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Early View

Original Research Article

Structure and function imaging in Primary ciliary dyskinesia: cross-sectional study with ^{129}Xe ventilation / ^1H anatomical MRI

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ERJ Open Research

Structure and function imaging in Primary ciliary dyskinesia: cross-sectional study with ^{129}Xe ventilation / ^1H anatomical MRI

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Summary

Most clinically stable children with PCD had visible ventilation abnormalities when assessed with ^{129}Xe MRI. ^{129}Xe derived VDP was abnormal in 75% of children who had a normal FEV₁.

Abstract

Background

In Primary ciliary dyskinesia (PCD), impaired mucociliary clearance leads to respiratory infections. Whilst FEV₁ is commonly used to evaluate lung function in PCD its sensitivity to detect abnormalities in mild disease is limited. This study explores the utility of ¹²⁹Xe ventilation and ¹H anatomical magnetic resonance imaging (MRI) to assess lung health in children with PCD.

Methodology

This prospective cohort study assessed clinically stable children with PCD (aged 5-18 years) using ¹²⁹Xe MRI, ¹H MRI and spirometry during one session. Ventilation defect percentage (VDP) was the primary metric calculated from ¹²⁹Xe MRI.

Results

35 children were assessed, mean age= 12.3 ± 3.3 years, FEV₁ z-score= -1.5 ± 1.7. Most children had visible ventilation abnormalities and 82.8% had an abnormal ¹²⁹Xe VDP, median (IQR) ¹²⁹Xe VDP=6.3 (1.3,16.1)%. 16 participants had a normal FEV₁ (z-score > -1.6), however 12 of these individuals had abnormal ¹²⁹Xe VDP. ¹H MRI found 55.9% participants had a region of collapse; 58.1% had bronchiectasis; and 50.0% had evidence of air trapping.

Conclusion

¹²⁹Xe MRI can identify early lung function abnormalities in children with PCD and demonstrates that the extent of ventilation distribution abnormalities in PCD varies between individuals. ¹H MRI provides complementary diagnostic information on structural abnormalities.

Introduction

Primary ciliary dyskinesia (PCD) is an inherited condition in which structural and/or functional ciliary abnormalities impair mucociliary clearance. People with PCD (PwPCD) experience chronic infections with exacerbations of the upper and lower respiratory tract, fueling inflammation and the development of bronchiectasis (1). Whilst forced expiratory volume in the first second (FEV₁) from spirometry is the most widely used clinical outcome measure in PCD, large variability in FEV₁ is seen during periods of disease stability (2). Additionally, FEV₁ can be insensitive to detect abnormalities; 73% of PwPCD with structural lung abnormalities on computerised tomography (CT) scan have an FEV₁ within the normal range (3). Whilst lung clearance index (LCI) from multiple breath washout is more sensitive to detect lung abnormalities than FEV₁ (4), the correlation of LCI with other outcomes remains unclear (5). As such, more sensitive outcome measures are needed to accurately assess lung abnormalities in PCD.

Hyperpolarised gas ventilation magnetic resonance imaging (MRI) using helium-3 (³He) or xenon-129 (¹²⁹Xe) (6) permits the visualisation and quantitative assessment of the distribution of ventilation within the lung in 3-dimensions. ¹²⁹Xe which is more readily available than ³He is now commonly used in hyperpolarised gas MRI research (6) and offers potential clinical application (7). Ventilation abnormalities, visible as areas without gas signal, are known as ventilation defects. Ventilation defect percentage (VDP), the proportion of the thoracic cavity with no gas signal, is the most commonly used metric. Hyperpolarised gas ventilation MRI has shown promise in evaluating lung ventilation defects in other chronic suppurative lung diseases (CSLDs) (8), but its application in PCD to date has been limited. VDP is a sensitive marker of lung health in cystic fibrosis (CF) (8-10) and other obstructive lung diseases (6). In a small study of children with PCD, ³He VDP was found to be more sensitive to detect abnormalities than LCI and FEV₁ (11), however there are no published data using ¹²⁹Xe MRI to assess PwPCD. Recognising that it is a safe, ionising radiation-free, and well tolerated technique (6), children with PCD and parent representatives were supportive of research to explore the utility of ¹²⁹Xe MRI.

^1H MRI of the lungs can identify structural abnormalities without ionising radiation and does not require access to hyperpolarised gas technology. Findings from ^1H Ultrashort echo-time (UTE) MRI sequences are increasingly comparable with CT imaging findings, and offer longitudinal assessment capability in CSLDs (12). In some centres ^1H MRI is used for routine clinical assessment of people with CF (13). In PCD, ^1H MRI can identify collapse, consolidation, mucus plugging and bronchiectasis (14) and agreement has been seen between abnormalities identified on ^1H MRI and high-resolution computerised tomography (HRCT) (15). Functional abnormalities have been identified in PwPCD previously using the Matrix pencil ^1H MRI (MP-MRI) method, but this had lower spatial resolution and was less sensitive than UTE structural MRI to detect abnormalities (3).

This study aimed to assess lung physiology and structure in children and young people with PCD using ^{129}Xe ventilation and ^1H anatomical MRI. This work is part of a larger study (16) which assessed lung health with ^{129}Xe MRI on the same day before and after airway clearance, this paper summarises the cross-sectional data from the study population.

Methods

This prospective study recruited children (aged 5 to 18 years) with PCD from four centres in the North of England between April 2022 and August 2023. For inclusion, participants needed to: have a confirmed PCD diagnosis according to the ERS guidelines (17), be clinically stable at the time of the assessment (not on treatment antibiotics for a chest exacerbation), have resting oxygen saturation $>90\%$ on air and not have any contraindications to MRI scanning. All eligible participants or their parents/guardians if under 16 years old were approached to take part by members of their direct clinical care team and provided written informed consent. The study was approved by the Oxford B Research ethics committee (21/SC/0197).

Lung imaging

Hyperpolarised ^{129}Xe MRI scanning was performed at a single site (University of Sheffield) on either a 1.5T GE HDxt, or 1.5T GE Artist MRI scanner (GE, Milwaukee, WI, USA) using a ^{129}Xe transmit-receive vest coil (Clinical MR Solutions, Brookfield,

WI, USA). ^{129}Xe MR images were acquired using a 3D steady state free precession (SSFP) sequence (18). The ^{129}Xe gas was polarised using an in-house polariser under regulatory licence (MS-18739) (19). Images were acquired during a breath-hold at an end-inspiratory tidal volume (EITV), following inhalation of a titrated mixture of ^{129}Xe and medical grade Nitrogen, administered via a Tedlar plastic bag from residual volume. The total volume of inhaled gas was calculated based on participant's height and ranged between 0.4 and 1.0 Litre (20). To estimate the thoracic cavity for image analysis, ^1H 3D spoiled gradient echo (SPGR) MR images were acquired pre and post the ^{129}Xe images during separate matched breath-holds at EITV. ^{129}Xe images were segmented using a semi-automated method (21) to calculate ^{129}Xe VDP and the ventilation heterogeneity index (VHI) (22) was also assessed.

^1H imaging sequences included:

- 3D SPGR acquired at two lung inflation levels: functional residual capacity plus inhaled bag volume (i.e. EITV), and residual volume (RV),
- Radial 3D UTE during free breathing (23),
- 2D SSFP at expiratory breath-hold,
- 2D T2-weighted Half-Fourier Acquisition Single-shot Turbo spin Echo imaging (HASTE) at expiratory breath-hold).

These were performed using an 8-channel or 16+ channel anterior array plus a spine ^1H array (HDxt and Artist MRI scanner respectively) (GE, Milwaukee, WI, USA) to provide high resolution anatomical images. A consultant paediatric radiologist reviewed all images to assess for typical PCD features including bronchiectasis, collapse, consolidation, and air trapping. Participants were imaged at three times on the same day (see OLS for study schedule). Baseline ^1H imaging was primarily used to assess for abnormalities, however where baseline images were not readable, ^1H MR images from later time points were used.

Further details on sequence details, image acquisition, processing, and calculation of ^{129}Xe MRI metrics are provided in the supplementary information.

Acquisition of non-imaging data

Clinical and diagnostic information were obtained prior to the study visit from medical notes and the clinical team. Spirometry was performed immediately prior to the first MRI scan with a 'PFT Pro' (Vyaire, Basingstoke, UK) in accordance with technical standards (24). All spirometry metrics were assessed using a z-score, with -1.64 used as the lower limit of normal (25). Quality of life was assessed on the same day using the age-appropriate version of the QOL-PCD online questionnaire: adolescent for aged 13-17 years and child and parent proxy for those aged under 13 years (26).

Statistical analysis

Statistical analysis was performed using GraphPad (Prism version 9.5.1, San Diego, US). Normal distribution was assessed using the Shapiro–Wilk test. Data are presented as mean \pm SD, if they are normally distributed, or median (range) otherwise. Group comparisons were completed using the un-paired t-test, Mann-Whitney U test (not normal distribution), parametric analysis of variance test or Kruskal-Wallis analysis of variance test, depending on data distribution and number of groups. Correlation was assessed using Spearman's rank-based correlation. Statistical significance was defined as $p < 0.05$. ^{129}Xe VDP of 1.2% and VHI of 9.5% were used as the upper limits of normal from a population of children without history of lung disease (27).

Results

93 individuals were assessed for eligibility; 15 did not meet inclusion criteria and 76 were approached. 36 individuals consented to participate of which 35 young people were assessed with ^{129}Xe and ^1H MRI and one individual withdrew due to intolerance of the scanner sounds. 35 participants were assessed with QOL-PCD in total: adolescent version (n=16), both child and parent-proxy versions (n=14), child version (n=3), parent-proxy version (n=2). Full details of screening and recruitment are provided in Figure 1. Baseline clinical demographics are provided in Table 1 and QOL-PCD data in the online supplementary material.

Spirometry

34 individuals performed technically acceptable spirometry (28) (Table 2), one data set was not technically acceptable and was excluded. 55.8% (n=19) had no evidence of airway obstruction (FEV_1/FVC z-score >-1.64). No correlation was seen between FEV_1 and age, age at diagnosis, number of antibiotic courses or lower respiratory symptoms/physical functioning QOL-PCD domains. No significant differences in FEV_1 were seen between ultrastructural defect groups, or those who had/had not recently grown *Pseudomonas aeruginosa*.

^{129}Xe MRI

A summary of ^{129}Xe derived metrics is provided in Table 2 and an image from each of the participants in Figure 2. Most children had visible ventilation abnormalities and 82.8% (n=29) had an abnormal VDP. Six children had very minimal ventilation abnormalities and were comparable to healthy children.

No difference in ^{129}Xe VDP was seen according to sex, or situs abnormalities. No correlation was seen between ^{129}Xe VDP and age, or age at diagnosis. ^{129}Xe VDP was highest in the inner dynein arm (IDA) defects or microtubular disorganisation (MTD) group, with a median (IQR) of 12.8% (1.5, 19.3%). The median (IQR) for the other groups were outer dynein arm defect (ODA), 5.8% (1.8, 7.1%); combined ODA and IDA defects, 9.2% (5.0, 19.6%); and normal ultrastructure with a genetic diagnosis of PCD, 0.4% (0.3, 2.5%). There was no statistically significant difference in ^{129}Xe VDP between the groups.

^{129}Xe VDP correlated moderately with FEV_1 ($r=-0.54$, $p=0.001$), FVC ($r=-0.42$, $p=0.01$) and FEV_1/FVC ($r=-0.57$, $p<0.001$) (Figure 3A-C). The number of individuals with abnormal ventilation was greater as assessed by ^{129}Xe VDP when compared to FEV_1 ; 16 participants had an FEV_1 within the normal range (z-score >-1.64) yet 12 of these individuals (75%) had raised VDP when compared to healthy children (^{129}Xe VDP $>1.2\%$). Two children had a normal VDP ($<2.0\%$) with abnormal spirometry: one with an obstructive pattern (FEV_1/FVC z-score <-1.64 and $\text{FEV}_1 <-1.64$) and one with a mixed pattern (FEV_1/FVC z-score <-1.64 and FVC <-1.64). Three children had no evidence of obstruction (FEV_1/FVC z-score >-1.64) but had a restrictive pattern on spirometry (FEV_1 and FVC z-score <-1.64). All of these children had an abnormal VDP (11.0, 11.3 and 27.7%).

Examples of cases in which only one of the two lung health measures were abnormal (either FEV_1 or ^{129}Xe VDP) are shown in Figure 4.

There was no correlation between ^{129}Xe VDP and the number of antibiotics administered in the last 12 months. Whilst ^{129}Xe VDP was higher in individuals who had cultured *Pseudomonas aeruginosa* (PsA) in the previous 12 months (11.0%) than those who had not (5.4%), this was not statistically significant ($p=0.15$). Across the three versions of the QOL-PCD, only the social function domain (adolescent version) showed a moderate correlation with ^{129}Xe VDP ($r=-0.57$, $p=0.02$).

25 participants (71.4%) had abnormally high ^{129}Xe VHI. ^{129}Xe VHI correlated moderately with spirometry: FEV_1 z-score ($r=-0.52$, $p=0.002$); FVC z-score ($r=-0.38$, $p=0.03$), and FEV_1/FVC z-score ($r=-0.58$, $p<0.001$) (Figure 3D-F). No correlation was seen between ^{129}Xe VHI and age, age at diagnosis, number of antibiotic courses, and lower respiratory/physical functioning QOL-PCD domains. No significant differences in ^{129}Xe VHI were seen between ultrastructural defect groups, or those who had or had not recently grown PsA.

¹H MRI assessment of lung structure

Baseline UTE scans were deemed readable for 24/34 (70.1%) participants, a further five participants (14.7%) had unreadable baseline UTE images, but scans from later timepoints within the day were readable. In the remaining five cases image acquisition issues permitted assessment for collapse but not bronchiectasis (n=3). Of those with readable images: 19/31 (61.3%) had bronchiectasis; 19/34 (55.9%) had a region of collapse; 17/34 (50.0%) participants had evidence of air trapping. All 19 with collapse identified had a region of collapse in their middle lobe (right middle lobe in situs solitus and left middle lobe in situs inversus). ¹²⁹Xe imaging was more sensitive to detect abnormalities than ¹H MRI; 4/6 (66.7%) of those with no structural abnormalities had both abnormal VDP and VHI (Figure 5). Figure 6 provides an example visual comparison of abnormalities identified on ¹H and ¹²⁹Xe MRI.

Discussion

This cross-sectional study found that ¹²⁹Xe MRI detects ventilation abnormalities in most children with PCD. 82.8% of the study population had abnormal ventilation assessed by ¹²⁹Xe VDP; 75% of individuals with no evidence of obstructive airways disease using spirometry had abnormal ventilation (VDP>1.2%). Whilst ventilation abnormalities were common, a small number of individuals with PCD did not have ventilation abnormalities. ¹H MRI also identified structural abnormalities: 55.9% of children with PCD had areas of collapse and 61.3% had bronchiectasis.

Our findings are consistent with the previous small study by Smith, West (11) which found VDP (using ^3He MRI) to be abnormal in children with PCD who had normal FEV_1 . With the finding that ^{129}Xe VDP is often abnormal in individuals with a normal FEV_1 , our results align with previous hyperpolarised gas MRI literature in CSLDs (9, 11, 20, 29-32). For the first time this study found a small number of children with PCD with abnormal FEV_1 , including one without restrictive lung disease, but normal ventilation on ^{129}Xe MRI (27). Whilst this study found a moderate correlation of ^{129}Xe VDP with FEV_1 in children with PCD, other studies have reported no correlation between ^{129}Xe VDP with FEV_1 (29), and a stronger correlation between ^3He VDP and FEV_1/FVC (11). In paediatric CF, the relationship between ^{129}Xe VDP and FEV_1 , is also variable (30, 33, 34). This variability in correlation between VDP and FEV_1 may arise from the differing study populations', or image acquisition methods. These findings illustrate that reliance on spirometry alone for the clinical assessment of lung health in CSLDs may lead to instances of false assurance of normal lung health. These studies show the importance of using a range of tools to assess different components of lung function in CSLDs.

The clinical picture in PCD is complex; neither spirometry nor ^{129}Xe metrics (VDP and VHI) correlated with age, age at diagnosis, or number of antibiotic courses in the last 12 months, and limited correlation was seen with quality of life assessed by the QOL-PCD. VDP was higher, and FEV_1 lower in those with IDA and ODA defects than those with normal ciliary ultrastructure. Previous research reported that individuals with IDA and MTD ciliary defects have the highest CT scan disease scores (35), and this study provides further evidence that ciliary ultrastructural defects may influence disease severity.

Bronchiectasis, bronchial wall thickening, mucus plugging, and atelectasis are common CT findings in people with PCD (36, 37). The frequency of abnormalities identified in children with PCD using ^1H MRI in this study is lower than previously reported: 100% structural abnormalities, 97% and 94% bronchiectasis or wall thickening (3, 14). It is unclear if the lower frequency seen may arise from differences in diagnostic inclusion criteria, image acquisition or a true variance. Finding the anatomical middle lobe, whether located on the left or right side of the body to be a common area of collapse in PCD, this study adds further evidence to the CT scan findings of middle lobe collapse reported by Jain, Padley (38) and highlights the importance of monitoring this area in children with PCD. These data also highlight the potential of ^1H MRI to reduce the number of CT scans of the lungs and provides a viable longitudinal lung imaging assessment method.

This is the largest cohort of children with PCD assessed by ^{129}Xe MRI to date, employing a prospective study design with standardised imaging and spirometry methods, an age-appropriate, validated QOL-PCD instrument and clinical data. This study used ^{129}Xe ventilation MRI, a highly sensitive technique, to provide quantitative and qualitative information on the range of functional abnormalities seen in PCD, and ^1H anatomical MRI to detect structural abnormalities. Although two children struggled with the MRI scanner noise, one of whom withdrew from the study, the scans were well tolerated by the vast majority of individuals, including those as young as 6 years old. As MRI is ionising radiation-free, it is well placed for assessing children and young people, especially longitudinally. The cross-sectional design precluded assessment of longitudinal changes or inferring causality. Future long-term follow-up studies to determine whether baseline ^{129}Xe VDP predicts key clinical outcomes including exacerbation frequency and lung function decline are needed. Further evaluation of healthy children with ^{129}Xe MRI will provide more robust paediatric reference values enabling multicentre research and clinical implementation of the findings. Whilst high-resolution CT (HRCT) scans were not included to ensure an ionising radiation-free protocol, studies permitting direct comparison of HRCT and ^1H MRI in children with CSLDs are warranted. As LCI is widely available, inclusion of LCI in the study protocol would have provided a valuable comparison of ^{129}Xe MRI with LCI. Future work from the ASPECT-PCD study (16) will evaluate the effects of airway clearance techniques using ^{129}Xe MRI.

Conclusion

^{129}Xe ventilation MRI can detect ventilation abnormalities in children with PCD, we also demonstrate that the extent of ventilation distribution abnormalities in PCD varies between individuals. Ventilation abnormalities are identifiable using ^{129}Xe ventilation MRI in children with PCD who have a normal FEV₁. ^1H MRI provides complementary diagnostic information on structural abnormalities. ^{129}Xe ventilation MRI offers a sensitive, safe, and well-tolerated imaging technique that is well placed for future longitudinal studies in children with PCD.

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Table 1: Participant demographics including ultrastructural defects. Data is presented as mean \pm SD or median (IQR) depending on if data is normally distributed. Ultrastructural defects within study population presented as n (%). IDA=inner dynein arm defect, ODA= outer dynein arm defect, MTD= microtubular disarrangement. *PCD diagnosis confirmed by genetics. PsA=Pseudomonas aeruginosa.

Clinical data (n=35)	Data reported as n (%), mean \pm SD, or median (IQR)
Sex	19 male (54%)
Age (years)	12.3 \pm 3.3
Age at diagnosis (years)	3 (0, 6)
Ultrastructural defect	
IDA or MTD only	9 (25.7%)
ODA only	8 (22.9%)
IDA and ODA	12 (34.3%)
No defect*	6 (17.1%)
Situs inversus	14 (40.0%)
No. of antibiotic courses in last 12 months (oral or IV, planned or unplanned)	2.5 (1,4)
PsA (cultured in last 12 months)	5 (14.3%)

Table 2: Spirometry and ^{129}Xe MRI metrics

	Data reported as mean \pm SD or median (IQR)	
Spirometry (n=34)	z-score	% predicted
FEV ₁	-1.5 \pm 1.7	81.3 \pm 20.6
FVC	-0.7 \pm 1.5	90.1 \pm 21.6
FEV ₁ /FVC	-1.5 \pm 1.3	87.2 \pm 11.7
^{129}Xe MRI (n=35)		
VDP (%)	6.3 (1.3,16.1)	
VHI (%)	11.3 \pm 3.2	

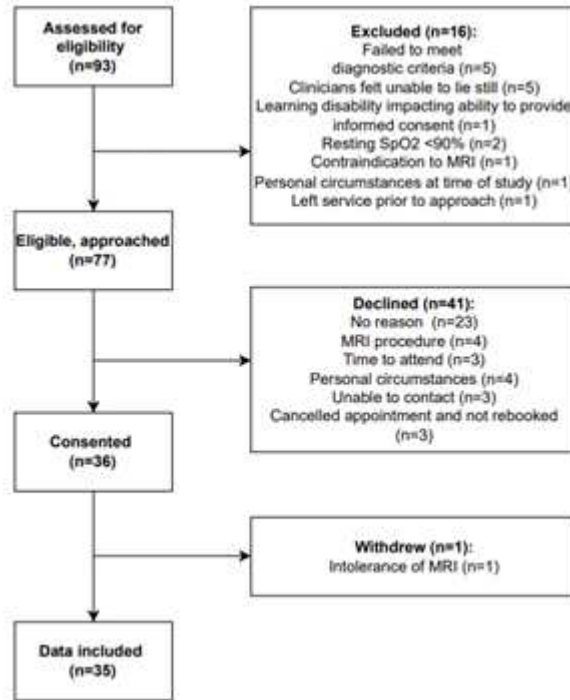


Figure 1

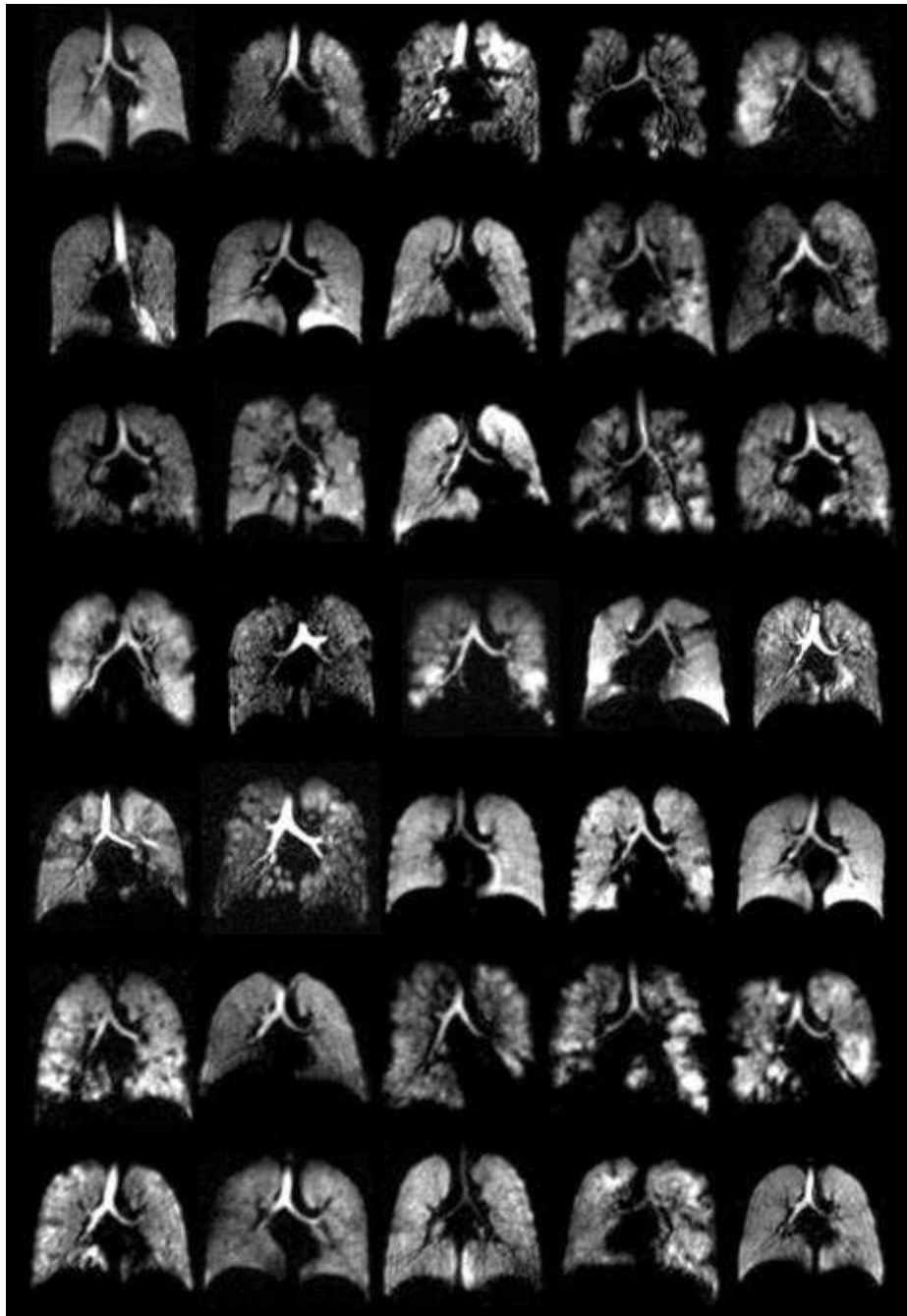


Figure 2

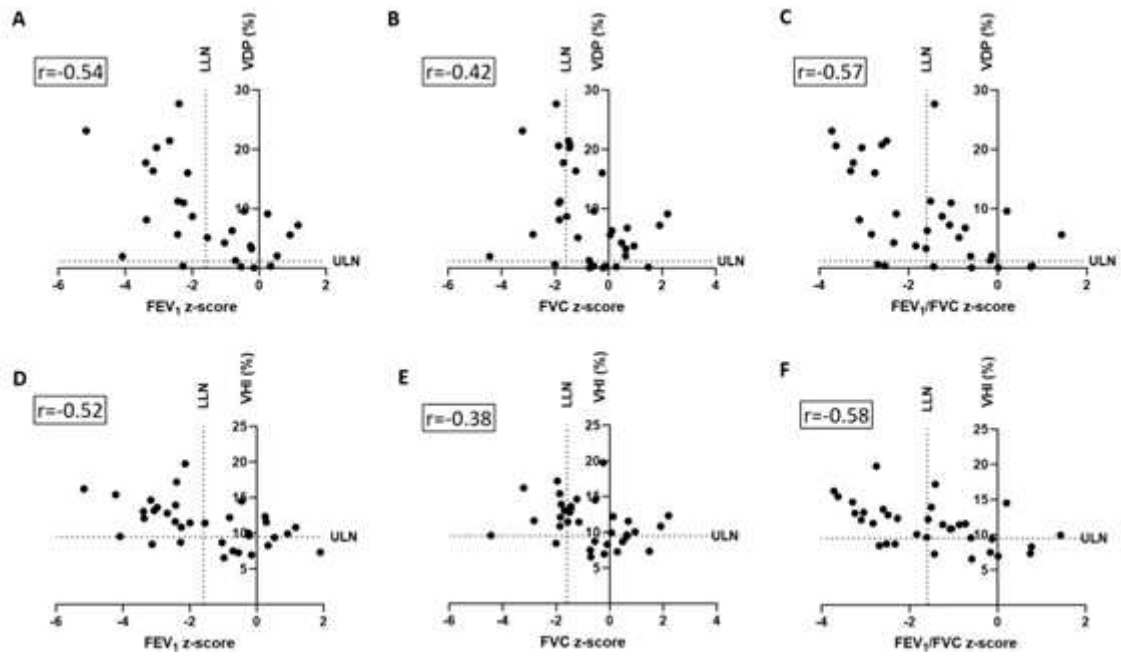


Figure 3

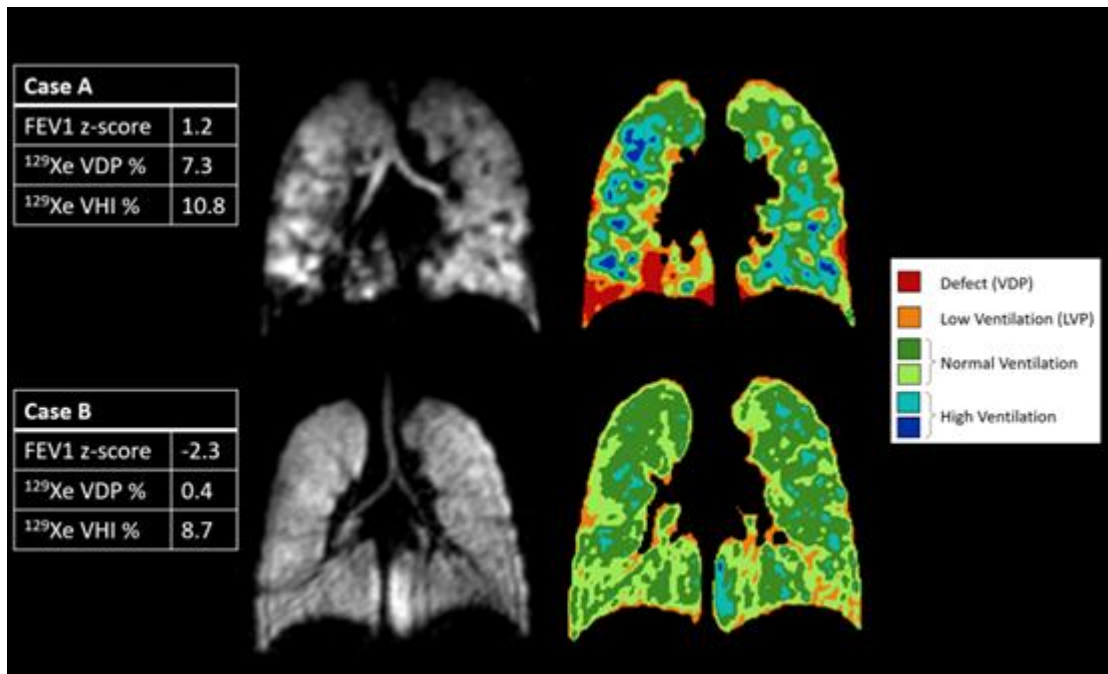


Figure 4

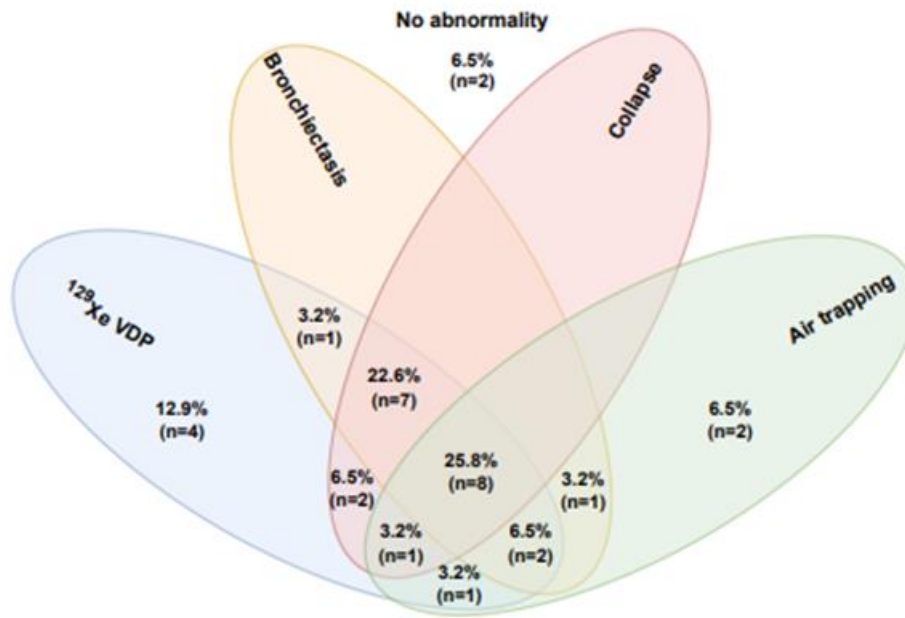


Figure 5

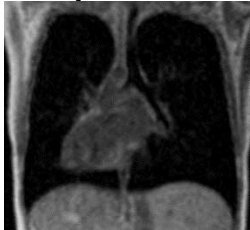
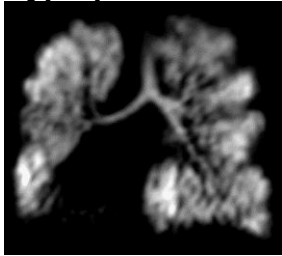
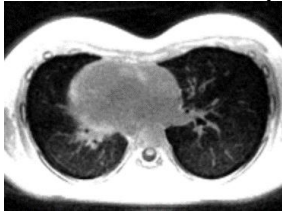




Figure 6

Image acquisition

Details of the imaging sequences are provided in Table S1.

Table S1: Overview of imaging sequences performed.

	<p>3D Spoiled Gradient Echo (SPGR)</p> <p>Description: T1 weighted images, taken at an end inspiratory tidal volume (EITV).</p> <p>Role: Images used for VDP calculation.</p> <p>Scan details: Plane: Coronal. Slice thickness 5.0mm, 36 slices per acquisition. Field of view (FOV) 40.0cm. Scan duration: Typically, 6 second breath hold at EITV. Coil: body transmit-receive coil.</p>
	<p>Hyperpolarised ¹²⁹Xe gas ventilation steady state free precession</p> <p>Description: Imaging of lung ventilation with ¹²⁹Xe as contrast.</p> <p>Role: Functional MRI scan. Signal from ¹²⁹Xe offers visualisation of the distribution of ventilation. Used to calculate metrics and for visual analysis of ventilation.</p> <p>Metrics: ¹²⁹Xe VDP and VHI.</p> <p>Scan details: Steady state free precession sequence. Plane: Coronal. Slice thickness: 10.0mm FOV 40.0cm. Scan duration: Typically, 6 second breath hold at EITV. Coil: ¹²⁹Xe transmit receive flex quad coil</p>
	<p>Ultra-short echo (UTE)</p> <p>Description: Proton density weighted sequence which provides high-resolution isotropic 3D images.</p> <p>Role: Structural MRI scan. Visualisation of the lung parenchyma with high-resolution, similar in contrast to CT.</p> <p>Scan details: Plane: Any (isotropic). resolution 1.3 mm isotropic. FOV 35cm. Scan duration: 8 to 9 minutes, relaxed breathing, data prospectively or retrospectively gated to expiration (i.e. at a lung volume near FRC). Coil: 8 channel array (HDxt) or 16+ channel anterior array + spine array.</p>
	<p>Steady State Free Precession (SSFP)</p> <p>Description: Offers a combination of T1 and T2 weighted contrast with hyper intense signal from long T2 components.</p> <p>Role: Structural MRI scan. Allows visualisation of anatomy and can visualise mucus.</p> <p>Scan details: Plane: Axial, 25 slices Slice thickness: 10mm. FOV 40.0cm. Scan duration: Typically, 8 second expiratory breath hold. Coil: 8 channel array.</p>
	<p>Half-Fourier-Acquired Single-shot Turbo Spin Echo (HASTE)</p> <p>Description: A rapid T2 weighted sequence.</p> <p>Role: Structural MRI scan. Allows visualisation of mucus. The acquisition speed reduces the impact of motion on the images.</p> <p>Scan details: Plane: Coronal. Slice thickness: 10mm, 18 slices. FOV 35.0cm. Scan duration: Typically, 8 second inspiratory breath-hold. Coil: 8 channel array</p>
<p>3D Spoiled Gradient Echo (SPGR) with isotropic voxel size</p>	



Description: T1 weighted images (as above), taken at EITV and approximately RV.

Role: At higher lung volumes images may visualise mucus, and at lower volumes air trapping.

Scan details: Plane: Any (isotropic) Slice thickness 3.0mm, 62 slices per acquisition. FOV 40.0cm. Scan duration: Typically, 6 second breath hold at TLC, EITV and RV. Coil: 8 channel array

¹²⁹Xe MRI image acquisition

Hyperpolarised gas ventilation images were acquired by inhaling a titrated mixture of ¹²⁹Xe and medical grade nitrogen, which were administered via a Tedlar plastic bag. An MRI scanner sequence which only images the ¹²⁹Xe gas is used in combination with a dedicated ¹²⁹Xe coil, to image the areas of the lung which have been ventilated by the polarised gas. Images were acquired during a single breath-hold, taken at an estimated end-inspiratory tidal volume (EITV). Participants were coached to perform at least two relaxed tidal breaths before then inhaling a fixed volume of the gas from a resting FRC lung volume. The total volume of inhaled gas was calculated based on the participant's height with the aim of achieving an approximate same lung inflation level across the group irrespective of the participant's size. This process of inhaling the volume of the bag from FRC provided an estimated volume of 60% of TLC, referred to as EITV. The volume of the gas, which ranged from 0.4 to 1.0 L comprised a scaled dose of ¹²⁹Xe balanced with medical grade Nitrogen as shown in Table S2, in line with previously published methods (1).

Table S2: Inhaled gas volume and composition height-based calculations.

Patient height (standing)	Total volume	Gas mixture composition
Over 160cm	1.0 L	500 ml ¹²⁹ Xe + 500 ml N ₂
150-160cm	800 ml	450 ml ¹²⁹ Xe + 350 ml N ₂
140-150cm	650 ml	400 ml ¹²⁹ Xe + 250 ml N ₂
130-140cm	500 ml	350 ml ¹²⁹ Xe + 150 ml N ₂
Less than 130cm	400 ml	300 ml ¹²⁹ Xe + 100 ml N ₂

Calculation of VDP and VHI

^1H matched anatomical images were registered to the ^{129}Xe ventilation images. Segmentation of the registered anatomical images was completed using in house software via MATLAB (MathWorks, Natick, MA) with a semi-automated process. An initial automated map of the estimated lung cavity was generated using deep-learning (2) or a combined approach employing deep-learning and fuzzy-c means clustering (3-5) which was then corrected as needed by manual editing to give the thoracic cavity volume (TCV). For individuals with situs inversus, images were flipped on a vertical axis prior to segmentation to optimise performance of the automated deep-learning segmentation. Images were flipped back to situs inversus orientation after the segmentation was completed. The ^{129}Xe ventilation signal was categorised using an automated linear “binning” process (6) in which voxels were sorted from low to high signal, to form a linear distribution and the signal was divided by the mean ventilation signal within the thoracic cavity. This ventilation distribution was then divided into six categories known as “bins” centred around the mean signal and with the width of each bin equal to one standard deviation. The two central bins were combined into a single normal ventilation bin, the two highest signal bins combined into a single high ventilation bin. Each voxel was therefore assigned into one of four bins: high ventilation, normal ventilation, low ventilation, or ventilation defect. ^{129}Xe VDP, the proportion of the lung with no ventilation, was calculated as the ventilation defect volume from bin 1 / TCV * 100. ^{129}Xe VHI was calculated using an in-house automated process which assesses each ventilated voxel with the eight in-plane adjacent voxels using an established method (7). From this, a map of the co-efficient of variation of signal intensity is produced and the interquartile range of the coefficient of variation of signal intensity is used as the ventilation heterogeneity index (VHI).

TableS3: MRI scan sequence order with details of coil used.

Order	Image sequence	Coil
1	^{129}Xe at EITV	^{129}Xe transmit-receive vest
2	SPGR at EITV	In-built transmit receive
3	HASTE	^1H array
4	SPGR at EITV	
5	SPGR at RV	
6	SSFP	
7	UTE	

Table S4: Study visit schedule.

Order and content of study visit	
1	Height and weight
2	Spirometry
3	Oxygen saturations (finger probe)
4	Pre-ACT MRI (0 hours)
5	ACT completion
6	MRI (+2 hours from baseline)
7	QOL-PCD
8	MRI (+6 hours from baseline)

Table S5: QOL-PCD scores

QOL-PCD Domain	Data reported as mean \pm SD or median (IQR)		
	Child (n=17)	Adolescent (n=16)	Parent proxy (n=16)
Physical functioning	69.4 \pm 21.0%	80.6 \pm 26.5%	71.3 \pm 26.8%
Emotional functioning	69.6 \pm 16.7%	80.2% \pm 16.9%	67.4 \pm 18.8%
Treatment burden	62.3 \pm 27.1%	72.2 \pm 23.3%	63.0 \pm 19.2%
Role	N/A	78.4 \pm 20.8%	N/A
Social functioning	71.0 \pm 30.2%	70.8 \pm 19.3 %	75.7 \pm 17.3%
Vitality	N/A	59.7 \pm 27.8%	N/A
Upper respiratory symptoms	63.1 \pm 26.7%	74.2 \pm 15.6%	47.9 \pm 18.4%
Lower respiratory symptoms	58.0 \pm 22.9%	75.6 \pm 15.8%	55.4 \pm 19.7%
Hearing	69.6 \pm 26.2%	93.2 \pm 9.2%	77.8 \pm 21.8%
Eating and weight	N/A	N/A	76.2 \pm 27.8%
Health perception	N/A	N/A	67.0 \pm 16.9%

Table S6: Diagnostic information for participants taken from clinical data: cilia ultrastructural abnormalities, genetic defects and nasal nitric oxide in ppb (parts per billion).

No	Ultrastructural abnormality	Genetic defect	Nasal nitric oxide (ppb)
1	IDA	Nil on record	29
2	Normal	DNAH11	213
3	ODA	DNAH5	Nil on record
4	IDA	CCDC39	14
5	IDA	HYDIN	<5
6	IDA+ODA	DNAH5	49

7	IDA+ODA	SPAG1	20
8	IDA+ODA	DNAH5	42
9	ODA	DNAH5	24
10	IDA+ODA	DNAH5	20
11	IDA+ODA	DNAAF5	<5
12	IDA+ODA	Nil on record	19
13	IDA+ODA	DNAH5	9
14	MTD	CCDC40	12
15	IDA+ODA	DNAAf5	5
16	IDA+ODA	CCDC103	49
17	Normal	CCDC103	537
18	IDA	Nil identified	51
19	Normal	LRRCG	680
20	MTD	Nil identified	30
21	ODA	DNAH5	67
22	MTD	RSPH4A	91
23	Normal	DNAH11	307
24	IDA + ODA	CCDC40	32
25	IDA	DNAH5	39
26	ODA	DNAH11	40
27	Normal	DRC1	116
28	IDA+ODA	LRRC6	<5
29	IDA+ODA	CCDC40	23
30	ODA	Nil on record	<5
31	ODA	DNAH5	18
32	Normal	DNAH11	30
33	ODA	Nil identified	355
34	IDA+ODA	DNAH5	22
35	ODA	DRC1	551

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