FOOTPRINTS study protocol: rationale and methodology of a 3-year longitudinal observational study to phenotype patients with COPD

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INTRODUCTION

Airflow limitation is a key characteristic of chronic obstructive pulmonary disease (COPD).1,2 It is caused by small airways disease, such as obstructive bronchiolitis, and structural changes related to parenchymal destruction (emphysema).1,2 Small airways disease and structural changes can occur alone or in combination with one another, with the degree of severity and relative contributions varying between individuals.3

COPD is associated with inflammation and tissue damage, which is thought to be linked to—among other aspects—an imbalance between neutrophil serine proteases (NSPs) and their inhibitors.3,4 The serine protease inhibitor alpha1-antitrypsin (A1AT) inactivates NSPs such as neutrophil elastase (NE), proteinase 3 (PR3) and cathepsin G

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ABSTRACT

Introduction A better understanding is needed of the different phenotypes that exist for patients with chronic obstructive pulmonary disease (COPD), their relationship with the pathogenesis of COPD and how they may affect disease progression. Biomarkers, including those associated with emphysema, may assist in characterising patients and in predicting and monitoring the course of disease. The FOOTPRINTS study (study 352.2069) aims to identify biomarkers associated with emphysema, over a 3-year period.

Methods and analysis The FOOTPRINTS study is a prospective, longitudinal, multinational (12 countries), multicentre (51 sites) biomarker study, which has enrolled a total of 463 ex-smokers, including subjects without airflow limitation (as defined by the 2015 Global Initiative for Chronic Obstructive Lung Disease (GOLD) strategy report), patients with COPD across the GOLD stages 1–3 and patients with COPD and alpha1-antitrypsin deficiency. The study has an observational period lasting 156 weeks that includes seven site visits and additional phone interviews. Biomarkers in blood and sputum, imaging data (CT and magnetic resonance), clinical parameters, medical events of special interest and safety are being assessed at regular visits. Disease progression based on biomarker values and COPD phenotypes are being assessed using multivariate statistical prediction models.

Ethics and dissemination The study protocol was approved by the authorities and ethics committees/institutional review boards of the respective institutions where applicable, which included study sites in Belgium, Canada, Denmark, Finland, Germany, Japan, Korea, Poland, Spain, Sweden, UK and USA; written informed consent has been obtained from all study participants. Ethics committee approval was obtained for all participating sites prior to enrolment of the study participants. The study results will be reported in peer-reviewed publications.

Trial registration number NCT02719184.

Strenghts and limitations of this study

► The study helps to address an unmet need to understand different chronic obstructive pulmonary disease (COPD) phenotypes and to expand on our understanding of biomarkers that are associated with emphysema, and hence COPD, in a 3-year longitudinal setting.

► A subset of patients with alpha1-antitrypsin deficiency provides information on an important but rarely studied subpopulation of patients that present with earlier onset and faster progression of emphysema.

► A regular sampling schedule is employed, with frequent visits and biomarker sample collections to allow early detection of changes, if present.

► MRI and sputum collection only occur at certain study sites and not necessarily at all planned times; hence, the smaller subsets of patients may not be representative of the entire population.

► Patients with the newly defined preserved ratio impaired spirometry phenotype are not included, nor are current smokers or healthy subjects who have never smoked, thus limiting the control groups for reference.
(Cat G) to protect lung tissue against protease-mediated damage. Although a number of gene defects have been associated with emphysema, α1-antitrypsin deficiency (A1ATD) is the most well established as a cause for COPD. It is associated with a higher risk of developing emphysema that tends to occur at a younger age and progresses faster than for patients without A1ATD. Subjects with the most severe A1ATD are usually homozygous for the ZZ allele. The resulting emphysema among patients with A1ATD is thought to be caused primarily by increased NSP activity in the lungs.

NE and PR3 are responsible for the degradation of critical components of the extracellular matrix, including elastin and fibronectin, and can produce other key features of COPD, such as mucous hypersecretion. When quantified, levels of NE, PR3 or their specific elastin degradation products have been associated with poorer disease outcomes in patients with COPD, such as incidence of exacerbations and higher risk of mortality. Other proteases and biomarkers, including desmosine, fibrinogen and C reactive protein, have also been associated with poorer disease outcomes in patients with COPD.

In the era of precision-based patient care, there is a need for a more in-depth understanding of COPD patient phenotypes, pathogenesis and progression. Biomarkers may assist in characterising patients and in predicting and monitoring the course of disease, in order to better enable the development of drugs aimed at slowing COPD progression. In fact, there are encouraging signs that therapies that modify NSP imbalance (eg, augmentation therapy for A1ATD) appear to alter emphysema progression, at least when considering blood and radiological biomarkers of disease.

To assess the correlation of physiological biomarkers with respect to their feasibility and rate of, disease progression.

METHODS AND ANALYSIS
Study design
The FOOTPRINTS study (registered on 21 March 2016) is a multinational, prospective, longitudinal, biomarker study in subjects without airflow limitation, patients with COPD and patients with COPD and A1ATD, taking place at 51 sites across 12 countries (Belgium, Canada, Denmark, Finland, Germany, Japan, Korea, Poland, Spain, Sweden, UK and USA). A complete list of study sites is detailed in online supplemental table 1.

The study consists of two consecutive periods (a screening period lasting up to 28 days and an observational period lasting 156 weeks) and includes seven site visits and additional phone interviews (figure 1). The observational period is completed at visit 7 or at the discontinuation visit, which is regarded as study completion. Patients are considered to be in the observational phase only after completing visit 2. All subjects who prematurely discontinue the study during the observational phase only after completing visit 2. All subjects who prematurely discontinue the study during the observational phase are followed for vital status and COPD exacerbation status (for patients with COPD and for patients with COPD and A1ATD only) via phone contact until 156 weeks after visit 2.

Calculation of sample size and power calculations for annual decline in forced expiratory volume in 1s (FEV1) and 15th percentile density assessed by CT are provided in online supplemental text 1 and tables 2 and 3.

Inclusion and exclusion criteria
Male and female participants who have signed an informed consent form consistent with the International Conference on Harmonisation Good Clinical Practice (GCP) guidelines prior to participation have been included, provided the following were applicable:

- Aged 40–70 years (30–70 years for patients with COPD and A1ATD).
Ex-smokers for ≥9 months with a smoking history of ≥20 pack-years (≥10 pack-years for patients with COPD and A1ATD).

Body mass index 18–35 kg/m² (≤30 kg/m² in the MRI subset).

Able to perform all study-related procedures.

Required to have been on stable therapy for 4 weeks prior to visit 1.

Participants were excluded if any of the following were applicable:

- Significant pulmonary disease (other than COPD) or other significant medical condition, for example, rheumatoid arthritis, severe liver disease or psoriasis.
- Documented history of asthma, including during childhood.
- Any respiratory tract infection or COPD exacerbation within 4 weeks prior to visit 1 or during the screening period prior to visit 2, if it was not possible to meet the rescheduling rules.
- Presence of an immunocompromising condition (eg, HIV or history of hepatitis B or C).
- Any treatment with phosphodiesterase-4 inhibitors and maintenance treatment with methylxanthines within 3 months or 6 half-lives prior to visit 1 and until visit 2; patients were permitted to initiate treatment following visit 2.
- Patients with COPD (with and without A1ATD) with newly added anti-inflammatory treatment (either respiratory or non-respiratory) or change in any therapy within 4 weeks prior to visit 1 and during the screening period between visit 1 and visit 2; patients were permitted to initiate treatment following visit 2.
- Any prior, current or planned A1AT augmentation therapy.
- Permitted and restricted medications and therapy are detailed in online supplemental table 4.
- Pregnant or lactating women.
- Previous participation in this study or participation in a parallel interventional clinical trial within 6 weeks prior to visit 1 or during the study.

Patient population and recruitment

The study is ongoing; however, enrolment is complete. In total, 463 patients were enrolled, who are described in more detail in table 1. Strategies for achieving adequate participant enrolment and retainment are described in online supplemental text 1.

The study started on 22 July 2016 when the first subject was screened, and the planned end date for the study is 29 March 2021.

Main outcome measures

The main outcome measures are change in lung density (assessed by CT), lung function decline, and number, duration and severity of exacerbations over the course of the study.
Planned analyses and assessments
Variables to be analysed include a range of soluble biomarkers, imaging biomarkers and clinical parameters, as well as medical events of special interest (MESI) and safety assessments. The pre-specified biomarkers being assessed are summarised in table 2, and the biomarker sampling schedule and overall trial design are illustrated in figure 1.

Blood and sputum are being collected at visits 2–7 for analysis of pre-specified soluble biomarkers, and blood is also being collected at visit 1 to perform a genetic test for A1ATD (figure 1). Total and differential cell counts in blood, which are being assessed within the safety laboratory assessments, are also being assessed as biomarkers. Sputum will be induced to generate sputum samples, including cell slides to assess total and differential cell count and supernatant to assess soluble biomarker levels (table 2).

Chest CT scans are being conducted annually at visits 2, 5, 6 and 7, following a specific low-dose protocol in order to minimise radiation exposure at each visit. A COPD-Gene phantom is being used before the first patient scan and then bi-monthly to monitor the stability of CT measurements for each scanner. MRI evaluation of the pulmonary vasculature and right ventricle is conducted in a subset of patients without airflow limitation and patients with COPD (not in patients with COPD and A1ATD) at visits 2, 4, 5, 6 and 7. MESI include respiratory disorders such as pneumonia, pulmonary fibrosis, bronchiectasis and COPD exacerbations, as well as any other disorders (eg, cardiovascular, metabolic/digestive and neurological/psychiatric) reported at the investigator’s discretion. COPD exacerbations are defined as new or increased lower respiratory events or symptoms related to the underlying COPD with duration ≥3 days. Assessment of safety is carried out at visits 1, 5, 6 and 7 unless otherwise indicated, including physical examination (plus assessment of height (visit 1 only) and weight), vital signs (blood pressure and pulse rate), safety laboratory parameters (from blood; assessed at every site visit (visits 1–7)) and ECG. Adverse events associated with any study procedure and MESI are collected at every site visit (visits 1–7; MESI also assessed via telephone contact between visits; figure 1). All adverse events are followed up until they have resolved, been sufficiently characterised or no further information can be obtained.

Rescheduling of clinic visits
Any deviations from the planned visit schedule are being documented. Rescheduling of certain clinic visits was permitted to promote participant retention. Clinic visits that involve a lung function test may be rescheduled twice within the permitted time windows due to violation of lifestyle restrictions. Subjects should refrain from strenuous activity for at least 12 hours prior to lung function testing (and throughout the testing period) and should avoid cold temperatures, environmental smoke, dust or areas with strong odours (eg, perfumes). Sputum induction at visit 2 can be rescheduled by up to 3 days; at all subsequent visits, it should not be repeated if the subject is unable to produce sputum or if the sputum sample quality is not acceptable. If chest CT and MRI (where applicable) cannot be performed on the day of the scheduled visit, the imaging assessment can be rescheduled up to 14 days after the visit. If a patient experiences a COPD exacerbation or respiratory tract infection during the screening period, visit 2 will be postponed until 4 weeks following recovery from the exacerbation or infection, and the screening period may be extended up to 8 weeks.

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Table 1: FOOTPRINTS study participants: all enrolled subjects

<table>
<thead>
<tr>
<th>Patient group</th>
<th>Rationale</th>
</tr>
</thead>
<tbody>
<tr>
<td>383 patients with COPD, including:</td>
<td>Included to provide data on increased lung protease levels and the contribution that this may have on the development and progression of emphysema in the absence of A1ATD.</td>
</tr>
<tr>
<td>▶ 123 patients with mild COPD*</td>
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<tr>
<td>▶ 130 patients with moderate COPD†</td>
<td></td>
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<tr>
<td>▶ 130 patients with severe COPD‡</td>
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<tr>
<td>18 patients with COPD and A1ATD§</td>
<td>Included as they present with an earlier onset of emphysema and a faster decline in lung function, as well as a faster change in lung density.</td>
</tr>
<tr>
<td>62 ex-smokers without airflow limitation</td>
<td>Included to provide a comparison for biomarkers between subjects with and without airflow limitation. Subjects were required to be healthy based on a complete medical history and to have: FEV₁ ≥80% of predicted normal and post-bronchodilator. FEV₁/forced vital capacity ≥lower limit of normal. A mean post DLco ≥70% of predicted normal at visit 1.</td>
</tr>
</tbody>
</table>

Total enrolled: 463 subjects

*Defined as GOLD stage 1, FEV₁ ≥80% predicted.
†Defined as GOLD stage 2, FEV₁ ≥50–<80% predicted.
‡Defined as GOLD stage 3, FEV₁ ≥30–<50% predicted.
§Defined as having a diagnosis of COPD and a documented A1ATD of ZZ genotype prior to visit 2.
A1ATD, alpha1-antitrypsin deficiency; COPD, chronic obstructive pulmonary disease; DLco, diffusing capacity of the lungs for carbon monoxide; FEV₁, forced expiratory volume in 1 s; GOLD, Global Initiative for Chronic Obstructive Lung Disease; ZZ, ZZ allele.

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Table 2  Summary of biomarker assessments

<table>
<thead>
<tr>
<th>Variables</th>
<th>Biomarker assessment</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Pre-specified soluble biomarkers</strong></td>
<td></td>
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<tr>
<td>Blood</td>
<td>▶ Total and differential cell counts.</td>
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<tr>
<td></td>
<td>▶ DNA analyses (to determine A1AT genotype).</td>
</tr>
<tr>
<td><strong>Induced sputum</strong></td>
<td>▶ Neutrophil elastase activity.</td>
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<tr>
<td></td>
<td>▶ Proteinase 3 activity.</td>
</tr>
<tr>
<td></td>
<td>▶ Cat G activity.</td>
</tr>
<tr>
<td><strong>Serum/plasma</strong></td>
<td>▶ Tissue turnover biomarkers (including neutrophil elastase-specific elastin fragment).</td>
</tr>
<tr>
<td></td>
<td>▶ Other protease-generated neoepitopes (including Cat S cleaved decorin).</td>
</tr>
<tr>
<td></td>
<td>▶ Soluble form of receptor for advanced glycation end products.</td>
</tr>
<tr>
<td></td>
<td>▶ Surfactant protein D.</td>
</tr>
<tr>
<td></td>
<td>▶ Lysyl oxidase-like 2.</td>
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<tr>
<td></td>
<td>▶ Biomarkers of systemic inflammation (including IL-6, high-sensitivity C reactive protein, white cell count and fibrinogen).</td>
</tr>
<tr>
<td><strong>Imaging biomarkers</strong></td>
<td></td>
</tr>
<tr>
<td><strong>CT scanning</strong></td>
<td>▶ Airway morphology</td>
</tr>
<tr>
<td></td>
<td>▶ Mean lumen diameter.</td>
</tr>
<tr>
<td></td>
<td>▶ % wall area.</td>
</tr>
<tr>
<td></td>
<td>▶ Mean segmental and subsegmental airway wall thickness.</td>
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<tr>
<td></td>
<td>▶ Square root of wall area of bronchus with internal perimeter of 10 mm (pi10).</td>
</tr>
<tr>
<td></td>
<td>▶ Emphysema (on inspiratory CT)</td>
</tr>
<tr>
<td></td>
<td>▶ PD15 of the CT histogram.</td>
</tr>
<tr>
<td></td>
<td>▶ PD15 adjusted for inspired lung volume.</td>
</tr>
<tr>
<td></td>
<td>▶ % lung voxels with attenuation ≤–950 HU (LAA-950).</td>
</tr>
<tr>
<td></td>
<td>▶ Air trapping (on expiratory CT)</td>
</tr>
<tr>
<td></td>
<td>▶ % lung voxels with attenuation ≤–856 HU (LAA-856).</td>
</tr>
<tr>
<td></td>
<td>▶ Registration-based parenchyma measurements</td>
</tr>
<tr>
<td></td>
<td>▶ % normal lung, % emphysema and % small airway disease.</td>
</tr>
<tr>
<td><strong>MRI</strong></td>
<td>▶ Conducted at 1.5 Tesla in a subset of subjects without airflow limitation and in patients with COPD and without A1ATD to assess functional lung parameters, including pulmonary blood flow.</td>
</tr>
<tr>
<td><strong>Clinical parameters/assessments</strong></td>
<td></td>
</tr>
<tr>
<td></td>
<td>▶ Spirometry to assess FEV1 and FVC.</td>
</tr>
<tr>
<td></td>
<td>▶ DLco and DLco per unit of alveolar volume.</td>
</tr>
<tr>
<td></td>
<td>▶ Body plethysmography (functional residual capacity, total lung capacity, inspiratory capacity, inspiratory vital capacity, residual volume and expiratory reserve volume).</td>
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<tr>
<td></td>
<td>▶ Pulse oximetry.</td>
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<tr>
<td></td>
<td>▶ Symptom questionnaires (modified Medical Research Council dyspnoea scale, COPD Assessment Test, St. George’s Respiratory Questionnaire).</td>
</tr>
<tr>
<td></td>
<td>▶ 6-min walk test.</td>
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<tr>
<td></td>
<td>▶ Body mass index and BODE index.</td>
</tr>
<tr>
<td></td>
<td>▶ Adverse events and MESI.</td>
</tr>
</tbody>
</table>

*Collected at selected sites.

A1AT, alpha1-antitrypsin; A1ATD, alpha1-antitrypsin deficiency; BODE, Body mass index, airway Obstruction, Dyspnea, and Exercise; Cat, cathepsin; COPD, chronic obstructive pulmonary disease; DLco, diffusing capacity of the lungs for carbon monoxide; FEV1, forced expiratory volume in 1 s; FVC, forced vital capacity; HU, Hounsfield Units; IL-6, interleukin-6; LAA, low attenuation areas; MESI, medical events of special interest; PD15, lung density at the 15th percentile point of the CT histogram.

**Statistical analysis**

Biomarker assessments and clinical outcomes measures are evaluated by regression models for repeated measures using respective point estimates and confidence interval estimates. The correlation between individual biomarkers, and between biomarkers and other clinical outcome measures, is being assessed. Multivariate statistical prediction models for disease progression based on biomarker values and COPD phenotypes are being developed and examined. Interim analyses will also be conducted by the trial team once data from the 52-week and 104-week assessments are available for ≥70% of subjects entered into the study. The purpose of this is to provide information on the use of MRI and soluble biomarker assessments.

**Information on steering committee**

A scientific steering committee, Emphysema Progression Biomarker Study (FOOTPRINTS study) Steering Committee, was established in January 2017 to discuss and evaluate key scientific data in the field of respiratory
medicine and to provide independent advice and recommendations to optimise and strengthen the FOOTPRINTS biomarker study. The steering committee comprises 13 external and internal scientific experts in respiratory medicine and chest radiology (online supplemental table 5) and is conducted in adherence to industry regulations on the cooperation of the pharmaceutical industry with the medical professions.27–29

Data collection and management
An interactive response technology system was used to track subject enrolment and to register clinic visits and phone calls. A central laboratory facility is handling all standard safety blood laboratory analyses. Spirometry is performed using an external vendor’s equipment to allow for centralised readings. ECGs are recorded for about a 10-s duration after the subject has rested for at least 5 min in a supine position, with electrode placement performed according to the Wilson method (precordial leads) and Goldberger and Einthoven (limb leads). Pulse oximetry will be performed to measure oxygen saturation; all recordings will be made using the same pulse oximetry measurement device per site on the same finger of the subject. Body plethysmography, diffusing capacity of the lungs for carbon monoxide (DLco), the 6-min walk test (6MWT) and sputum induction and processing (at selected sites only) are performed according to internationally accepted guidelines as follows. Body plethysmography and DLco measurement are being performed according to American Thoracic Society (ATS)/European Respiratory Society guidelines, after salbutamol administration.30 The 6MWT is being conducted according to ATS guidelines.31 Inspiratory and expiratory CT scans are being performed according to the specifications provided in online supplemental table 6. Sputum induction and processing are performed according to the sputum manual issued by the vendor for sputum assessment and analysis. All samples collected for pre-specified biomarker assessments are prepared by the site according to instructions given in the investigator site file until shipment to the central contract research organisation. Additional blood samples (whole blood, plasma and serum) will be taken at visits 2–7 for future unspecified analyses, while potential leftovers from pre-specified blood analysis and induced sputum will also be stored and used for this purpose; these samples will be stored for up to 15 years after the final study report has been signed. Biomarkers are analysed by the sponsor or contractors of the sponsor. Data management and statistical evaluation are conducted by the sponsor or contractors of the sponsor. Source documents are filed at the investigator’s site; these data are transferred to the electronic case report form. Missing data may be imputed if necessary.

A clinical trial monitor, appointed by the sponsor, ensures the good running of the study and directs the clinical study team in the preparation, conduct and reporting of the study, in accordance with the sponsor’s standardised operating procedures, GCP guidelines and current legislation.

Medical information obtained for individual subjects during the study is considered confidential, and this will be ensured by using subject identification code numbers. Further information on confidentiality is detailed in the informed consent form (online supplemental file 2).

Patient and public involvement
Patients or the public were not involved in the design, conduct, reporting or dissemination plans of our research.

ETHICS AND DISSEMINATION
This study is being performed in accordance with the International Conference on Harmonisation Tripartite Guideline for GCP, the Japanese GCP regulations and local regulations. The study protocol has received approvals from the following institutional ethics committees (IECs)/institutional review boards (IRBs): Belgium: Ethische commissie onderzoek UZ/KU Leuven; Canada: Hamilton Integrated Research Ethics Board, Ontario; Research Ethics Board University of Cardiology and Pneumology Institute of Quebec; Health Research Ethics Board, University of Alberta; University of Saskatchewan Biomedical Research Ethics Board; Denmark: De Videnskabsetiske Komitéer for Region Hovedstaden, Hillerød; Finland: Varsinais-Suomen sairaanhoitopiirin Ettätiin toimitunta; Germany: Ethikkommission Landesärztekammer Hessen, Frankfurt; Ethikkommission Schleswig-Holstein, Bad Segeberg; Ethikkommission der Medizinischen Hochschule Hannover, Hannover; Japan: the IRB of Osaka City University Hospital, Osaka; the IRB of Kishiwada City Hospital, Kishiwada; the IRB of Showa University Hospital, Shinagawa; the IRB of Kagoshima University Hospital, Kagoshima; Republic of Korea: The Catholic University of Korea, Eunpyeong St. Mary’s Hospital IRB, Seoul; Seoul Metropolitan Government Seoul National University Boramea Medical; Korea University Guro Hospital IRB, Seoul; Konkuk University Medical Center IRB, Seoul; Poland: Bioethics Committee at Regional Medical Chamber, Białystok; Spain: CEIm Hospital Clínico, Barcelona; CEICIllesBalears, Palma; CEIC Hospital Universitari Vall d’Hebrón, Barcelona; CEIC-Parc de Salut Mar, Barcelona; CEIC–Hospital de Bellvitge, Barcelona; Sweden: Regionala etikprövningsnämnden i Lund, Lund; UK: North West – Greater Manchester East Research Ethics Committee; USA: University of Alabama at Birmingham Western Institutional Review Board, Puyallup; Baylor College of Medicine Institutional Review Board for Baylor, Houston; University of California San Diego UCSD Human Research Protection Program, La Jolla; Temple University Hospital Western Institutional Review Board, Puyallup; University of Utah Health Sciences Center University of Utah IRB, Salt Lake City; University of California Los Angeles Western Institutional Review Board, Puyallup; Chesapeake IRB, Columbia; Advarra, Inc, Columbia; University of Iowa Human
Subjects Office/Institutional Review Board, Iowa City; Johns Hopkins Institutional Review Board Reed Hall – B 130, Baltimore; National Jewish Health Western Institutional Review Board, Puyallup.

In the respective countries, the ethics approval preceded the study participant’s enrolment. Written informed consent is obtained from all study participants. Notably, any protocol amendments will be initiated only after all required legal documentation has been reviewed and/or approved by the respective IRB/IEC, as applicable.

The study findings will be reported in due course at research conferences and in peer-reviewed journals. The study protocol and clinical study data will be available to be shared after publication of the primary manuscript in a peer-reviewed journal. These data will be available on reasonable request; further details will be provided in the primary manuscript.

**DISCUSSION**

COPD is characterised by progressive airflow limitation and a decline in lung function. It has also been demonstrated that emphysema, a major cause of airflow limitation in COPD, is independently associated with a rapid decline in lung function. Current pharmacological treatment options for COPD and emphysema do not prevent the progressive decline of FEV1, the most commonly used marker of disease severity and progression in COPD. However, FEV1 is poorly correlated with symptoms and some other measures of disease progression.

The heterogeneity of COPD extends to a molecular level, a concept familiar in asthma, where endotype-directed therapy is now well accepted; hence, there is a need for biomarkers that can assist with diagnosis, risk stratification and assessment of therapeutic interventions. A recent systematic review by Fermont et al reported that biomarkers and clinical parameters including 6MWT, fibrinogen, C reactive protein, white cell count and interleukin-6, are associated with clinical outcomes in COPD and worth further investigation. A systematic review of CT density as a radiological biomarker came to similar conclusions. It is anticipated that biomarkers may facilitate a better understanding of the prognostic of COPD, help to guide treatment options and allow for a greater degree of precision medicine as treatment options.

So far, few studies have addressed the potential of biomarkers to predict the longitudinal outcomes of COPD, and this has been assessed for a limited number of biomarkers only. The COPDGene study, which has enrolled over 10,000 smoking (with or without COPD) and non-smoking subjects, is predominantly focused on identifying genetic determinants for COPD susceptibility, while the UK-based British Lung Foundation early COPD study is currently enrolling smokers with normal lung function or mild lung function abnormalities to assess the very early stages of COPD development. The Evaluation of COPD Longitudinally to Identify Predictive Surrogate End-points (ECLIPSE) study and the Subpopulations and Intermediate Outcomes in COPD Study (SPIROMICS) have enrolled subjects with COPD and smoking/non-smoking control subjects, similar to the COPDGene study. Other longitudinal studies include the German COPD and SYstemic consequences-Comorbidities NETwork (COSYCONET) study and the Canadian Cohort Obstructive Lung Disease (CanCOLD) study. All of these studies plan to analyse a range of soluble biomarkers and clinical outcomes and conduct CT imaging to help identify COPD phenotypes.

This prospective, longitudinal study was designed to investigate, over a 3-year period, biomarkers in different biofluids, imaging biomarkers and clinical parameters that may be associated with emphysema. A key strength of the FOOTPRINTS study is that it actively recruited a subset of patients with A1ATD. This is an important but rarely studied subpopulation of patients, which has not been actively recruited for in other longitudinal studies, such as the COPDGene, ECLIPSE, SPIROMICS and UK early COPD cohort studies. Patients with A1ATD present with earlier onset of emphysema and could help us to better understand possible markers for emphysema progression, in particular the role of NSP imbalance conditional on A1AT levels. While A1ATD is associated with a higher risk of emphysema and COPD, increased levels of serum A1AT occur in response to inflammation. Interestingly, increased levels of A1AT have also been associated with worse outcomes in patients with COPD. A recent analysis of data from the Hokkaido COPD cohort study examined the association of circulating A1AT levels with the clinical course of COPD patients without A1ATD and reported that higher A1AT levels were associated with worse outcomes, including more emphysema, a worse systemic inflammation status and higher 10-year mortality. Our findings in patients with A1ATD could help us to further understand the complex relationship between A1AT levels and COPD disease progression.

Compared with the COPDGene study, in which assessments were scheduled at baseline, 5 years and 10 years (ongoing), the FOOTPRINTS study has more frequent visits and biomarker sample collections, similar to the ECLIPSE and SPIROMICS studies. In particular, the regular collection of sputum is a major asset of the FOOTPRINTS study; sputum collection is scheduled to occur more frequently than in both the COPDGene and SPIROMICS studies and with a similar frequency to the ECLIPSE study, with four collections scheduled in the first year of the study and subsequent collections scheduled at year 2 and year 3. Another key strength and differentiator of the FOOTPRINTS study is that annual chest CT scans are being performed; compared with the other longitudinal studies discussed here, and to the best of our knowledge, the FOOTPRINTS study will provide the most regular CT imaging data. This, in addition to the frequent biomarker sampling, will allow for more robust measures of short-to-medium-term decline, critical for informing clinical trials of emphysema-targeted
medicinal products. The FOOTPRINTS study is also assessing a wider range of non-genetic biomarkers and clinical parameters across the entire study population compared with the COPDGene study, which is focused primarily on genetic analyses of DNA samples. For example, the FOOTPRINTS study is assessing NE, PR3 and Cat G activity, which could help us to further understand the role of these biomarkers in patients with COPD, including those with A1ATD and poorer disease prognosis. In addition, FOOTPRINTS is conducting MRI assessments, which could be valuable in improving our understanding of changes in vasculature and perfusion in the lung that may precede changes in emphysema. Due to the small number of patients included in the subset analysis and the challenges associated with obtaining meaningful MRI data, these findings are expected to be limited.

Although the FOOTPRINTS study is not focusing on frequent exacerbators, it is anticipated that a considerable number of adverse events, MESI and hospital admissions will be detected during the 3 years of follow-up. The wide range of biomarkers measured in sputum and blood may help identify subjects at risk for such events. In addition, the longitudinal biosampling in FOOTPRINTS will offer a unique opportunity to study the impact of exacerbations on the evolution of pathogenic processes.

It is anticipated that the results of the FOOTPRINTS study will complement, as well as expand on, the data generated in other longitudinal studies, such as COPDGene, ECLIPSE and SPIROMICS. It is envisaged that the results of this study will increase our understanding of COPD phenotypes and the underlying pathophysiology of emphysema progression, which may be of use when developing drugs to reduce COPD advancement. In addition, biomarkers associated with ongoing emphysema destruction of lung parenchyma may be identified, which could assist in predicting and monitoring patients’ COPD disease course.

A potential limitation of the FOOTPRINTS study is that sputum collection and MRI only occur at certain study sites. As such, the smaller subset of subjects may not be sufficiently large to represent the entire population. In addition, potential for selection bias at site and sufficient size to represent the entire population. In sites. As such, the smaller subset of subjects may not be.

In summary, biomarkers may help to characterise patients with COPD, allow for better monitoring and prediction of the disease course and enable an increased understanding of COPD itself. Subsequently, this should help us to develop drugs to reduce disease progression. The FOOTPRINTS longitudinal study is investigating biomarkers in biofluids, imaging biomarkers and clinical parameters associated with emphysema over a 3-year period to increase the understanding of COPD patient phenotypes, pathogenesis and disease progression.

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