

SPECT imaging of Lysyl Oxidase-Like 2 in a model of idiopathic pulmonary fibrosis

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1	SPECT imaging of Lysyl Oxidase-Like 2 in a model of idiopathic pulmonary fibrosis
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- 1 Abstract
- 2
- Non-invasive imaging of idiopathic pulmonary fibrosis (IPF) remains a challenge. The aim of this study was to
 develop an antibody-based radiotracer targeting Lysyl Oxidase-like 2 (LOXL2), an enzyme involved in the
- 5 fibrogenesis process, for SPECT/CT imaging of pulmonary fibrosis.
- 6 The bifunctional chelator DOTAGA-PEG₄-NH₂ was chemoenzymatically conjugated to the murine antibody
- 7 AB0023 using microbial transglutaminase, resulting in a degree of labeling (number of chelators per antibody)
- 8 of 2.3. Bio-layer interferometry (BLI) confirmed that the binding affinity of DOTAGA-AB0023 to LOXL2 was
- 9 preserved with a dissociation constant of 2.45 ± 0.04 nM. DOTAGA-AB0023 was then labeled with ¹¹¹In and *in*
- *vivo* experiments were carried out in a mice model of progressive pulmonary fibrosis induced by intratracheal
 administration of bleomycin. [¹¹¹In]In-DOTAGA-AB0023 was injected in three groups of mice (control, fibrotic
- 12 and treated with nintedanib). SPECT/CT images were recorded over 4 days p.i. and an *ex vivo* biodistribution
- 13 study was performed by gamma counting. A significant accumulation of the tracer in the lungs of the fibrotic
- 14 mice at D18 post-bleomycin. Interestingly, the tracer uptake was found selectively upregulated in fibrotic lesions
- 15 observed on CT scans. Images of mice that received the antifibrotic drug nintedanib from D8 up to D18 showed
- 16 a decrease in [¹¹¹In]In-DOTAGA-AB0023 lung uptake associated with a decrease in pulmonary fibrosis
- 17 measured by CT scan.
- 18 In conclusion, we report the first radioimmunotracer targeting the protein LOXL2 for nuclear imaging of IPF.
- 19 The tracer showed promising results in a preclinical model of bleomycin-induced pulmonary fibrosis, with high
- 20 lung uptake in fibrotic areas and accounted for the antifibrotic activity of nintedanib.
- 21
- 22 Keywords
- 23 Idiopathic Pulmonary Fibrosis, Lysyl Oxidase-Like 2, SPECT/CT, DOTAGA, Site-specific conjugation
- 24
- 25
- 26

Table of Contents Graphic



1 Introduction

2

3 Idiopathic pulmonary fibrosis (IPF) is a chronic and irreversible lung disease characterized by an alteration of

- 4 the pulmonary architecture, caused by the excessive and progressive deposition of extracellular matrix (ECM) in
- 5 the lung parenchyma [1]. This aberrant accumulation of ECM leads to the destruction of the lung structure and
- 6 ultimately to a decline in lung function. Although this disease is rare, with an estimated incidence of 0.09 to 1.30
- 7 per 10,000 patients worldwide, its prevalence is increasing, and its prognosis is poor, with a median patient
- 8 survival time of five years after diagnosis [2]. Therapeutic options remain limited to pirfenidone and nintedanib
- 9 which only delay the progression of the disease without stopping it [3].
- Despite significant progress, IPF diagnosis is still difficult. While a standard high-resolution computed tomography (HRCT) can confirm the diagnosis of IPF in cases where an interstitial pneumonia (UIP) pattern is present, the majority of patients may need a lung biopsy, which carries a high risk of morbidity and mortality [4]. Additionally, one of the primary imaging criteria for UIP, honeycombing, indicates an advanced stage of the illness. However, early diagnosis is important in IPF because delayed diagnosis is linked to worse outcomes and higher mortality [5]. In order to support early diagnosis, which could significantly reduce existing treatment
- 16 delays, imaging biomarkers for early/active fibrosis are required.
- 17 Major clinical problems also include the absence of instruments to track the progression of the disease and the 18 effectiveness of anti-fibrotic therapy. IPF progression is unpredictable; whereas some patients decline quickly 19 and pass away within months, others experience a gradual decline and limited progression. Additionally, 20 considering how differently IPF patients respond to pirfenidone and nintedanib from one another, those who do 21 not respond to one drug may benefit from the other. The early prediction of treatment success is a crucial 22 therapeutic concern given the potentially crippling adverse effects linked to these treatments. The first step 23 towards a tailored management of IPF patients would therefore be to develop novel imaging predictive 24 biomarkers to predict/monitor fibrosis progression and treatment success.
- 25 Lysyl Oxidase-like 2 (LOXL2), a copper-dependent amine oxidase, plays a key role in a wide range of fibrotic 26 disorders, including pulmonary fibrosis. LOXL2 catalyzes the post-translational oxidative deamination of the ε -27 amino group of lysine in collagen and other proteins of the extracellular matrix into an aldehyde group (allysine). 28 This aldehyde can react with other lysine or allysine residues in nearby proteins, allowing cross-linking of 29 collagen to take place [6,7]. A number of studies have shown that the expression of LOXL2 in pulmonary 30 fibroblasts increases drastically in case of fibrosis promoting the development of the disease [8–12]. Preclinical 31 studies, conducted with a murine monoclonal antibody, AB0023, have shown that LOXL2 blockade inhibits 32 partially LOXL2 activity, leading to a decrease in ECM production and reduced bleomycin-induced pulmonary 33 fibrosis in vivo [11]. In addition, LOXL2 was found increased in fibrotic tissues and in sera from patients with 34 IPF in association with increased risk for disease progression [7]. Further, an upregulation of LOXL2 has been 35 shown in active area, fibroblastic foci, of diseased IPF lung tissue, with relatively low expression in healthy lung 36 tissue. As a consequence LOXL2 is considered a major driver of fibrosis [11,13]. Therefore, simtuzumab, a 37 humanized monoclonal antibody targeting LOXL2, was evaluated in phase 2 clinical trials [11,14–17]. 38 Unfortunately, simtuzumab failed at clinical stage, with no significant therapeutic efficacy observed on patients 39 [18–21]. These results are yet not fully explained, but may be related to insufficient inhibitory potency of the 40 antibody or an inadequate selection of patients included in the trials.

1 The protein LOXL2 has not yet been explored as an imaging target. Wuest et al. reported in 2015 a [¹⁸F]F-2 labeled tracer targeting Lysyl Oxidase (LOX), developed from an oligopeptide substrate of this enzyme (with a 3 Km in the sub-millimolar range) [22]. However, this tracer is a pan-LOX substrate, and has shown rapid washout 4 from LOX-expressing tissues and some degree of non-specific binding. A LOXL2-specific radiotracer could 5 provide valuable information to facilitate in the staging of IPF patients, monitoring disease progression and 6 response to treatments. In addition, it could improve the selection of patients most likely to respond to new 7 LOXL2 inhibitors currently in development [23-26]. Herein, we report the development and preclinical 8 evaluation of a radiotracer targeting LOXL2, based on the murine antibody AB0023. The antibody was 9 conjugated to the metal chelator DOTAGA, radiolabeled with ¹¹¹In and evaluated in a mice model of pulmonary 10 fibrosis, induced by intratracheal administration of bleomycin, by SPECT/CT and biodistribution studies [27]. 11

12

13 Materials and methods

14 Reagents and equipment specifications

AB0023 was produced by Proteogenix (France) and Biointron (China). DOTAGA-PEG₄-NH₂ was kindly given by Chematech (France). PNGase F was supplied by Promega (USA) in 20 mM Tris-HCl (pH 7.5), 50 mM NaCl and 5 mM EDTA at a concentration of 10 000 U/mL. mTGase was provided by Zedira (Germany) in PBS (Phosphate Buffer Saline) 0.1 M, pH 7.4, at a concentration of 280.8 U/mL. Recombinant human LOXL2 (fused with a polyhistidine tag at the C-terminus) was purchased from Sino Biological (China) and was reconstituted in ultrapure water.

21

22 HPLC analyses were conducted on a Thermo Scientific MAbPac column (4 µm, 50 x 2.1 mm) with the 23 following eluents: A: H₂O + 0.08% FA + 0.02% TFA and B: MeCN + 0.08% FA + 0.02% TFA and a gradient 24 program: 20% of B for 0.5 min, 20% to 55% of B in 5.5 min, 55% to 80% in 0.5 min, 80% of B for 1 min, 80% 25 to 20% in 0.1 min, 20% of B for 2.5 min, at a flow rate of 0.5 mL/min at 80 °C. High-resolution mass 26 spectrometry analyses were recorded on a Orbitrap Exploris 240 mass spectrometer (Thermo Scientific) 27 equipped with an electrospray ionization source (H-ESI II). Spectra were deconvoluted on BioPharma Finder 28 software (Thermo Scientific). Ultrapure water produced by a PURELAB Ultra system from ELGA was used 29 throughout (18.2 MΩ.cm). UV spectroscopy was done on Mettler Toledo UV/Vis excellence UV5Nano.

30

Size-exclusion chromatography (SEC-HPLC) analyses were performed on a JASCO HPLC system LC-2000 analytical series equipped with a Superdex 200 5/150 GL column. Instant thin-layer chromatography (iTLC or radio-TLC) was performed using sheets impregnated with silica gel (iTLC-SG; Agilent), eluted with 0.1 M citrate buffer pH 6.0, and analyzed on an AR-2000 radio-TLC plate reader (Bioscan Inc.).

35

36 Synthesis of the bioconjugate

37 AB0023 was first deglycosylated using PNGase F. 2 µL of PNGase F (20 U) was added to 500 µg of AB0023

- 38 (3.4 nmol, in solution at 3.5 g/L in PBS pH 7.4). The mixture was incubated 8 h at 37°C in a Thermomixer (700
- 39 rpm). PNGase F was removed from the mixture by ultrafiltration on an Amicon Ultracel-50 kDa (Merck-
- 40 Millipore) and the purified solution was centrifuged (4000 rpm, 20 min, 4 °C) to concentrate the product at 9.0

- $1 \qquad g/L. \ To \ perform \ the \ bioconjugation, \ DOTAGA-PEG_4-NH_2 \ (89 \ equivalents, \ 246 \ nmol, \ 4.1 \ \mu L \ of \ a \ 60 \ mM \ stock$
- 2 solution in PBS pH 7.4) was added to 400 μ g of deglycosylated AB0023 (1 equivalent, 2.76 nmol, in solution at
- 3 9.0 g/L in PBS pH 7.4) followed by 4.7 μL (1.32 U) of mTGase (280.8 U/mL). The reaction mixture was
- 4 incubated 6 h at 37 °C on a Thermomixer (700 rpm) and monitored by mass spectrometry. The bioconjugate was
- 5 purified by SEC-FPLC on an Äkta Pure 25 M chromatography system (GE Healthcare Life Sciences) equipped
- 6 with a Superdex 200 10/300 GL column, using AcONH₄ 0.1 M pH 5.4 as eluant. This eluant allowed the
- 7 bioconjugate to be isolated directly in the radiolabeling buffer without further buffer exchange. The protein yield
- 8 was calculated by measuring the absorbance at 280 nm by UV spectroscopy ($\varepsilon_{280} = 1.4 \text{ mL.mg}^{-1}.\text{cm}^{-1}$).
- 9

10 **BLI measurements**

- 11 The affinity of AB0023 for rhLOXL2 was studied by biolayer interferometry (BLI) on a Sartorius Octet[®] R2
- 12 system. All experiments were performed at 30 °C with 1000 rpm shaking. Ni-NTA Dip and Read[™] (Sartorius)
- 13 biosensors were employed. LOXL2 was diluted in PBS to obtain a final concentration of 5 μ g/mL. LOXL2 and
- 14 AB0023 were used as the ligand and the analyte respectively in the experiments. PBS was used as control.
- 15 Biosensors were first dipped into wells containing PBS during 10 min. The loading time of LOXL2 on the
- 16 biosensors was 600 s. Two washes of the biosensors with PBS during 60 s were performed before the association
- 17 step. Association time of AB0023 to LOXL2 was set at 600 s. Dissociation time was set at 900 s in PBS.
- 18 Raw sensorgrams were fitted using a 1:2 bivalent analyte model in OctetTM data analysis HT software v 12.2.
- 19 Data were corrected with an alignment of the Y axis to the average of baseline step, an alignment of the inter-
- 20 step to the association step and a Savitzky-Golay fittering.
- 21

22 Radiochemistry

- 476 μ L of a [¹¹¹In]InCl₃ solution at 840 MBq/mL (400 MBq) was added to 708 μ g of bioconjugate (1.9 mg/mL in AcONH₄ 0.1M pH 5.4). 47.6 μ L of AcONH₄ 1 M was further added to neutralize the acidity of the [¹¹¹In]InCl₃ solution. The mixture was incubated 1 h at 37 °C in a Thermomixer (700 rpm). The reaction was monitored by radio-TLC. Unincorporated ¹¹¹In was scavenged by adding 47 μ L of a solution of 50 mM EDTA in 0.1 M AcONH₄ pH 5.4. No further purification was needed. The final radiolabeling purity was measured by radio-TLC.
- 28

29 Animal experiments

- 30 All animal studies were conducted in accordance with the legislation on the use of laboratory animals (directive
- 31 2010/63/EU) and were approved by accredited Ethical committee (C2ea Grand Campus n°105) and the French
- 32 Ministries of Research and Agriculture (project #12816).
- 33 Eight-week-old C57/Bl6 mice received at D0 a single intratracheal injection of 2 mg/kg of bleomycin (BLM,
- 34 Santa Cruz biotechnology, USA) or NaCl (Control) under anesthesia (3% isoflurane). BLM-induced lung
- 35 fibrosis is characterized by an initial inflammatory phase lasting from D0 to D7 which moves on to a progressive
- 36 fibrotic phase from D7 to D23. BLM instillation was performed by a unique expert operator. Animal weights
- 37 were monitored daily from D0 to D7 in order to select mice that responded to BLM (selection of animals with at
- 38 least 10% weight loss from D0 to D7). For each experiment, approximately 20% of animals were excluded.
- 39 When indicated, animals were treated with nintedanib (OfevTM, 60mg/kg) by daily gavage during the fibrotic
- 40 phase of BLM-induced lung fibrosis from D7 to D18 as recommended [28].

2 In vivo imaging

- 3 One pilot study was performed on NaCl- and BLM-receiving mice at D18 post-BLM by SPECT/CT imaging
- 4 with [¹¹¹In]In-DOTAGA-AB0023 (NaCl n=6, BLM D18 n=6). Mice were anesthetized through isoflurane
- 5 (1.5%) inhalation for intravenous injection (tail vein) of $[^{111}In]In$ -DOTAGA-AB0023 (10 MBq, 25 µg/mouse).
- Mice were then maintained under anesthesia (1.5% isoflurane) and placed on an imaging heated bed inside a
 NanoSPECTTM-CT (Mediso, Hungary). A CT scan of a lung-centered region was obtained (500 ms, 45 kV, 180
- 8 projections, pitch 1, binning 1:4) followed by SPECT acquisition with 90–120 s per projection frame resulting in
- 9 acquisition times of 45–60 min of the same region or a PET acquisition (30 minutes) of the same region (lung-
- 10 centered, 250-700 keV). Images were taken 24 h, 48 h, 72 h and 96 h post-injection (p.i.) of [¹¹¹In]In-DOTAGA-
- 11 AB0023.
- 12 In another experiment, imaging of lung fibrosis was performed on NaCl- (n=6) and BLM-receiving mice treated
- 13 with vehicle (n=6) or nintedanib (n=5) by SPECT/CT with [¹¹¹In]In-DOTAGA-AB0023 at D18 following the
- 14 same imaging protocol as described above.
- 15 For each experiment, after the last imaging (96 h p.i.), mice were sacrificed and major organs (blood, bladder,
- 16 kidneys, spleen, liver, heart, muscle and lungs) were harvested for *ex vivo* quantification using a γ -counter
- 17 (Wizard, Perkin Elmer).
- 18

19 Image analysis

- All SPECT/CT fusion images were obtained using the VivoQuantTM software (Invicro, USA). Each image was visually interpreted and 3D regions of interest (3DROI) corresponding to the lungs were manually drawn for CT quantification and to determine their radioactivity content. Injected doses per animal were measured at the time of injection in MBq. Lung radioactivity content was expressed in MBq, converted to percentage of injected dose per gram of tissue (% ID/g). All images were decay corrected for quantification. In addition, a semi-automatic segmentation of 3DROI was performed on CT scans as follows: normal lung density (-800 to -100 HU)
- corresponding to aerated lung areas and high lung density (-100 to 300 HU) corresponding to nonaerated/fibrotic lung areas as previously described [29]. This semi-automatic segmentation allowed the independent quantification of the radioactivity content of [¹¹¹In]In-DOTAGA-AB0023 in normal (aerated) and
- 29
- 30 31

33

32 **Results**

34 Conjugation of DOTAGA to AB0023

high (non-aerated) density lung tissue respectively.

35 The recombinant murine antibody was deglycosylated using PNGase F and then conjugated to DOTAGA-PEG₄-

36 NH₂ using microbial transglutaminase (Fig. 1). DOTAGA-AB0023 was obtained with 40% protein recovery

37 yield over two steps. Native AB0023, deglycosylated AB0023 and the final bioconjugate and were analyzed by

- 38 RP-HPLC-MS (Fig. 2A, 2B and 2C). After deglycosylation, the deconvoluted spectrum revealed a major peak at
- 39 144,974 Da corresponding to deglycosylated AB0023 (Fig. 2B). Two minor peaks (145,103 Da and 145,232 Da)
- 40 were detected, corresponding to different degrees of C-terminal lysine clipping [30]. After conjugation, the
- 41 deconvoluted spectrum showed a major peak at 146,329 Da corresponding to the covalent grafting of two

- 1 DOTAGA-PEG₄-NH₂ and a degree of labeling (DOL) of 2 (Fig. 2C). The additional peaks at 145,651 Da and
- 2 147,009 Da corresponded to DOL 1 and 3 respectively. LC-MS/MS analysis of DOTAGA-AB0023 after tryptic
- 3 digestion enabled us to identify heavy chain residues Q153 and Q300 as those conjugated to DOTAGA-PEG₄-
- 4 NH_2 . The average DOL after 6 h of incubation at 37 °C was 2.3.



Fig. 1: Synthesis of the bioconjugate 1) PNGase F (20 U), PBS pH 7.4, 37 °C, 8 h; 2) DOTAGA-PEG₄-NH₂ (89 eq), mTGase (1.32 U), PBS pH 7.4, 37 °C, 6 h





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8

13 Biolayer interferometry

14 Results of the BLI measurements demonstrated high affinity binding between native AB0023 and rhLOXL2

- 15 with a K_D of 0.89 \pm 0.02 nM. Deglycosylated AB0023 and DOTAGA-AB0023 showed K_D equal to 2.43 \pm 0.08
- 16 nM and 2.45 \pm 0.04 nM respectively (Fig. 3). R² values for each measurement were >0.99.



Fig. 3: Kinetic analysis of (A) native AB0023 (B) deglycosylated AB0023 and (C) DOTAGA-AB0023

1

4 Radiolabeling

Radiolabeling yield of [¹¹¹In]In-DOTAGA-AB0023 reached 96% after 1 h at 37 °C with a specific activity equal
to 570 MBq/mg (85 MBq/nmol) (supplemental fig. 3 A). This tracer proved to be stable *in vitro* over 7 days, in
human plasma at 37°C (supplemental fig. 3 B).

8

9 In vivo imaging of LOXL2 can detect BLM-induced lung fibrosis

NaCl and BLM-receiving mice underwent SPECT/CT imaging with the LOXL2 targeting agent [¹¹¹In]In-10 DOTAGA-AB0023 at D18 (Fig. 4A). SPECT/CT images were performed 24 h, 48 h, 72 h and 96 h post-11 12 injection of [¹¹¹In]In-DOTAGA-AB0023 showing that the best imaging time to differentiate NaCl- from BLMreceiving mice was 96 h post-injection (Fig. 4AB). The lung uptake of [¹¹¹In]In-DOTAGA-AB0023 was 13 14 significantly increased in BLM-treated mice at D18 compared to non-fibrotic controls at 48 h, 72 h and 96 h (Fig. 4B). Ex vivo radioactivity quantification of the lungs confirmed the increase in [¹¹¹In]In-DOTAGA-15 16 AB0023 lung uptake 96 h p.i. in BLM-treated animals (Fig. 4C). In addition, lung to blood ratio and lung to 17 muscle ratio demonstrated a clear increase in BLM-receiving mice compared with controls (Fig. 4DE). 18 Similarly, lung CT of BLM-receiving mice showed an increase in fibrotic consolidations compared with control 19 mice with an increase in mean lung density (MLD) and a decrease of aerated areas in BLM-treated mice 20 compared with controls at D18 (Fig. 4FG). The global biodistribution of [111In]In-DOTAGA-AB0023 demonstrated that uptake was not different between NaCl- and BLM-receiving mice except in the lungs (Fig. 21 4H). Time curves of lung/blood and lung/muscle ratio confirms that [¹¹¹In]In-DOTAGA-AB0023 is significantly 22 23 increased in fibrotic lungs with an optimal signal-to-background ration at 96 h post-injection (Supplemental fig. 24 6A, B). Interestingly, discrimination of aerated and non-aerated lung areas by semi-automatic segmentation on 25 CT scans demonstrated that [¹¹¹In]In-DOTAGA-AB0023 SPECT signal was only significantly increased in non-

- 1 aerated lung areas in BLM-receiving mice compared with control mice, showing the specificity of [¹¹¹In]In-
- 2 DOTAGA-AB0023 for fibrotic areas (Fig. 4I). Interestingly, [¹¹¹In]In-DOTAGA-AB0023 lung uptake at 96 h
- 3 significantly correlated with MLD measured on CT (Fig. 4J).





Fig. 4: [¹¹¹In]In-DOTAGA-AB0023 is able to detect advanced BLM-induced lung fibrosis. A/ Representative lung
SPECT/CT images with [¹¹¹In]In-DOTAGA-AB0023 of NaCl- and BLM-receiving mice at D18 at 24h, 48h, 72h and 96h
p.i.. B/ Graph represents the [¹¹¹In]In-DOTAGA-AB0023 lung uptake in %ID/g measured on SPECT images of NaCl- and
BLM-receiving mice at D18. Results are presented as mean ± SEM, NaCl n=6, BLM n=6. Difference between groups were
compared using Mann-Whithney non-parametric t-test, **p<0.01, ***p<0.001. C/ Graph represents the [¹¹¹In]In-IDTAGA-AB0023 lung uptake in %ID/g measured on SPECT images of NaCl- and

1 Results are presented as mean ± SEM, NaCl n=6, BLM n=6. Difference between groups were compared using Mann-Whithney non-parametric t-test, **p<0.01. D/ Graph represents the lung/blood ratio of [¹¹¹In]In-DOTAGA-AB0023 2 3 uptake in %ID/g measured by gamma-counting 96h p.i. of NaCl- and BLM-receiving mice at D18. Results are presented as 4 mean ± SEM, NaCl n=6, BLM n=6. Ratios are normalized to 1 for the NaCl group. Difference between groups were 5 compared using Mann-Whithney non-parametric t-test, **p<0.01. E/ Graph represents the lung/muscle ratio of [¹¹¹In]In-6 DOTAGA-AB0023 uptake in %ID/g measured by gamma-counting 96h p.i. of NaCl- and BLM-receiving mice at D18. 7 Results are presented as mean \pm SEM, NaCl n=6, BLM n=6. Ratios are normalized to 1 for the NaCl group. Difference 8 between groups were compared using Mann-Whithney non-parametric t-test, **p<0.01. F/ Graph represents the mean lung 9 density quantified on CT images of NaCl- and BLM-receiving mice at D18. Results are presented as mean \pm SEM, NaCl n=6, 10 BLM n=6. Difference between groups were compared using Mann-Whithney non-parametric t-test, **p<0.01. G/ Graph 11 represents the percentage of aerated lung tissue measured on CT images of NaCl- and BLM-receiving mice at D18. Results 12 are presented as mean \pm SEM, NaCl n=6, BLM n=6. Difference between groups were compared using Mann-Whithney non-13 parametric t-test, **p<0.01. H/ Graph represents the [¹¹¹In]In-DOTAGA-AB0023 uptake in %ID/g measured by gamma 14 counting in blood, heart, kidney, bladder, liver, spleen, muscle and lungs of NaCl- and BLM-receiving mice at D18. 15 Difference between groups were compared using Mann-Whithney non-parametric t-test, ***p<0.001. I/ Graph represents the 16 [¹¹¹In]In-DOTAGA-AB0023 lung uptake in %ID/g of NaCl- and BLM-receiving mice at D18 in aerated and non-aerated lung 17 areas (segmented on CT images). Results are presented as mean ± SEM, NaCl n=6, BLM n=6. Stars (*) are representative of 18 comparison of each group with NaCl group and hashs (#) are representative of statistical comparison of BLM aerated group 19 with BLM non-aerated group. Difference between groups were compared using Mann-Whithney non-parametric t-test, 20 *p<0.05, **p<0.01 and ***p<0.001. J/ Correlation between mean lung densities (HU) measured on CT images and

21 [¹¹¹In]In-DOTAGA-AB0023 lung uptake (%ID/g) of corresponding lungs measured on SPECT/CT.

22 In vivo imaging of LOXL2 is a useful tool to monitor the anti-fibrotic efficacy of nintedanib

23 NaCl and BLM-receiving mice treated or not with nintedanib underwent SPECT/CT imaging with [¹¹¹In]In-DOTAGA-AB0023 at D18 (Fig. 5A). We confirmed that the lung uptake of [¹¹¹In]In-DOTAGA-AB0023 was 24 25 increased in BLM-receiving mice compared to control at D18 (Fig. 5AB). Interestingly, in BLM-receiving mice nintedanib treatment dramatically decreased [¹¹¹In]In-DOTAGA-AB0023 lung uptake at D18 (Fig. 5AB). These 26 27 results were confirmed by gamma counting at 96 h p.i. in which [¹¹¹In]In-DOTAGA-AB0023 lung uptake was 28 significantly higher in BLM-receiving mice and was decreased by nintedanib (Fig. 5C). In addition, lung to 29 blood ratio demonstrated a clear increase in BLM-receiving mice compared with BLM mice treated with 30 nintedanib and controls (Fig. 5D). Similarly, MLD measured on CT significantly increased in BLM-receiving mice and was decreased by nintedanib treatments (Fig. 5E). The global biodistribution of [¹¹¹In]In-DOTAGA-31 32 AB0023 demonstrated that uptake was not different between NaCl-, BLM-receiving mice and BLM mice treated 33 with nintedanib except in the lungs (Fig. 5F). Time curves of lung/blood and lung/muscle ratio confirm that 34 nintedanib significantly reduced [¹¹¹In]In-DOTAGA-AB0023 lung uptake (Supplemental fig. 6D, E). 35 Interestingly, immunostainings performed on a separate subset of animals demonstrated ex vivo than nintedanib 36 was able to diminish the BLM-induced upregulation of LOXL2 expression in the lungs (Supplemental Fig. 7).

- 37 Most interestingly, the analysis of publicly available RNA sequencing datasets (GSE53845) demonstrated that
- 38 LOXL2 gene expression was upregulated in IPF patients compared with healthy controls (Supplemental Fig. 8).
- 39 These findings further highlight the potential of LOXL2 as a theranostic target in IPF.





Fig. 5: Nintedanib reduces lung fibrosis which can be detected by [¹¹¹In]In-DOTAGA-AB0023 SPECT imaging A/ 2 3 Representative lung SPECT/CT images with [¹¹¹In]In-DOTAGA-AB0023 of NaCl-, BLM-receiving mice and BLM-4 receiving mice treated with nintedanib at D18 at 24h, 48h, 72h and 96h p.i.. B/ Graph represents the [¹¹¹In]In-DOTAGA-5 AB0023 lung uptake in %ID/g measured on SPECT images at 96h p.i. of NaCl- and BLM-receiving mice at D18. Results are 6 presented as mean ± SEM, NaCl n=6, BLM n=6, BLM+nintedanib n=6. Difference between groups were compared using 7 Kruskal-Wallis non-parametric ANOVA, *p<0.05, **p<0.01. C/ Graph represents the [¹¹¹In]In-DOTAGA-AB0023 lung 8 uptake in %ID/g measured by gamma-counting 96h p.i. of NaCl-, BLM-receiving mice and BLM-receiving mice treated with 9 nintedanib at D18. Results are presented as mean ± SEM. Difference between groups were compared using Kruskal-Wallis

- 1 non-parametric ANOVA, *p<0.05, ***p<0.001. D/ Graph represents the lung/blood ratio of [¹¹¹In]In-DOTAGA-AB0023
- 2 uptL2ke in %ID/g measured by gamma-counting 96h p.i. of NaCl-, BLM-receiving mice and BLM-receiving mice treated
- 3 with nintedanib at D18. Results are presented as mean ± SEM, Difference between groups were compared using Kruskal-
- 4 Wallis non-parametric ANOVA, *p<0.05, ***p<0.001. E/ Graph represents the mean lung density quantified on CT images
- 5 of NaCl-, BLM-receiving mice and BLM-receiving mice treated with nintedanib at D18. Results are presented as mean ±
- 6 SEM. Difference between groups were compared using Kruskal-Wallis non-parametric ANOVA, *p<0.05, ***p<0.001. F/
- 7 Graph represents the [¹¹¹In]In-DOTAGA-AB0023 uptake in %ID/g measured by gamma counting in blood, heart, kidney,
- 8 bladder, liver, spleen, muscle and lungs of NaCl-, BLM-receiving mice and BLM-receiving mice treated with nintedanib at
- 9 D18. Difference between groups were compared using Kruskal-Wallis non-parametric ANOVA, *p<0.05, ***p<0.001.
- 10 11

12 **Discussion**

13 14 The identification of biomarkers of disease progression and therapy efficacy has become a hot topic in the lung 15 fibrosis field in the last decade. The heterogeneous nature of IPF progression and the variability of response 16 observed in patients under anti-fibrotic treatments is a major clinical concern leading to poorer outcome. While 17 most of the attention has been focused on the discovery and evaluation of circulating biomarkers [31], in vivo 18 molecular imaging recently emerged as a valuable tool to non-invasively assess the expression of various 19 proteins involved in fibrosis progression [32]. In this context, our study aimed at developing an imaging probe 20 targeting LOXL2. LOXL2 is a major protein involved in fibrogenesis as it is responsible for the cross-linking of 21 the excessive amount of collagen in the fibrotic lung, thus leading to lung tissue stiffening and increased 22 mechanical stretch which is a contributor of disease progression [33]. Our strategy was based on the 23 bioconjugation of a murine anti-LOXL2 antibody, AB0023, with the chelator DOTAGA which has proven to be 24 a valuable alternative to the standard DOTA [34]. To achieve reproducible results and control the degree of 25 labeling, we opted for a site-specific enzymatic bioconjugation strategy using mTGase. This enzyme reacts on 26 glutamine-295 residues, forming isopeptidic bonds with amine-functionalized probes [35]. PNGase F was first 27 used to remove the glycan chains on asparagine-297 residues, leaving access to mTGase to react. Higher DOL 28 than expected appeared during conjugation step, suggesting that there are two more reactive positions on this 29 antibody. However, the conjugation of DOTAGA on AB0023 did not affect its affinity for LOXL2.

30 In clinic, the imaging of IPF is currently limited to high-resolution computed tomography (HRCT) both for 31 diagnosis and monitoring, with little impact on mortality and therapeutic decision in IPF [36-38]. Indeed, lesions 32 detected on HRCT may reflect a too advanced stage of the disease in patients for effective therapeutic 33 intervention. In addition, while HRCT may inform on the extent of lung fibrosis at a given time, it does not 34 account for activated molecular mechanisms that could be triggered for therapeutic purposes. Moreover, the 35 recent approval of pirfenidone and nintedanib implies the implementation of strategies, which are not yet 36 available, to monitor efficacy of these novel therapies in order to improve patient management and clinical 37 outcome in IPF. Despite the fact that these two molecules have been shown to reduce lung function decline and 38 to improve patients' survival in several clinical trials [39], their use is currently limited due to a high 39 interpersonal variability among IPF patients. The lack of tools to accurately monitor pirfenidone and nintedanib 40 efficacy in patients is another reason limiting their efficacy in the clinical practice. Considering the disabling side 41 effects which may be associated to these therapies, the early prediction of treatment efficacy is a major clinical

1 issue. As a consequence, identifying novel imaging predictive biomarkers to predict/monitor therapeutic efficacy 2 would be the first step towards a personalized management of IPF patients. Our results confirm previous reports 3 demonstrating an increase in LOXL2 expression in the fibrotic lungs upon BLM stimulation [11]. As expected for a full length antibody, which has slow distribution times, the optimal imaging time of [¹¹¹In]In-DOTAGA-4 5 AB0023 was between 72 h and 96 h post-injection. Our results are comparable to recent studies highlighting the 6 interest of molecular imaging to follow other biomarkers. John et al. developed a specific peptide to the integrin 7 $\alpha_V \beta_6$ (A20FMDV2) radiolabeled with ¹¹¹In to measure $\alpha_V \beta_6$ expression in BLM-treated mouse lungs by 8 SPECT/CT [40]. Overexpression of $\alpha_V \beta_6$ integrin by alveolar epithelial cells plays a crucial role in the 9 pathogenesis of pulmonary fibrosis by favoring TGF-B1 activation. Lung uptake of ¹¹¹In-A20FMDV2 was 10 increased in BLM-treated mice compared to control mice. Quantitative analysis demonstrated that SPECT 11 signals correlate with the levels of collagen, $\alpha_V \beta_6$ protein, and messenger RNA of the β_6 subunit [40]. Very 12 recently the lung uptake of the same peptide modified with fluorine-18 for PET imaging ([¹⁸F]FB-A20FMDV2) 13 has been shown to be increased in a small cohort of patients with IPF compared with healthy controls [41]. 14 Following the same concept, Désogère et al. developed a peptide-based PET probe ([⁶⁸Ga]Ga-CBP8) which 15 targets collagen type I for lung fibrosis imaging [42]. The authors demonstrate high specificity of [⁶⁸Ga]Ga-16 CBP8 for pulmonary fibrosis in vivo in bleomycin-treated mice. In addition, [⁶⁸Ga]Ga-CBP8 signals correlated 17 with lung collagen content and showed sensitivity to monitor response to treatment in bleomycin-induced lung 18 fibrosis [42]. Similarly, imaging of Fibroblast Activation Protein-a (FAP), a membrane marker of fibroblasts 19 activation, with radiolabeled quinoline-based small molecule (FAPI) has been shown to noninvasively detect 20 disease activity in a murine model of pulmonary fibrosis [43] Interestingly, PET imaging with [⁶⁸Ga]Ga-21 radiolabeled FAPI has the potential to detect the early onset of fibrotic injury to promote early diagnosis. A recent clinical study confirms these findings by demonstrating an increase in [⁶⁸Ga]Ga-FAPI lung uptake in IPF 22 23 patients compared with healthy volunteers [44]. Contrary to our study, most of the developed tracers for lung 24 fibrosis imaging rely on small peptides instead of full length antibodies. While small peptides present the 25 advantage to display a short biodistribution profile favorable for imaging purposes, they also present much lower 26 accumulation rates in targeted organs which may worsen the interpretation of images due to a weak signal-tobackground ratio. In our study, the lung accumulation of [¹¹¹In]In-DOTAGA-AB0023 in fibrotic lungs is 27 approximately three to four fold higher than in non-fibrotic lungs. In addition, accumulation of [111In]In-28 29 DOTAGA-AB0023 is approximately three-fold higher in the lungs compared with blood and muscles which are 30 not expressing LOXL2. These results demonstrate a high signal-to-background ratio allowing an accurate 31 evaluation of [¹¹¹In]In-DOTAGA-AB0023 lung accumulation in the context of fibrosis. Nevertheless, the 32 evaluation of imaging properties of smaller fragments of AB0023 (e.g. Fab or F(ab')) may be of interest in 33 future studies. Despite promising preclinical results, the inhibition of LOXL2 with simtuzumab, the humanized 34 monoclonal antibody corresponding to AB0023, failed in a large phase 2 study (RAINIER trial) that examined 35 the efficacy of simtuzumab for the treatment of IPF. Benefit of simtuzumab over placebo was not shown for the 36 endpoints of progression-free survival, all-cause hospitalizations, and mortality as well as decline in FVC % 37 predicted, diffusing capacity of the lung (DLCO) % predicted, and 6-min walk distance. This trial enrolled a 38 large number of patients (544) which were stratified based on LOXL2 serum levels to increase chances of 39 success. Nevertheless, whether the level of circulating LOXL2 is correlated with its lung expression remains to 40 be fully elucidated. In this context, imaging tools such as [¹¹¹In]In-DOTAGA-AB0023 may be of great value to

- 1 stratify patients based on LOXL2 lung expression levels to select patients more likely to respond to anti-LOXL2
- 2 therapies. This strategy may represent a first step towards precision medicine for the treatment of patients with
- 3 IPF. It has been suggested that the failure of the RAINIER trial was due to poor tissue penetration in the human
- 4 IPF lung thus leading to a non-optimal LOXL2 inhibition [45]. Our results demonstrate that [¹¹¹In]In-DOTAGA-
- 5 AB0023 was able to reach and specifically remain in the lungs in fibrotic areas where LOXL2 is most likely
- 6 overexpressed. Such tools may be of interest to assess tissue penetration of full-length antibodies such as 7 simtuzumab in human IPF lungs. Additionally, LOXL2 is only one of the family of LOXL enzymes that are
- simtuzumab in human IPF lungs. Additionally, LOXL2 is only one of the family of LOXL enzymes that are
 involved in collagen crosslinking, and the non-LOXL2 enzymes were not susceptible to selective allosteric
- 9 inhibition by simtuzumab. For instance, LOXL1 has been demonstrated to also play a crucial role in lung fibrosis
- 10 [46] suggesting that therapeutic strategies involving co-inhibition of LOXL2 with other crosslinking enzymes
- 11 such as LOXL1 may be of interest.
- Finally, our study reveals that in our preclinical model nintedanib seems to decrease lung LOXL2 expression. Clinicians are currently unable to identify responders versus non-responders before starting antifibrotic treatments nintedanib and pirfenidone, and do not know whether declining lung function during treatment indicates a lack of response to one of the drugs or whether lung function decline has been significantly attenuated and would have declined more rapidly without antifibrotic treatment. In this context, the use of an imaging agent such as [¹¹¹In]In-DOTAGA-AB0023 may represent a non-invasive strategy to longitudinally monitor efficacy of antifibrotic treatments.
- 19

1 Conclusion

2

The use of an enzymatic site-specific conjugation strategy has led to the reliable and reproducible production of the first anti-LOXL2 SPECT tracer. The radioimmunoconjugate showed promising results in a preclinical model of bleomycin-induced lung fibrosis, with a high lung uptake. The ability of AB0023 to monitor the response to nintedanib was also investigated and demonstrate encouraging outcomes. Our study offers a first proof of concept of the use of a non-invasive and accurate strategy to monitor LOXL2 expression that may help, in the future, patients' stratification for potential clinical trials assessing anti-LOXL2 strategies in IPF, a first step toward personalized management of IPF patients.

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Supporting information file

HPLC-MS analysis of reduced DOTAGA-AB0023, HPLC-MS/MS analysis of DOTAGA-AB0023 after tryptic digestion, Radio-iTLC chromatogram and *in vitro* stability in plasma of [¹¹¹In]In-DOTAGA-AB0023, Individual SPECT and CT images, Time-curves for lung/blood and lung/muscle ratios, IHC of LOXL2, LOXL2 mRNA expression in healthy controls and IPF patients.

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