



ORIGINAL ARTICLE

Progression of early structural lung disease in young children with cystic fibrosis assessed using CT

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See Editorial, p 471

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ABSTRACT

Background Cross-sectional studies implicate neutrophilic inflammation and pulmonary infection as risk factors for early structural lung disease in infants and young children with cystic fibrosis (CF). However, the longitudinal progression in a newborn screened population has not been investigated.

Aim To determine whether early CF structural lung disease persists and progresses over 1 year and to identify factors associated with radiological persistence and progression.

Methods 143 children aged 0.2–6.5 years with CF from a newborn screened population contributed 444 limited slice annual chest CT scans for analysis that were scored for bronchiectasis and air trapping and analysed as paired scans 1 year apart. Logistic and linear regression models, using generalised estimating equations to account for multiple measures, determined associations between persistence and progression over 1 year and age, sex, severe cystic fibrosis transmembrane regulator (CFTR) genotype, pancreatic sufficiency, current respiratory symptoms, and neutrophilic inflammation and infection measured by bronchoalveolar lavage.

Results Once detected, bronchiectasis persisted in 98/133 paired scans (74%) and air trapping in 178/220 (81%). The extent of bronchiectasis increased in 139/227 (63%) of paired scans and air trapping in 121/264 (47%). Radiological progression of bronchiectasis and air trapping was associated with severe CFTR genotype, worsening neutrophilic inflammation and pulmonary infection.

Discussion CT-detected structural lung disease identified in infants and young children with CF persists and progresses over 1 year in most cases, with deteriorating structural lung disease associated with worsening inflammation and pulmonary infection. Early intervention is required to prevent or arrest the progression of structural lung disease in young children with CF.

INTRODUCTION

Studies using limited slice high resolution CT (HRCT) scans have demonstrated that infants and young children with cystic fibrosis (CF) diagnosed clinically have dilated airways compared with control infants,¹ with bronchiectasis detected in 30% and air trapping in 45%.² CT-detected changes have been reported in infants without clinically apparent lung disease,³ suggesting that CT-detected structural lung abnormalities may be an early and

Key messages

What is the key question?

- Does early cystic fibrosis (CF) structural lung disease detected by CT persist?

What is the bottom line?

- CT-detected structural lung disease persists and progresses in the majority of young children with CF and is associated with neutrophilic inflammation and pulmonary infection.

Why read on?

- These observations represent the first opportunity to track early CF lung disease using CT. Most children in this study did not have clinically apparent lung disease but had progressive structural lung changes that were associated with pulmonary inflammation and infection. These data support the concept of early intervention to prevent or arrest progressive lung disease.

sensitive marker of the disease status. Following detection by newborn screening, infants with CF at a median age of 3 months have significant lung changes detectable by limited slice HRCT with 19% of infants showing bronchial dilatation and 67% air trapping.⁴ A cross-sectional analysis of children aged from 2 months to 6 years demonstrated that prevalence and extent of CT-detected structural changes increases during the preschool years.⁵ Neutrophilic inflammation and pulmonary infection, particularly *Pseudomonas aeruginosa*, are implicated as contributors to structural lung disease.^{4,5}

Improvement in CT-detected indicators of structural lung disease has been observed following treatment of pulmonary exacerbations in children with CF younger than 4 years of age,⁶ and children with non-CF bronchiectasis may demonstrate spontaneous improvement of CT abnormalities,⁷ suggesting that while bronchiectasis is commonly detected in infants and preschool children with CF, a proportion of these children might have reversible airway dilatation. Chest CT has been proposed as an endpoint for clinical trials in young children with CF, therefore longitudinal changes in CT appearance need to be fully understood, particularly changes that occur over the likely duration of early interventional studies.

We hypothesised that early structural lung disease would not be reversible and that persistence and progression of CT-detected abnormalities would be associated with airway inflammation and pulmonary infection. We tested these hypotheses by analysing CT scans acquired 1 year apart in children up to 6 years of age from a newborn screened population and determining associations between persistence and progression of structural changes with simultaneously acquired clinical data including cystic fibrosis transmembrane regulator (CFTR) genotype and symptoms, and bronchoalveolar lavage (BAL)-detected pulmonary infection and inflammation.

METHODS

Study population

This retrospective analysis included 143 subjects born between 2000 and 2008, diagnosed with CF following newborn screening and enrolled in the Australian Respiratory Early Surveillance Team for Cystic Fibrosis (AREST CF) early surveillance program (ESP) in Perth and Melbourne, Australia. All aspects of the ESP have been approved by the human research ethics committees at each centre. Parents provided written consent at each assessment. Clinic participation rates for the AREST CF ESP were over 90%, with 85% of study children undergoing annual CT and bronchoscopy with BAL for the duration of the ESP. Pancreatic sufficiency was considered as a single assessment of faecal pancreatic elastase levels at diagnosis, if available, or by requirement for pancreatic enzyme replacement therapy for children transferred to the clinic from other centres. CFTR genotype was classified as 'severe' if both mutations were class I, II or III, or 'mild' CFTR genotype if at least one mutation was from class IV or V, based on previously published classifications.^{8–14} The presence of current respiratory symptoms was determined as a dichotomous variable by history and examination by the treating physician on the day of the scan, guided in part by the Kanga score.¹⁵

Early surveillance program

The ESP commences soon after diagnosis (median age 3.6 months) with annual CT and bronchoscopy with BAL up to 6 years. The AREST CF ESP was introduced into the Perth clinic in 1999 and into the Melbourne clinic in 2005 and was made available to all children under the age of 6 years in the clinics at that time.

Chest CT

Chest CT scanning was introduced in 2005 and data used in the present study were acquired between 2005 and 2010. The median age at the first ever CT scan was 1.0 year (25th–75th percentile 0.3–2.6 years, range 0.2–5.4 years). Each child in this study had at least two assessments a median of 12 months apart (25th–75th percentile 11–13 months, range 6–18 months) and all available pairs of chest CTs were included in the analysis. Chest CT and BAL were performed as previously described.^{4–5} Briefly, children were anaesthetised and intubated using a cuffed tube, with a recruitment manoeuvre to reduce procedure related atelectasis.^{4–5} A volume controlled limited slice scan was performed with three slices taken at end-inspiration (25 cm H₂O) and end-expiration (0 cm H₂O) as previously described,^{4–5} with lung volume controlled by the anaesthetist, using a Philips Brilliance 64 (Philips Medical Systems, Eindhoven, The Netherlands) CT scanner in Perth and a Somatron Sensation 16 (Siemens Medical Solutions, Erlangen, Germany) CT scanner in Melbourne. The CT settings used for this limited slice technique are described in the online supplement.

BAL-detected infection

BAL was performed immediately following CT, with three aliquots of 1 ml/kg normal saline instilled into and aspirated from the right middle lobe and one aliquot into the lingula or worst affected lobe identified on CT. The first recovered aliquot from each lobe was sent for microbiological processing to detect bacterial and fungal infections (while viral infections were detected using immunofluorescence or viral culture, they were uncommon (eight infections in nine children) and therefore not included in analyses). BAL samples were considered 'infected' if there was a positive culture for at least one predominant organism at any density, 'mixed oral flora' if mixed oral flora only were cultured without a predominant organism, or 'uninfected' with a negative culture. Details about specific infection with *P. aeruginosa* are given in the online supplement.

BAL-detected inflammatory analyses

The second and third aliquots from the right middle lobe were pooled and processed for inflammatory analyses as previously described,^{4–5–16} including total and differential cell count, interleukin 8 (IL-8) level measured using ELISA and detection of free neutrophil elastase (NE) activity. The lower limit of detection for IL-8 and NE are 100 pg/ml and 200 ng/ml, respectively. A value half the lower limit of detection was assigned to samples below the limit of detection for statistical analyses.

CT analysis

Images were scored in batches of 5–20 as they became available by a single scorer (CPM) as previously published^{4–5} on a soft copy reporting station (Agfa Impax TM, Agfa-Gevaert, Mortsel, Belgium) using standard lung window protocols. While the scorer was aware of the diagnosis of CF, scans were assessed without knowledge of the outcome of any previous scans, clinical status, microbiology or inflammatory results. Each scan was divided into six zones (left and right upper, middle and lower zones), corresponding with each axial slice. Each zone was scored for the presence and extent of bronchiectasis and air trapping using a previously reported modification of a CF specific CT scoring system.^{5–17} Briefly, bronchiectasis was scored on inspiratory images, defined as a bronchus-to-artery ratio >1, or non-tapering bronchus visible in transverse plane. Air trapping was scored on expiratory images, defined as geographical regions of reduced density. Extent in each zone was defined as 0=absent, 1=affecting ≤50% of airways (bronchiectasis) or lung field (air trapping), and 2=affecting >50%. Extent ranged from 0 (absent) to 12 (affecting >50% in all zones). Intra-observer reliability has been previously reported.⁵ Zones were excluded from analysis if anaesthetic related atelectasis obscured airways in >one-third of the visible segments. If >2 zones were excluded from analysis, the scan was excluded from the study. Demographics and summary data from initial ('t-1') and subsequent ('t') assessments are presented in table 1.

Statistical analysis

Each child contributed up to five scans for analysis (49 children contributed two scans, 46 contributed three scans, 32 contributed four scans and 16 contributed five scans). Data were analysed as scan pairs, with bronchiectasis and air trapping during the scan at time *t* ('subsequent scan') as the outcome of interest, in comparison with the scan at *t*-1 ('initial scan'), obtained at a median of 1 year prior to time *t*. Each scan could be represented as both the scan at time *t* in one pair and as the scan at *t*-1 in a subsequent pair. Variables that were not normally distributed had a natural logarithmic transformation applied. As

Table 1 Demographics of the population and summary data (n=143 children)

	Initial scan (t-1) (n=301)	Subsequent scan (t) (n=301)
Demographics		
Age (years)	2.00 (0.97–3.24)	3.01 (1.98–4.25)
Sex (% female)	51% (73/143)	
Severe CFTR genotype (%)*	90% (114/126)	
Pancreatic sufficiency (%)	16% (23/143)	
Respiratory symptoms (%)	20% (59/301)	22% (63/291)
Pulmonary inflammation		
Inflammatory response score	-0.18 (-1.03–0.84)	-0.05 (-1.04–0.87)
Total cell count ($\times 10^3$ /ml)	293 (13–490)	300 (176–476)
Neutrophil count ($\times 10^3$ /ml)	46 (17–139)	59 (16–160)
Per cent neutrophils (%)	19% (9–37)	22% (9–42)
IL-8 level (pg/ml)	740 (220–2040)	1050 (340–2260)
Detectable neutrophil elastase	29% (86/298)	31% (94/300)
Pulmonary infection		
Uninfected (%)	33% (99/298)	30% (91/300)
Mixed oral flora only (%)	30% (90/298)	25% (76/300)
Infected (%)†	37% (109/298)	44% (133/300)
Bacterial infection	32% (95/298)‡	38% (115/300)§
Fungal infection	10% (30/298)¶	15% (46/300)**
Structural lung disease		
Detectable bronchiectasis (%)	44% (133/301)	62% (187/301)
Extent of bronchiectasis	0 (0–2)	1 (0–3)
Detectable air trapping (%)	73% (220/301)	74% (222/301)
Extent of air trapping	2 (0–5)	2 (0–6)

Data represented as median (25th–75th percentile) or percentage (absolute numbers) as appropriate.

*Complete data for CFTR genotype are not available for all subjects as both genes have not been identified in all children and some CFTR mutations have unknown effects on CFTR function.

†Some infections included both bacterial and fungal organisms.

‡Including 26 infections with *Pseudomonas aeruginosa*, 26 infections with *Staphylococcus aureus*, 21 infections with *Haemophilus influenzae*, eight infections with *Streptococcus pneumoniae* and 33 infections with other bacteria.

§Including 30 infections with *P aeruginosa*, 40 infections with *S aureus*, 32 infections with *H influenzae*, 11 infections with *S pneumoniae* and 39 infections with other bacteria.

¶Including 19 infections with *Aspergillus* species, seven infections with *Candida* and seven infections with other fungi.

**Including 32 infections with *Aspergillus* species, 12 infections with *Candida* and 10 infections with other fungi.

CFTR, cystic fibrosis transmembrane regulator; IL-8, interleukin 8.

neutrophilic inflammatory variables (total cell count, neutrophil count, IL-8 level and NE activity) were highly correlated, they were summarised, guided by a principal component analysis, into a single 'inflammatory response score' as the marker of

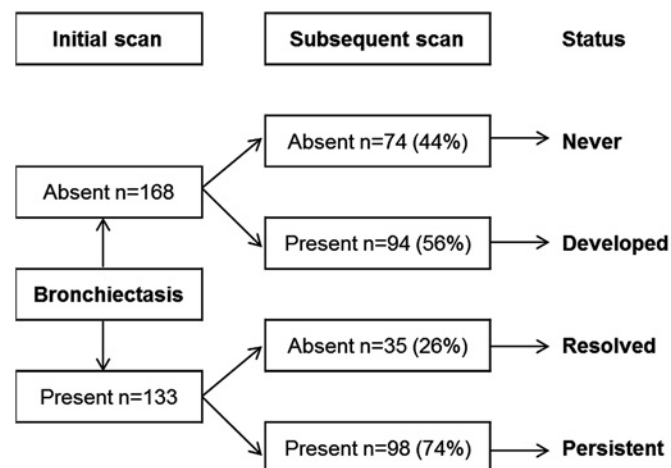
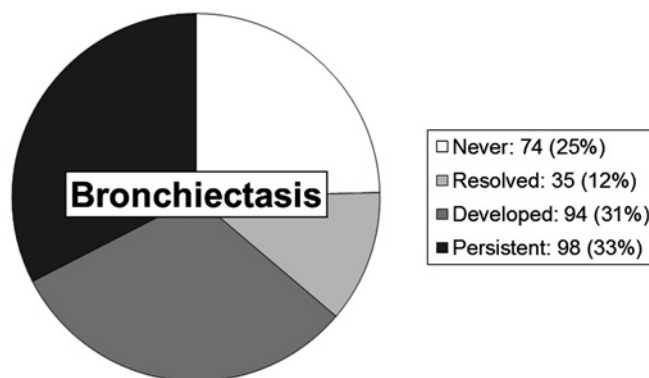
neutrophilic inflammation. Further details are given in the online supplement. Analysis of infection compared 'infected' and 'mixed oral flora' with 'uninfected'.

Logistic regression models were used to measure the associations between the persistence of structural lung disease and age, sex, severe CFTR genotype, pancreatic sufficiency, current respiratory symptoms, neutrophilic inflammation and pulmonary infection, assessed where appropriate at both time t and at time $t-1$. The response variables in the models were the presence of structural lung disease at time t excluding children who did not have bronchiectasis or air trapping present at time $t-1$. Age at time $t-1$ was included as a covariate in all analyses.

To examine the associations between the progression of structural lung disease and factors outlined above, linear regression models were used. The data for the extent of structural lung disease were positively skewed and therefore natural logarithmic transformations were applied to the data. The response variables in the models were the log-transformed extent of structural lung disease at time t . The models were adjusted for the extent of structural lung disease at time $t-1$, age at time $t-1$ and severe CFTR genotype and excluded children with no detectable bronchiectasis or air trapping at either time t or time $t-1$.

Further multivariate analyses were performed for both persistence and progression to examine the effects of changes in infection and inflammation status on the presence or change in extent of structural lung disease. The models were adjusted for the extent of structural lung disease at time $t-1$ (if the outcome was the extent of structural lung disease at time t), age at time $t-1$, severe CFTR genotype (for progression analyses), infection status at times $t-1$ and t , and inflammation status at times $t-1$ and t . The models were chosen to control for the confounding effects of age, severe CFTR genotype (for progression analyses), inflammation and infection.

Repeated measurements on the same child were allowed for in both persistence and progression analyses by using generalised estimating equations. This approach was selected due to the ease of interpretation of results as it focuses on marginal parameters. Similar results were obtained using other models to account for repeated measures. Data were analysed using Stata V.11.0 (Stata Corporation, College Station, Texas, USA). As we did not want the interpretation of our findings to be influenced by the number of tests performed, we have presented all analyses as previously suggested.¹⁸

**Figure 1** Bronchiectasis status at initial and subsequent scans.

RESULTS

Bronchiectasis

There were 27 (19%) children who never demonstrated bronchiectasis during the surveillance period of this study. Bronchiectasis was never present in 74 pairs (25%), and was detected at either the initial or subsequent scan in 227 pairs (75%) as shown in figure 1. Bronchiectasis was detected at the initial scan in 133 scan pairs. Of these 133 scan pairs, bronchiectasis persisted at the subsequent scan in 98 (74%) (median age 3.1, 25th–75th percentile 1.9–4.2, range 0.2–5.5 years), while 35 (26%) showed apparent resolution of bronchiectasis (median age 2.0, 25th–75th percentile 1.2–3.3, range 0.3–5.4 years). Associations between persistence of bronchiectasis and clinical factors, CT factors, inflammation and infection are shown in table 2.

Bronchiectasis was detected at either the initial scan or subsequent scan (or both) in 227 paired CT scans. Of these, the extent of bronchiectasis increased from the initial to the subsequent scan in 139 scan pairs (63%), with extent unchanged in 24 scan pairs (11%), and extent decreased in 59 scan pairs (27%). After excluding children who never had detectable bronchiectasis, the median extent of bronchiectasis increased from 1 (25th–75th percentile 0–3) at the initial scan to 2 (25th–75th percentile 1–4) at the subsequent scan. Associations between progression of bronchiectasis and clinical factors, CT factors, inflammation and infection are shown in table 3.

Air trapping

Air trapping was never present in 37 pairs (12%) and was detected at either the initial or subsequent scan in 264 pairs

(88%) as shown in figure 2. Air trapping was detected at the initial scan in 220 scan pairs. Of these 220 scan pairs, air trapping persisted at the subsequent scan in 178 pairs (81%) (median age 1.97, 25th–75th percentile 0.97–3.17, range 0.2–5.5 years), while 42 pairs (19%) (median age 2.24, 25th–75th percentile 1.24–3.44, range 0.3–5.4 years) showed apparent resolution of air trapping. Associations between persistence of air trapping and clinical factors, CT factors, inflammation and infection are shown in table 4.

Air trapping was detected at either the initial scan or subsequent scan (or both) in 264 (88%) paired CT scans. Of these, the extent of air trapping increased from the initial to the subsequent scan in 121 scan pairs (46%), with extent unchanged in 36 scan pairs (14%), and extent decreased in 107 scan pairs (41%). After excluding children who never had detectable air trapping, the median extent of air trapping at both the initial scan (25th–75th percentile 0–5) and the subsequent scan (25th–75th percentile 0–6) was 3. Associations between progression of air trapping and clinical factors, CT factors, inflammation and infection are shown in table 5.

DISCUSSION

This is the first longitudinal assessment of CT-detected structural lung disease in young children with CF detected by newborn screening. The observations provide a unique insight to the early development of CF lung disease and factors associated with the persistence and progression of structural changes. We found that neutrophilic inflammation and pulmonary infection were independently associated with progression of

Table 2 Associations with persistence of bronchiectasis

Clinical factors	Change in bronchiectasis status		Model 1*		Model 2†	
	Transient	Persistent	OR (95% CI)	p Value	OR (95% CI)	p Value
Age	2.0 (1.2–3.3)	3.1 (1.9–4.0)	1.23 (0.91 to 1.66)	0.169	1.30 (0.95 to 1.79)	0.101
Sex (female)	53% (19/35)	57% (56/98)	1.17 (0.53 to 2.57)	0.703	–	–
Severe CFTR genotype	88% (28/32)	96% (86/90)	3.03 (0.69 to 13.29)	0.142	–	–
Pancreatic sufficiency	20% (7/35)	10% (10/98)	0.51 (0.17 to 1.49)	0.219	–	–
Respiratory symptoms at the initial scan	20% (7/35)	30% (29/98)	1.61 (0.63 to 4.12)	0.320	–	–
Respiratory symptoms at the subsequent scan	9% (3/33)	29% (27/98)	3.86 (1.06 to 14.00)	0.040	–	–
CT factors						
Detectable air trapping at the initial scan	69% (24/35)	86% (84/98)	2.72 (1.08 to 6.86)	0.034	–	–
Air trapping extent at the initial scan	3 (0–6)	4 (2–9)	2.83 (1.22 to 6.54)	0.015	–	–
Detectable air trapping at the subsequent scan	57% (20/35)	80% (79/98)	1.18 (1.02 to 1.36)	0.029	–	–
Air trapping extent at the subsequent scan	1 (0–6)	4.5 (1–9)	1.24 (1.07 to 1.44)	0.004	–	–
Change in air trapping extent	0 (–1–2)	0 (–1–4)	1.21 (1.02 to 1.45)	0.032	–	–
Bronchiectasis extent at the initial scan	2 (1–6)	3 (1–7)	1.85 (0.73 to 4.68)	0.257	–	–
Bronchoalveolar lavage factors at the initial scan						
Uninfected	34% (12/35)	23% (22/97)	(Comparator)		(Comparator)	
Mixed oral flora	26% (9/35)	25% (24/97)	1.55 (0.54 to 4.44)	0.412	1.73 (0.50 to 5.96)	0.387
Infected	40% (14/35)	53% (51/97)	1.85 (0.73 to 4.68)	0.195	1.51 (0.46 to 5.00)	0.501
Detectable neutrophil elastase	38% (13/35)	42% (41/97)	1.16 (0.52 to 2.58)	0.725	–	–
Inflammatory response score	–0.01 (–0.80–0.88)	0.55 (–0.74–1.44)	1.17 (0.90 to 1.53)	0.238	0.99 (0.69 to 1.41)	0.949
Bronchoalveolar lavage factors at the subsequent scan						
Uninfected	40% (14/35)	22% (12/98)	(Comparator)		(Comparator)	
Mixed oral flora	20% (7/35)	20% (20/98)	4.33 (1.28 to 14.68)	0.018	3.34 (0.86 to 12.90)	0.080
Infected	40% (14/35)	67% (66/98)	6.23 (2.29 to 16.93)	<0.001	4.92 (1.51 to 16.00)	0.008
Detectable neutrophil elastase	23% (8/35)	49% (48/98)	3.22 (1.32 to 7.84)	0.010	–	–
Inflammatory response score	–0.20 (–1.36–0.69)	0.78 (–0.43–1.96)	1.43 (1.09 to 1.88)	0.010	1.30 (0.92 to 1.82)	0.134
Change in inflammatory response score	0.27 (–1.49–1.04)	0.35 (–0.86–1.33)	1.50 (1.09 to 2.04)	0.011	–	–

Data given as median (25th–75th percentile) or percentage (raw data) as appropriate.

–Not included in model. Bold, significant p-values (<0.05).

*Model 1: Each variable analysed in a separate model after adjusting for age at the initial scan.

†Model 2: Combined model adjusted for age at initial scan, infection status at the initial and subsequent scans and inflammation status at the initial and subsequent scans. CFTR, cystic fibrosis transmembrane regulator.

Table 3 Associations with progression of bronchiectasis

Clinical factors	Model 1*		Model 2†	
	Regression coefficient‡ (95% CI)	p Value	Regression coefficient‡ (95% CI)	p Value
Age	1.07 (1.00 to 1.14)	0.041	1.06 (1.01 to 1.12)	0.020
Sex (female)	1.08 (0.90 to 1.30)	0.408	—	—
Severe CFTR genotype	1.54 (1.19 to 1.99)	0.001	1.28 (1.04 to 1.59)	0.021
Pancreatic sufficiency	0.81 (0.63 to 1.05)	0.106	—	—
Respiratory symptoms at the initial scan	1.13 (0.88 to 1.45)	0.325	—	—
Respiratory symptoms at the subsequent scan	1.39 (1.16 to 1.67)	<0.001	—	—
CT factors				
Detectable air trapping at the initial scan	1.31 (1.09 to 1.58)	0.004	—	—
Air trapping extent at the initial scan	1.05 (1.01 to 1.09)	0.008	—	—
Detectable air trapping at the subsequent scan	1.41 (1.12 to 1.77)	0.004	—	—
Air trapping extent at the subsequent scan	1.09 (1.05 to 1.13)	<0.001	—	—
Change in air trapping extent	1.08 (1.05 to 1.12)	<0.001	—	—
Bronchoalveolar lavage factors at the initial scan				
Uninfected	(Comparator)		(Comparator)	
Mixed oral flora	0.91 (0.72 to 1.15)	0.439	0.85 (0.67 to 1.10)	0.218
Infected	1.16 (0.92 to 1.48)	0.212	1.00 (0.79 to 1.27)	0.973
Detectable neutrophil elastase	1.04 (0.84 to 1.30)	0.692	—	—
Inflammatory response score	1.07 (1.01 to 1.13)	0.016	1.02 (0.96 to 1.08)	0.508
Bronchoalveolar lavage factors at the subsequent scan				
Uninfected	(Comparator)		(Comparator)	
Mixed oral flora	1.35 (1.08 to 1.68)	0.008	1.23 (0.97 to 1.56)	0.091
Infected	1.57 (1.23 to 2.01)	<0.001	1.38 (1.07 to 1.79)	0.014
Detectable neutrophil elastase	1.44 (1.14 to 1.80)	0.002	—	—
Inflammatory response score	1.18 (1.10 to 1.26)	<0.001	1.14 (1.08 to 1.22)	<0.001
Change in inflammatory response score	1.18 (1.11 to 1.26)	<0.001	—	—

—Not included in model. Bold, significant p-values (<0.05).

*Model 1: Each variable analysed in a separate model after adjusting for age at the initial scan and severe CFTR genotype.

†Model 2: Combined model adjusted for age at initial scan, severe CFTR genotype, infection status at the initial and subsequent scans and inflammation status at the initial and subsequent scans.

‡Regression coefficients have been back-transformed from the logarithmic scale.

CFTR, cystic fibrosis transmembrane regulator.

bronchiectasis, with neutrophilic inflammation associated with progression of air trapping. Importantly, neutrophilic inflammation at the initial scan was associated with subsequent progression of structural lung disease and might therefore be an early predictor of significant future lung disease. These findings are in agreement with earlier cross-sectional studies that implicated neutrophilic inflammation and pulmonary infection as risk factors for structural lung disease.^{4, 5} We also identified an association between air trapping at the initial scan and subsequent persistence and progression of bronchiectasis, which may

offer some insights into the pathophysiology and natural history of different structural lung disease components. Further analyses are required to identify whether early air trapping can predict those children who will later develop bronchiectasis at an early age.

In addition to associations with neutrophilic inflammation and pulmonary infection, we identified an independent association between severe CFTR genotype and progression of bronchiectasis and air trapping. Previous studies have identified associations between mild CFTR genotypes and lower rates of

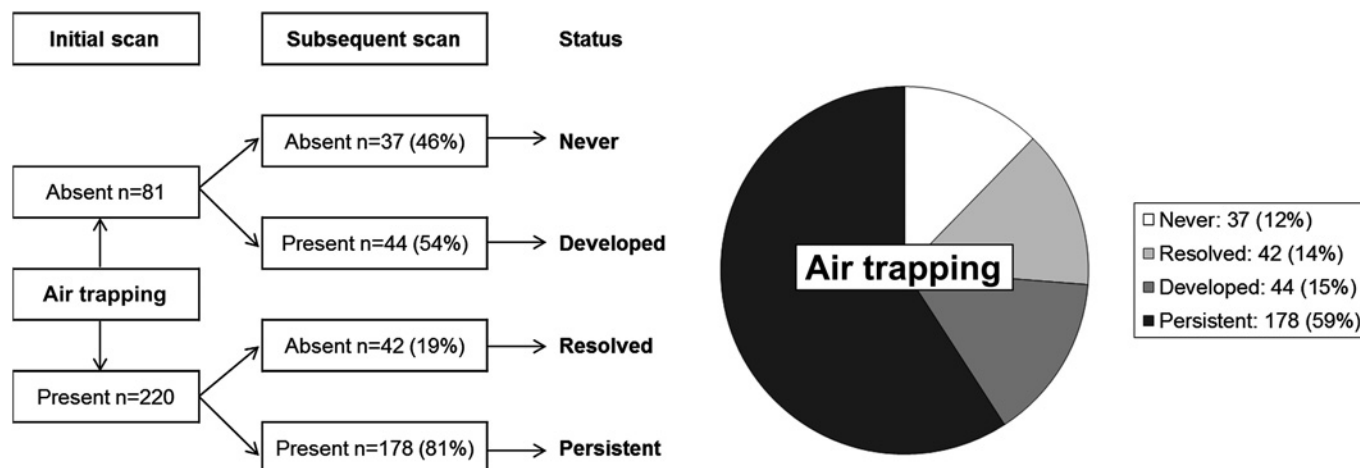


Figure 2 Air trapping status at initial and subsequent scans.

Table 4 Associations with persistence of air trapping

Clinical factors	Change in air trapping status		Model 1*		Model 2†	
	Transient	Persistent	OR (95% CI)	p Value	OR (95% CI)	p Value
Age	2.2 (1.2–3.4)	2.0 (1.0–3.2)	0.86 (0.70 to 1.07)	0.175	0.76 (0.59 to 0.97)	0.028
Sex (female)	50% (21/42)	54% (97/178)	1.17 (0.58 to 2.37)	0.657	–	–
Severe CFTR genotype	88% (29/33)	93% (160/172)	1.83 (0.51 to 6.57)	0.355	–	–
Pancreatic sufficiency	17% (7/42)	11% (19/178)	0.53 (0.20 to 1.42)	0.208	–	–
Respiratory symptoms at the initial scan	19% (8/42)	20% (36/178)	1.18 (0.51 to 2.74)	0.697	–	–
Respiratory symptoms at the subsequent scan	13% (5/40)	21% (36/172)	1.95 (0.73 to 5.17)	0.180	–	–
CT factors						
Detectable bronchiectasis at the initial scan	52% (22/42)	48% (86/178)	0.99 (0.49 to 2.02)	0.986	–	–
Bronchiectasis extent at the initial scan	1 (0–3)	0 (0–5)	1.19 (0.98 to 1.43)	0.072	–	–
Detectable bronchiectasis at the subsequent scan	60% (25/42)	68% (121/178)	1.69 (0.83 to 3.44)	0.152	–	–
Bronchiectasis extent at the subsequent scan	1 (0–4)	1 (0–7)	1.22 (1.04 to 1.42)	0.013	–	–
Change in bronchiectasis extent	0 (–1–4)	1 (0–4)	1.18 (1.00 to 1.40)	0.048	–	–
Air trapping extent at the initial scan	2 (1–5)	4 (2–7)	1.34 (1.13 to 1.60)	0.001	–	–
Bronchoalveolar lavage factors at the initial scan						
Uninfected	43% (18/42)	31% (55/177)	(Comparator)		(Comparator)	
Mixed oral flora	26% (11/42)	27% (47/177)	1.32 (0.57 to 3.06)	0.523	0.89 (0.35 to 2.30)	0.814
Infected	31% (13/42)	42% (75/177)	1.95 (0.87 to 4.37)	0.105	1.49 (0.57 to 3.92)	0.416
Detectable neutrophil elastase	22% (9/41)	34% (61/177)	1.75 (0.81 to 3.81)	0.156	–	–
Inflammatory response score	–0.43 (–1.04–0.74)	0.07 (–1.02–1.19)	1.20 (0.98 to 1.49)	0.083	1.02 (0.79 to 1.32)	0.885
Bronchoalveolar lavage factors at the subsequent scan						
Uninfected	40% (17/42)	25% (44/177)	(Comparator)		(Comparator)	
Mixed oral flora	26% (11/42)	22% (39/177)	1.39 (0.58 to 3.35)	0.461	1.33 (0.49 to 3.63)	0.572
Infected	33% (14/42)	53% (94/177)	2.83 (1.26 to 6.36)	0.012	2.54 (0.96 to 6.71)	0.060
Detectable neutrophil elastase	21% (9/42)	37% (66/178)	2.13 (0.97 to 4.67)	0.060	–	–
Inflammatory response score	–0.32 (–1.05–0.65)	0.47 (–0.63–1.33)	1.23 (0.98 to 1.53)	0.071	1.10 (0.85 to 1.44)	0.470
Change in inflammatory response score	0.11 (–1.05–1.17)	0.45 (–1.81–1.34)	1.19 (0.94 to 1.49)	0.150	–	–

–Not included in model Bold, significant p-values (<0.05).

Data given as median (25th–75th percentile) or percentage (raw data) as appropriate.

*Model 1: Each variable analysed in a separate model after adjusting for age at the initial scan.

†Model 2: Combined model adjusted for age at initial scan, infection status at the initial and subsequent scans and inflammation status at the initial and subsequent scans.

CFTR, cystic fibrosis transmembrane regulator.

infection with *P aeruginosa* and other pathogens,^{14 19} and chest radiography and lung function decline is faster in patients with severe CFTR mutations²⁰ with poorer lung function seen in those with the most severe mutations.²¹ However, close correlations between clinical pulmonary phenotypes and CFTR genotype have been elusive.¹³ To our knowledge, no other studies have examined the link between CFTR genotypes and structural lung disease so early in life in a newborn screened population. The causal pathways that link CFTR dysfunction with structural lung disease are poorly understood, but studies such as ours provide information regarding the earliest pathobiological events in the lung, and determining relations with structural changes are likely to be a key to defining these pathways. Further studies in larger populations are required to confirm this association.

There are some limitations to this study. To determine whether structural lung disease was common in infancy we initially chose a three-slice HRCT method as a screening tool using a low dose protocol, before more extensive imaging techniques with low radiation exposures had been adequately validated. In practice, our estimates of persistence and progression of bronchiectasis are likely to have been underestimated because limited slice scans underestimate this element of structural change,²² and this limitation is likely to have resulted in misclassification of apparently resolved bronchiectasis. Thus, using this radiological method, we cannot say with certainty whether early bronchiectasis truly resolved in any infants or preschool children, though it is interesting to note that of the 35 pairs that showed apparent resolution of bronchiectasis, 13

had further annual scan(s) and of these, 12 children (92%) demonstrated recurrence of bronchiectasis. Despite this, we demonstrated that children with apparent resolution of bronchiectasis were less likely to have worsening neutrophilic inflammation and infection at the subsequent scan. Scans were assessed by an experienced paediatric thoracic radiologist (CPM) as previously described.^{4 5} While we were unable to make the size and resolution of each scan identical and it was not possible to blind the scorer to the approximate age of the child, we do not believe that an age bias is likely because at the time of the assessment the scorer was unaware of whether the images represented the initial or subsequent scan of any given pair from children in a relatively wide age range. While the CT scans were available to treating clinicians and evidence of lung disease might lead to more aggressive clinical management, this would be expected to reduce the likelihood of demonstrating persistence and progression as we have done so in this study.

The results of this study have implications for the clinical management of young children with CF. We observed that neutrophilic inflammation at the initial scan was associated with subsequent progression of both bronchiectasis and air trapping. This observation suggests a possible role for anti-inflammatory therapies early in life. However, the effectiveness of anti-inflammatory agents for preventing, delaying or reversing structural lung disease in young children with CF is unknown and needs to be investigated in appropriately designed, prospective early intervention trials. Importantly, respiratory symptoms at the initial scan were not associated with increased rates of persistence or progression of

Table 5 Associations with progression of air trapping

Clinical factors	Model 1*		Model 2†	
	Regression coefficient‡ (95% CI)	p Value	Regression coefficient‡ (95% CI)	p Value
Age	0.97 (0.92 to 1.02)	0.209	0.92 (0.88 to 0.98)	0.004
Sex (female)	1.07 (0.91 to 1.27)	0.419	—	—
Severe CFTR genotype	1.40 (1.14 to 1.71)	0.001	1.40 (1.07 to 1.81)	0.012
Pancreatic sufficiency	0.87 (0.69 to 1.10)	0.240	—	—
Respiratory symptoms at the initial scan	1.00 (0.83 to 1.21)	0.983	—	—
Respiratory symptoms at the subsequent scan	1.18 (0.97 to 1.42)	0.091	—	—
CT factors				
Detectable bronchiectasis at the initial scan	1.05 (0.88 to 1.26)	0.575	—	—
Bronchiectasis extent at the initial scan	1.05 (1.01 to 1.09)	0.008	—	—
Detectable bronchiectasis at the subsequent scan	1.39 (1.16 to 1.65)	<0.001	—	—
Bronchiectasis extent at the subsequent scan	1.08 (1.05 to 1.11)	<0.001	—	—
Change in bronchiectasis extent	1.07 (1.04 to 1.10)	<0.001	—	—
Bronchoalveolar lavage factors at the initial scan				
Uninfected	(Comparator)		(Comparator)	
Mixed oral flora	0.95 (0.76 to 1.17)	0.609	0.84 (0.68 to 1.03)	0.098
Infected	1.05 (0.87 to 1.27)	0.613	0.86 (0.68 to 1.03)	0.171
Detectable neutrophil elastase	1.13 (0.96 to 1.33)	0.147	—	—
Inflammatory response score	1.08 (1.02 to 1.15)	0.005	1.06 (0.99 to 1.13)	0.124
Bronchoalveolar lavage factors at the subsequent scan				
Uninfected	(Comparator)		(Comparator)	
Mixed oral flora	1.09 (0.83 to 1.43)	0.548	1.06 (0.80 to 1.42)	0.671
Infected	1.32 (1.06 to 1.64)	0.014	1.15 (0.87 to 1.51)	0.321
Detectable neutrophil elastase	1.48 (1.26 to 1.75)	<0.001	—	—
Inflammatory response score	1.11 (1.05 to 1.18)	<0.001	1.09 (1.02 to 1.17)	0.017
Change in inflammatory response score	1.10 (1.03 to 1.17)	0.004	—	—

—Not included in model. Bold, significant p-values (<0.05).

*Model 1: Each variable analysed in a separate model after adjusting for age at the initial scan and severe CFTR genotype.

†Model 2: Combined model adjusted for age at initial scan, severe CFTR genotype, infection status at the initial and subsequent scans and inflammation status at the initial and subsequent scans.

‡Regression coefficients have been back-transformed from the logarithmic scale.

CFTR, cystic fibrosis transmembrane regulator.

bronchiectasis or air trapping, and therefore symptoms cannot be relied upon to identify children at risk of further deterioration in structural lung disease. While respiratory symptoms at the subsequent scan were associated with persistence and progression of bronchiectasis, the overall prevalence of symptoms was low in this cohort and most children with persistent or progressive disease did not have respiratory symptoms at the time of their subsequent scan. Associations between pulmonary infection at the second scan of a pair and structural lung disease emphasise the importance of early detection of infection since infection is an important determinant of inflammation²³ and eradication, particularly of *P aeruginosa*, is usually successful when detected early.²⁴

This study contributes to our understanding of CT-detected structural lung disease in young children and has implications for clinical trials in young children with CF. CT as an outcome measure for clinical trials for infants and pre-school children with CF shows great promise, particularly since traditional outcome measures used by regulatory agencies are not appropriate for young children.²⁵ The findings from this study suggest a low rate of spontaneous resolution of structural changes with evidence of disease progression over 1 year. Therefore, CT could provide direct measurements of CF lung disease for use in short term (1–3 years) intervention studies of potentially disease modifying therapies in infants and preschool children, groups currently excluded from early intervention studies due to the absence of acceptable trial endpoints. Further research using CT as an outcome measure should focus on the use of ultra-low dose volumetric CT scans to improve our understanding of lung disease in CF and to develop strategies to

optimise image quality, minimise radiation dose and determine the best method to quantify the extent of early CT detected structural lung disease in young children. While we acknowledge the small inherent risk associated with exposure to ionising radiation using chest CT, the limited slice approach used in this paper involves a very low radiation exposure of approximately 0.28–0.40 mSv as previously reported.^{4,5} Using the latest scanning techniques and software we are now able to obtain high quality low dose full inspiratory and expiratory volumetric scans that provide more complete information about structural lung disease, and research to determine the best methods for quantification of structural lung disease from these scans is ongoing.

In conclusion, we have shown that early structural lung disease, measured by limited slice chest CT, persists in most children once detected, with progression of disease associated with worsening neutrophilic inflammation, pulmonary infection and severe CFTR genotype. Since lung disease can commence very early in life, and early inflammation is associated with subsequent disease progression, interventions to prevent structural lung disease or arrest progression must be introduced soon after diagnosis.

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Competing interests None.

Ethics approval The study was approved by the ethics committees of the Princess Margaret Hospital, Perth, and Royal Children's Hospital, Melbourne.

Contributors Study design: LSM, CPM, CLG, PJR, CFR, SCR, PDS, SMS; literature review: LSM; data collection: LSM, CPM, CLG, AREST CF; data analysis: LSM, JP, CLG, NHdK; data interpretation: LSM, CLG, NHdK, PJR, CFR, SCR, PDS, SMS; manuscript preparation: LSM, SMS; manuscript review and approval: all authors including AREST CF.

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ONLINE SUPPLEMENT

Progression of early structural lung disease in young children with cystic fibrosis assessed using computed tomography (Mott et al)

METHODS

CT Settings

Supplementary Table – Representative CT Settings

CT settings	Perth		Melbourne	
	Inspiratory	Expiratory	Inspiratory	Expiratory
Tube voltage (kVp)	120	120	120	120
Tube current (mA)	140	140	40	48
Current time-length product (mAs)	70	70	30	36
Rotation time (msec)	500	500	750	750
Slice thickness (mm)	0.63	0.63	1.00	1.00
Collimation (mm)	2x0.625	2x0.625	2x1.00	2x1.00
Estimated CTDI (mGy) *	22.0	22.0	5.0	6.1
Estimated DLP (mGy*cm) *	8.0	8.0	3.0	3.6
Estimated radiation dose (mSv) **	0.1-0.2	0.1-0.2	0.1	0.1

Abbreviations: kVp peak kilovoltage, mA milliamperes, mAs milliamperes seconds, msec

milliseconds, mGy milliGray, mSv milliSievert, CTDI CT dose index, DLP dose length product

* Values expressed relative to a 16cm diameter CT dosimetry phantom

** Radiation doses estimated using age specific DLP conversion coefficients for children age 1 to 5 years (Thomas KE, Wang B. Age-specific effective doses for pediatric MSCT examinations at a large children's hospital using DLP conversion coefficients: a simple estimation method. *Pediatric*

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Creation of the Inflammatory Response Score

As neutrophilic inflammatory variables (total cell count, neutrophil count, IL-8 level and NE activity) were highly correlated, they were summarised, guided by a principle component analysis, into a single “inflammatory response score” as the marker of neutrophilic inflammation. First, the 4 variables underwent natural logarithmic transformation for approximate normality since the data were positively skewed. Second, principal component analysis was performed on the correlation matrix and the first principal component, which accounted for 65% of the total variability, was extracted from the data. This principal component was used as the summary measure called the “inflammatory response score”. Inflammatory response scores ranged from -5.51 to 6.14 with a median value of -0.05. Higher inflammatory response scores were interpreted as worse neutrophilic inflammation.

Effects of Specific Infection with P. aeruginosa

Current infection with *P. aeruginosa* was determined from BAL culture results, with a positive culture considered with infection at any density. Ever infection with *P. aeruginosa* was determined from annual BAL cultures from diagnosis.

RESULTS

Supplement to Table 1 – Demographics

	Initial scan (<i>t</i> -1) n=301	Subsequent scan (<i>t</i>) n=301
PULMONARY INFECTION		
Current <i>P. aeruginosa</i>	9% (26/298)	10% (30/300)
Ever <i>P. aeruginosa</i>	17% (50/298)	23% (70/300)

Supplement to Table 2 – Associations with Persistence of Bronchiectasis

	Change in Bronchiectasis Status		Model 1*	
	Transient	Persistent	Odds ratio (95% CI)	p-value
Current <i>P. aeruginosa</i> (initial scan)	11% (4/35)	14% (14/97)	1.40 (0.42-4.63)	0.580
Ever <i>P. aeruginosa</i> (initial scan)	17% (6/35)	30% (29/97)	1.87 (0.69-5.06)	0.217
Current <i>P. aeruginosa</i> (subsequent scan)	9% (3/35)	15% (15/98)	1.87 (0.50-6.91)	0.350
Ever <i>P. aeruginosa</i> (subsequent scan)	20% (7/35)	38% (37/98)	2.26 (0.88-5.77)	0.090

* Adjusted for age at the initial scan

Supplement to Table 3 – Associations with Progression of Bronchiectasis

	Model 1*	
	Regression coefficient ‡ (95% CI)	p-value
Current <i>P. aeruginosa</i> (initial scan)	1.03 (0.73-1.45)	0.884
Ever <i>P. aeruginosa</i> (initial scan)	1.10 (0.86-1.42)	0.433
Current <i>P. aeruginosa</i> (subsequent scan)	1.35 (0.99-1.84)	0.058
Ever <i>P. aeruginosa</i> (subsequent scan)	1.27 (1.02-1.59)	0.034

* Adjusted for age at the initial scan and severe CFTR genotype

‡ Regression coefficients have been back-transformed from the logarithmic scale

Supplement to Table 4 – Associations with Persistence of Air Trapping

	Change in Air Trapping Status		Model 1*	
	Transient	Persistent	Odds ratio (95% CI)	p-value
Current <i>P. aeruginosa</i> (initial scan)	2% (1/42)	12% (21/177)	4.65 (0.75-28.76)	0.098
Ever <i>P. aeruginosa</i> (initial scan)	5% (2/42)	21% (38/177)	6.75 (1.50-30.32)	0.013
Current <i>P. aeruginosa</i> (subsequent scan)	5% (2/42)	12% (21/177)	2.47 (0.61-9.91)	0.203
Ever <i>P. aeruginosa</i> (subsequent scan)	10% (4/42)	29% (51/177)	4.36 (1.42-13.41)	0.010

* Adjusted for age at the initial scan

Supplement to Table 5 – Associations with Progression of Air Trapping

	Model 1*	
	Regression coefficient ‡ (95% CI)	p-value
Current <i>P. aeruginosa</i> (initial scan)	1.30 (1.05-1.62)	0.014
Ever <i>P. aeruginosa</i> (initial scan)	1.25 (1.05-1.49)	0.012
Current <i>P. aeruginosa</i> (subsequent scan)	1.20 (0.94-1.52)	0.131
Ever <i>P. aeruginosa</i> (subsequent scan)	1.23 (1.01-1.49)	0.040

* Adjusted for age at the initial scan and severe CFTR genotype

‡ Regression coefficients have been back-transformed from the logarithmic scale