made between patients with severe and mild CFTR genotype configurations using the student T-test. Pearson's correlation coefficients were calculated to assess associations between indices of left ventricular function and recorded variables.

Results

Between the years 2000 and 2015, a TTE was performed in 100 genotyped patients (Table 1). The study was performed to exclude pulmonary hypertension (n=17), valve dysfunction (n=20), endocarditis (n=6), and causes for arrhythmias (n=7), neurological events (n=10) or as part of routine lung transplant workup (n=40). Patients were 31.5 ± 11.2 years old and had moderately severe lung disease (FEV₁ 47.3 ± 23.4% predicted) but without pulmonary hypertension (RVSP 36.1 ± 12.4 mmHg). Mean left ventricular ejection fraction (LVEF) was low to normal ($61.7 \pm 7.4\%$).

One hundred patients had at least one CF-causing CFTR mutation identified and were studied in the primary analysis by genotype severity (Table 1). Seventy-nine patients had mutations of both alleles from the CFTR classification groups 1-3 (severe genotype), and 21 had a single allelic mutation (milder genotype). Of the 21 patients with mild genotype, 13 did not have a second CFTR mutation identified. Patients with a severe genotype were younger than those with a mild genotype (31.1 \pm 10.0, 44.0 \pm 16.8 years; P < .0001).

	TOTAL COHORT	MILD GENOTYPE	SEVERE GENOTYPE	P VALUE	NORMATIVE VALUES
Ν	100	21	79	_	
Female, n (%)	52 (52)	11 (52.4)	41 (51.9)		
Age	33.8 (12.8)	44.0 (16.8)	31.1 (10.0)	<.0001	
BSA (m²)	1.68 (0.19)	1.73 (0.27)	1.66 (0.16)	.16	
SBP	120 (21.1)	127 (15.9)	118 (22.1)	.309	
DBP	72.9 (13.0)	78.5 (15.8)	71.5 (12.2)	.176	
HR	86.6 (15.7)	81.2 (15.7)	88.0 (15.6)	.251	
FEV_1 (% predicted)	46.8 (23.3)	51.8 (25.1)	45.2 (22.6)	.26	
LVDD (mm)	45.6 (5.2)	45.8 (6.0)	45.5 (5.0)	.84	50.2 (4.1)
LVSD (mm)	30.0 (4.4)	28.9 (3.9)	30.3 (4.5)	.18	32.4 (3.7)
FS (%)	34.1 (6.6)	36.9 (6.3)	33.3 (6.6)	.03	25-43
LVEF (%)	61.1 (6.1)	63.5 (4.9)	60.5 (6.3)	.15	62.0 (2.5)
LA area (cm ²)	15.6 (3.9)	18.0 (4.2)	14.9 (3.6)	.002	≤20
LA volume (mL)	42.3 (19.1)	51.0 (18.7)	39.9 (18.7)	.03	18 – 58
E/A	1.43 (0.56)	1.39 (0.76)	1.45 (0.49)	.66	1.4 (0.3)
RVSP (mmHg)	33.5 (9.4)	35.8 (9.4)	32.9 (9.4)	.33	28.9 (7.1)
TAPSE (cm)	1.91 (0.36)	2.09 (0.38)	1.86 (0.33)	.02	2.4 (0.35)
RAP (mmHg)	5.79 (3.33)	4.84 (3.17)	6.01 (3.36)	.17	0 – 5

Table 1. Patient characteristics and echocardiographic indices of cardiac function for total cohort and by CFTR genotype severity.

BSA: body surface area; FEV₁: forced expiratory volume in 1 second; LVDD: left ventricular diastolic diameter; LVSD: left ventricular systolic diameter; FS: left ventricle fractional shortening; LVEF: left ventricular ejection fraction; LA: left atria; E/A: mitral ratio of peak early to late diastolic filling velocity; RVSP: right ventricular filling pressure; TAPSE: tricuspid annular plane systolic excursion; RAP: right atrial pressure.

Data are presented as mean (\pm *SD*) unless otherwise stated. Normal values are presented as either mean (\pm *SD*), normal range or lower limit of normal.^{16–19} *P* values are given for comparisons between mild and severe genotype groups.

No significant difference was observed between patients with severe and mild genotype in right ventricular systolic pressure (RVSP) (32.9 ± 9.4 mmHg versus 35.8 ± 9.4 mmHg; P=.33) or FEV₁ ($45.2 \pm 22.6\%$ versus $51.8 \pm 25.1\%$ predicted; P=.26). There was a wide range of FEV₁ in both groups; however, other factors such as effective airway clearance, medication compliance and influence of the unknown gene can affect FEV₁ performance. The severe genotype group had a significantly lower FS (33.3 ± 6.6 versus 36.9 ± 6.3 %; P < .05) and a lower but non-significant LVEF (60.5 ± 6.3 versus $63.5 \pm 4.9\%$; P=.15). They also had a lower LA area (14.9 ± 3.6 versus 18.0 ± 4.2 cm²; P < .01) and LA volume (39.9 ± 18.7 versus 51.0 ± 18.7 mL; P < .05).

Univariate analysis using Pearson's correlation identified a linear association between FS and TAPSE (r=0.2726, P<.05); however, there was no association between FS and LA area, LA volume or FEV₁ (Table 2).

Discussion

In this study, we found that CF patients with a severe genotype had a lower left ventricular fractional shortening and TAPSE

than their counterparts with a mild genotype. This finding was independent of lung function severity based on ppFEV1 or nutritional changes; however, the specific mechanism by which classification 1-3 CFTR mutations might effect change in cardiac function was not part of this study.

Previous studies show left atrial dimensions reflect volume status,²⁰ and here, the lower LA area and LA volume in the severe genotype group favours a hypovolaemic state, previously reported in CF,²¹ causing a decrease in preload and reducing FS. The trend towards a negative association between heart rate and FS also supports decreased preload as a potential mechanism. Systolic and diastolic blood pressure (BP) between both groups did not differ implying afterload changes were either not influenced by genotype differences or compensatory mechanisms were able to moderate their impact at rest. Potential confounders for altered FS might be myocardial fibrosis, although no regional wall changes were seen on Echo, while autonomic dysfunction in CF is not common.¹² TAPSE was also lower (although within normal range) in the severe group identifying early RV changes, while LVEF did not differ between groups. Huang et al²² have recently showed that single

Table 2. Univariate analysis of fractional shortenin	g.
--	----

VARIABLE	FRACTIONAL SHORTENING		
	CORRELATION COEFFICIENT	P VALUE	
TAPSE (cm)	0.273	.012	
RVSP (mmHg)	-0.181	.07	
HR	-0.167	.08	
Iron	-0.141	.15	
Thyroxine	-0.1	.4	
LA area (cm ²)	-0.065	.41	
Age	0.05	.53	
LA volume (mL)	-0.047	.62	
FEV1 (% predicted)	0.037	.73	
RAP (mmHg)	0.019	.81	
BSA (m²)	0.014	.86	

BSA: body surface area; FEV₁: forced expiratory volume in 1 second; LA: left atria; RVSP: right ventricular filling pressure; TAPSE: tricuspid annular plane systolic excursion; RAP: right atrial pressure; HR: heart rate.

plane assessment was more sensitive than volumetric data in an intensive care unit (ICU) setting where volume changes may affect measurement.

Normal CFTR function is necessary for optimal cardiac function. Research has shown that in failing hearts, the CFTR adenosine triphosphate (ATP) binding cassettes are almost completely absent.²³ Murine studies describe reduced cardiac contraction by CFTR inhibition and blunting of the effect with stimulation of calcium-activated chloride channels.²⁴ Impaired CFTR activity on cardiac myocytes in patients with CF might directly result in subclinical cardiac dysfunction either through a direct myocardial impairment or a remodelling effect after injury.²⁵ Poor expression of mature CFTR protein has been reported in heart failure patients without CF in comparison to control subjects suggesting CFTR deficiency has a role not only in depolarisation but also contractility.²⁴ One potential explanation for the role of CFTR might be that in response to strain (failing respiratory function, or cardiac ischaemia) normally functioning chloride channels assist in repair of injured myocardium. Genetically or functionally (antibiotics, smoking) hindered CFTR may result in myocardial dysfunction.

The use of strain rate echocardiography in CF patients has unmasked subclinical LV dysfunction not present on resting techniques.^{7,8} These findings have also been supported by work using equilibrium radionuclide cineangiography.⁹ Exercise also exposes CF cardiac dysfunction; using an acetylene rebreathe technique during cardio pulmonary exercise testing (CPET), Van Iterson et al¹¹ showed impaired cardiac function contributed to CF exercise limitation. Adrenergic stimulation itself may also induce dysfunction in primed CFTR-defective myocardium, with evidence that albuterol (selective B2-agonist) impaired cardiac output (6.7 + / -0.5 vs 9.1 + / -0.3 mL/min/m2) compared to healthy individuals.²⁶

The difference in FS and TAPSE between high-risk and low-risk CF genotypes indicates a change in myocardial function as identified in the literature;²⁷ however, the variability of this effect based on genotype has not been identified. We know that severe genotypes are associated with less CFTR activity than milder genotypes and have reduced survival.¹⁴ A recent retrospective analysis of the United States' Cystic Fibrosis Foundation (CFF) patient registry (25,753 patients) found sweat chloride concentration at CF diagnosis predicted longterm survival, due mainly to the differences in CFTR genotype. Patients with residual CFTR function (classification 4-5) had better survival compared with CFTR genotypes with severely reduced CFTR function (classification 1-3).¹⁴

There are some limitations to the current study. This is a retrospective study, and cardiorespiratory stability is difficult to assess accurately as this relied on review of physiological variables in the patient's medical record, while identifying contemporaneous FEV_1 with echocardiography is also limited by this design. However, these values are assessed regularly in the routine care of adult CF patients limiting the time lag. In 13 subjects, the second CFTR mutation was unidentified, it is unlikely but possible that the second mutation was a class 1-3 mutation in these 13 patients which may influence the findings. We have minimised this limitation in these subjects by only including those with pancreatic sufficiency at transition to adult CF care in the mild genotype group.

There is a growing appreciation that CF is associated with subclinical ventricular dysfunction. In this study, we found a greater compromise in cardiac function associated with severe CF genotypes versus mild genotypes, while severity of lung disease was unrelated. Recently, CFTR potentiator and corrector treatment has improved patient lung function and quality of life; however, the impact on cardiac disease in CF is unknown. Further clinical trials are needed to delineate the impact of cardiac CFTR dysfunction on CF patients and if CFTR modulators might improve cardiac function in CF patients with severe genotypes.

Authors' Note

The authors M.P., D.T.K., D.M.K., T.K., and J.W. have no financial relationships or conflicts of interest to declare with respect to this manuscript.

ORCID iD

Dominic Keating (D https://orcid.org/0000-0001-6150-6927

REFERENCES

- 1. Elborn JS. Cystic fibrosis. Lancet. 2016;388:2519-2531.
- Rand S, Prasad SA. Exercise as part of a cystic fibrosis therapeutic routine. Expert Rev Respir Med. 2012;6:341–351; quiz 352.
- Williams CA, Saynor ZL, Tomlinson OW, Barker AR. Cystic fibrosis and physiological responses to exercise. *Expert Rev Respir Med.* 2014;8:751–762.

- Wasserman K. Diagnosing cardiovascular and lung pathophysiology from exercise gas exchange. *Chest.* 1997;112:1091–1101.
- Edgeworth D, Keating D, Ellis M, et al. Improvement in exercise duration, lung function and well-being in G551D-cystic fibrosis patients: a double-blind, placebo-controlled, randomized, cross-over study with ivacaftor treatment. *Clin Sci* (*Lond*). 2017;131:2037–2045.
- Warth JD, Collier ML, Hart P, et al. CFTR chloride channels in human and simian heart. *Cardiovasc Res.* 1996;31:615–624.
- Labombarda F, Pellissier A, Ellafi M, et al. Myocardial strain assessment in cystic fibrosis. JAm Soc Echocardiogr. 2011;24:1037–1045.
- Sellers ZM, McGlocklin L, Brasch A. Strain rate echocardiography uncovers subclinical left ventricular dysfunction in cystic fibrosis. J Cyst Fibros. 2015;14:654–660.
- Benson LN, Newth CJ, DeSouza M, et al. Radionuclide assessment of right and left ventricular function during bicycle exercise in young patients with cystic fibrosis. *Am Rev Respir Dis.* 1984;130:987–992.
- Ionescu AA, Ionescu AA, Payne N, Obieta-Fresnedo I, Fraser AG, Shale DJ. Subclinical right ventricular dysfunction in cystic fibrosis. A study using tissue Doppler echocardiography. *Am J Respir Crit Care Med.* 2001;163:1212–1218.
- Van Iterson EH, Wheatley CM, Baker SE, Morgan WJ, Snyder EM. The relationship between cardiac hemodynamics and exercise tolerance in cystic fibrosis. *Heart Lung*. 2016;45:283–290.
- Szollosi I, King SJ, Wilson JW, Naughton MT. Tachycardia in adults with cystic fibrosis is associated with normal autonomic function. *Intern Med J.* 2011;41:455–461.
- US CF Foundation, JHU, The Hospital for Sick Children. The clinical and functional translation of CFTR (CFTR2). http://www.cftr2.org. Accessed November 14, 2016.
- McKone EF, Goss CH, Aitken ML. CFTR genotype as a predictor of prognosis in cystic fibrosis. *Chest.* 2006;130:1441–1447.
- McKone EF, Emerson SS, Edwards KL, Aitken ML. Effect of genotype on phenotype and mortality in cystic fibrosis: a retrospective cohort study. *Lancet*. 2003;361:1671–1676.
- Lang RM, Bierig M, Devereux RB, et al. Recommendations for chamber quantification. Eur J Echocardiogr. 2006;7:79–108.

- Lang RM, Badano LP, Mor-Avi V, et al. Recommendations for cardiac chamber quantification by echocardiography in adults: an update from the American Society of Echocardiography and the European Association of Cardiovascular Imaging. JAm Soc Echocardiogr. 2015;28:1–39.e14.
- Kou S, Caballero L, Dulgheru R, et al. Echocardiographic reference ranges for normal cardiac chamber size: results from the NORRE study. *Eur Heart J Cardiovasc Imaging*. 2014;15:680–690.
- Armstrong DW, Tsimiklis G, Matangi MF. Factors influencing the echocardiographic estimate of right ventricular systolic pressure in normal patients and clinically relevant ranges according to age. *Can J Cardiol.* 2010;26: e35–e39.
- Agarwal R, Bouldin JM, Light RP, Garg A. Inferior vena cava diameter and left atrial diameter measure volume but not dry weight. *Clin J Am Soc Nephrol.* 2011;6:1066-1072.
- Holland AE, Wilson JW, Kotsimbos TC, Naughton MT. Metabolic alkalosis contributes to acute hypercapnic respiratory failure in adult cystic fibrosis. *Chest.* 2003;124:490–493.
- Huang SJ, Ting I, Huang AM, Slama M, McLean AS. Longitudinal wall fractional shortening: an M-mode index based on mitral annular plane systolic excursion (MAPSE) that correlates and predicts left ventricular longitudinal strain (LVLS) in intensive care patients. *Crit Care.* 2017;21:292.
- Solbach TF, Paulus B, Weyand M, Eschenhagen T, Zolk O, Fromm MF. ATPbinding cassette transporters in human heart failure. *Naunyn Schmiedebergs Arch Pharmacol.* 2008;377:231–243.
- 24. Sellers ZM, De Arcangelis V, Xiang Y, Best PM. Cardiomyocytes with disrupted CFTR function require CaMKII and Ca(2+)-activated Cl(-) channel activity to maintain contraction rate. *J Physiol*. 2010;588:2417–2429.
- Jiang K, Jiao S, Vitko M, et al. The impact of cystic fibrosis transmembrane regulator disruption on cardiac function and stress response. J Cyst Fibros. 2016;15:34–42.
- 26. Van Iterson EH, Karpen SR, Baker SE, Wheatley CM, Morgan WJ, Snyder EM. Impaired cardiac and peripheral hemodynamic responses to inhaled β_2 -agonist in cystic fibrosis. *Respir Res.* 2015;16:103.
- Sayyid ZN, Sellers ZM. Technological advances shed light on left ventricular cardiac disturbances in cystic fibrosis. J Cyst Fibros. 2017;16:454–464.