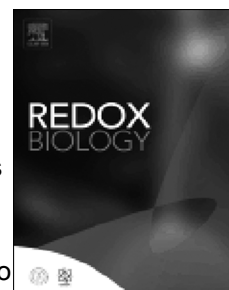# Journal Pre-proof

Monitoring ferroptosis *in vivo*: Iron-driven volatile oxidized lipids as breath biomarkers

Yuta Matsuoka, Yoshinori Katsumata, Po-sung Chu, Rei Morikawa, Nobuhiro Nakamoto, Kohta Iguchi, Ken Takahashi, Tadayuki Kou, Ryo Ito, Kojiro Taura, Shujiro Yazumi, Hiroaki Terajima, Gen Honjo, Genki Ichihara, Yuki Muramoto, Kazuki Sato, Rae Maeda, Kazuhiro Hata, Naoya Toriu, Motoko Yanagita, Masaki Tajima, Sidonia Fagarasan, Ken-ichi Yamada, Yuki Sugiura

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Abstract

Ferroptosis, an iron-dependent cell death mechanism characterized by excessive lipid peroxidation, has been implicated in numerous human diseases and organ pathologies. However, current detection methods necessitate invasive tissue sampling to assess lipid peroxidation, making noninvasive detection of ferroptosis in human subjects extremely challenging. In this study, we employed oxidative volatolomics to comprehensively characterize the volatile oxidized lipids (VOLs) produced during ferroptosis. Polyunsaturated fatty acid-derived VOLs were generated via iron-dependent LPO and released extracellularly as ferroptosis progressed. These VOLs were specifically generated during hepatic ferroptosis in mouse models of acetaminophen-induced liver injury and metabolic dysfunction-associated steatohepatitis (MASH) and were also detectable in the exhaled breath of patients with MASH. Specific VOLs released upon iron-dependent LPO are potential markers of ferroptosis *in vivo* and may facilitate noninvasive monitoring of cellular health in humans.

3

Introduction

Ferroptosis is a regulated cell death mechanism driven by iron-dependent lipid peroxidation (LPO) [1]. To date, ferroptosis has been implicated in a wide range of human diseases and organ pathologies[2], prompting numerous drug discovery studies targeting this process. To elucidate the role of ferroptosis in specific human diseases or pathological conditions, it is essential to detect LPO within damaged tissues [3]. Currently, the assessment of LPO in animal models and human tissues requires invasive tissue sampling followed by analysis using fluorescence probes or mass spectrometry. However, obtaining tissue samples from human subjects is frequently challenging, representing a major bottleneck in ferroptosis research.

During ferroptosis induction, LPO progresses within cellular membranes, leading to the fragmentation of membrane lipids. This produces volatile oxidized lipids (VOLs) that diffuse from cells and tissues, enter the circulation, and are released as part of exhaled breath [4]. Accelerated lipid fragmentation can result in plasma membrane rupture and ferroptosis[5-7]. Given that VOLs are generated alongside the progression of ferroptosis, they can serve as noninvasive ferroptosis markers. Hence, detecting ferroptosis-associated VOLs in exhaled breath may enable the noninvasive evaluation of disease progression.

The profile of VOLs produced during ferroptosis and in damaged tissues remains poorly defined, primarily due to the lack of established methodologies for elucidating the mechanisms responsible for small volatile molecule formation, which could be exploited to establish *in vitro* and *in vivo* measurement systems. Therefore, despite the considerable potential of VOLs as noninvasive biomarkers, there has been minimal effort to comprehensively analyze the chemical structures of VOLs produced by ferroptotic cells and damaged tissue.

To address this research gap, we performed oxidative volatolomics for a high-throughput analysis of VOLs released during ferroptosis. Three VOLs released from cells undergoing iron-dependent LPO were identified and confirmed as biomarkers for monitoring ferroptosis in liver disease. Notably, the levels of these markers were also higher in the breath of patients with metabolic dysfunction-associated steatotic liver disease (MASLD) than in healthy individuals, highlighting their potential as noninvasive markers for disease monitoring.

These molecules serve as noninvasive markers for evaluating iron-dependent LPO, a key hallmark of ferroptosis[3], at the human level, and are valuable for monitoring pathological conditions associated with ferroptosis.

98 **Materials and Methods**

99 *Materials*

100   Linoleic acid (FA18:2), arachidonic acid (FA20:4), eicosapentaenoic acid (FA20:5), and

101 docosahexaenoic acid (FA22:6) were purchased from Nacalai Tesque, Inc. (Kyoto, Japan). 2,2′-

102 Azobis (2-methylpropionamidine) dihydrochloride (AAPH) and *N*-acetyl cysteine (NAC) were

103 obtained from Wako Pure Chemical Industries (Osaka, Japan), and hemin was procured from

104 Tokyo Chemical Industry Co., Ltd. (Tokyo, Japan), and acetaminophen (APAP) was purchased

105 from Sigma-Aldrich (St. Louis, MO, USA). (1S,3R)-RSL3, FINO2, ML162, liproxstatin-1,

106 staurosporine, 15-HpETE, 12-HpETE, 9-HpODE, 13-HpODE, arachidonic acid-d5 (FA20:4-

107 d5), linoleic acid-d4 (FA18:2-d4), arachidonic acid-d8 (FA20:4-d8), and docosahexaenoic

108 acid-d5 (FA22:6-d5)were obtained from Cayman Chemical (Ann Arbor, MI, USA).

109 Acetonitrile (LCMS grade, $\geq$ 99.9%), isopropanol (LCMS grade, $\geq$ 99.9%), methanol (LCMS

110 grade, $\geq$ 99.9%), water (LCMS grade, $\geq$ 99.9%), and ammonium formate (Wako 1st grade)

111 were purchased from Wako Pure Chemical Industries, Ltd.

112

113 *Cell culture*

114   HepG2, Calu-1, and HEK293 cells were purchased from American Type Culture Collection

115 (Manassas, VA, USA). HepG2 and HEK293 cells were cultured in high-glucose Dulbecco's

116 Modified Eagle Medium (DMEM), whereas Calu-1 cells were cultured in McCoy's 5A medium.

117 All growth media were supplemented with 10% fetal bovine serum (FBS) and 100 U/mL of

118 penicillin and streptomycin. All cell lines were maintained at 37 °C in a humidified incubator

119 with a 5% CO_2 atmosphere, passaged for < 6 months, and were not further tested or

120 authenticated by the authors.

121

122 *Cytotoxicity assay*

123   Cells (1×10^4 cells/well) were plated in 96-well flat-bottom plates and cultured for 24 h

124 before exposure to various concentrations of compounds at 37 °C. The amount of LDH in the

125 supernatant was measured using a cytotoxicity LDH assay kit-WST (Dojindo Molecular

126 Technologies, Inc., Kumamoto, Japan) following the manufacturer's instructions. For the MTT

127 assay, 15 µL of 5 mg/mL MTT solution (Nacalai Tesque, 23547-21) was added to each well

128 and incubated at 37 °C for 1 h. The supernatant was removed, and 100 µL of dimethyl sulfoxide

129 (DMSO) was added to each well. Absorbance was measured at 490 nm (LDH) and 570 nm

130 (MTT assay) using a 96-well plate reader (Infinite F50; Tecan Trading AG, Switzerland).

131

132 *Isotope labeling of VOLs generated during ferroptosis*

133   Screening-1 ($^{18}O_2$/$H_2^{18}O$ labeling): HepG2 cells (1×10^6 cells) were seeded into culture

134 flasks and incubated for 24 h. Subsequently, the culture medium was replaced with DMEM

135 prepared using $H_2^{18}O$ and supplemented with 10% FBS. The flask was then hermetically sealed,

136 and $^{18}O_2$-enriched air (composition: N_2 74%, $^{18}O_2$ 21%, and CO_2 5%) generated using the
137 CUBE GM-X3 (FCON CO., LTD., Kochi, Japan) was introduced into the flask at a flow rate
138 of 50 mL/min for 20 min via a syringe, thereby replacing the internal atmosphere with $^{18}O_2$ air.
139 Subsequently, a ferroptosis inducer (with or without a ferroptosis inhibitor) was added, and
140 after a defined incubation period, the headspace gas above the cell culture was collected.
141 Sampling was conducted using the SP209-1000 Dual device (GL Sciences Inc., Tokyo, Japan)
142 at a flow rate of 50 mL/min for 30 min, and the volatiles were trapped in inert-coated
143 biomonitoring tubes (Markes International, Llantrisant, UK). The collected samples were
144 subsequently analyzed using thermal desorption–gas chromatography/high-resolution mass
145 spectrometry (TD-GC/HRMS).
146 Screening-2 (d-labeled PUFA labeling); HepG2 cells (5×10^5 cells) were seeded into culture
147 flasks and incubated for 24 h. Subsequently, d-labeled PUFAs (FA20:4-d5, FA18:2-d4, FA20:4-
148 d8, and FA22:6-d5) were added at a final concentration of 20 µM, followed by additional 24-
149 hour incubation. After this treatment, the culture medium containing the D-labeled PUFAs was
150 replaced with standard DMEM, and ferroptosis induction, gas collection, and analysis were
151 performed in the same manner as described in Screening-1.
152

VOL analysis by TD-GC/HRMS

154 The TD injection system was a TD100-xr (Markes International, California, USA) equipped
155 with Tenax TA focusing trap. The sample path was set to 230 °C, and the flow rate was adjusted
156 to 10:1 in the split mode. The cold focusing trap was set to −30 °C, and trap desorption was
157 performed at 280 °C. A TRACE 1610 Gas Chromatograph (Thermo Fisher Scientific, San Jose,
158 CA, USA) equipped with a TG-5SILMS column (60 m, 0.25 mm ID, 0.25 µm FT, Thermo
159 Fisher Scientific) was used. The carrier flow was set to 1 mL/min using helium. The ramped
160 oven program was set as follows: 30 °C (5 min hold), 5 °C/min to 150 °C, 10 °C/min to 280 °C
161 (20 min hold).
162 An Orbitrap Exploris GC 240 mass spectrometer (Thermo Fisher Scientific) was used. Data
163 acquisition was performed in full scan mode (m/z 35–450) with a resolving power of 60,000
164 (FWHM) at m/z 200. HRMS data were analyzed using Compound Discoverer software
165 (Thermo Fisher Scientific).
166 Unlabeled and labeled raw data were employed to distinguish between isotope-unlabeled
167 and isotope-labeled mass spectra, and the retention times at which the labeled spectra were
168 detected were extracted. Next, unlabeled raw data were employed to identify the components
169 corresponding to the extracted retention times from the component table obtained through
170 deconvolution, NIST spectrum search, high-resolution filtering, and the retention index.
171 TD-GC/HRMS was employed for VOL analysis, as shown in **Figures 1b, c,** and **e**.
172

Hydroperoxy fatty acid (HpFA)-derived VOLs, radicals and lipid fragments analysis

174   HpFA/hemin-induced oxidation was performed by incubating a solution containing 10 µM
175   HpFAs, 10 µM hemin, and 10 µM NBD-Pen in PBS (pH 7.4) at 37 °C for 15 min. For the VOL
176   analysis, the reaction was conducted in an SPME vial. After the designated reaction period, the
177   volatile compounds were analyzed using SPME-GC/MS. For radical and lipid fragment
178   analyses, the mixture was extracted using the modified Bligh and Dyer method. The extracted
179   solution was dried under a nitrogen gas stream, and the residue was dissolved in methanol (200
180   µL) and stored at −80 °C before injection into the LC/HRMS system.
181

182   *VOL analysis by SPME-GC/MS*
183   The GC/MS system used was an Agilent 5977 B GC/MSD. An SPME autosampler was used
184   as a PAL-RTC multifunctional sampler for gas chromatography. Chromatographic separation
185   was achieved using an HP-5ms column (60 m × 0.25 mm, I.D. × 0.25 µm), with helium as the
186   carrier gas at a flow rate of 1.5 mL/min. The injector temperature was maintained at 250 °C.
187   The column oven temperature program began at 35 °C, was held for 4 min, increased to 150 °C
188   at a rate of 5 °C/min, increased to 280 °C at a rate of 10 °C/min, and was finally held for 10
189   min. Electron ionization (EI) method employing scan (test-tube experiments) and SIM (cell
190   culture experiments) analyses was used. GC/MS analysis was controlled using MassHunter
191   software, with autosampler operations managed using PAL Sample Control software.
192   VOL extraction was performed using a SPME Arrow fiber (SARR15-DVB/CWR120/20-P
193   1). The vial was agitated at 40 °C and 500 rpm for 900 s, followed by the extraction of VOLs
194   from the headspace for 600 s under the same conditions. Subsequently, the SPME arrow fiber
195   was inserted into the GC/MS injector and desorbed at 250 °C for 60 s.
196   The identified VOLs were analyzed using MassHunter Workstation Qualitative Analysis
197   software. Compounds were identified using the NIST MS Search ver. 2.4 and the NIST Mass
198   Spectral Library. Mass spectra from the experiments were compared with those of the selected
199   candidate compounds and were further verified by comparing the GC/MS results with those of
200   the standard samples of these candidates. Peaks that were not identified in the library search
201   were designated VOL-n (number). Additionally, peak area values were computed from the
202   mass chromatograms of the most intense fragment peaks for each compound. SPME-GC/MS
203   was employed for VOL analysis, as shown in **Figures 1i, 2b-d, f-g, 3d, f, g, i-k, 4e, l-n, and**
204   **Extended Data Figures 4** and **5**.
205

206   *Oxidation of standard PUFAs using AAPH/hemin*
207   In an SPME vial, PUFAs (500 µM), AAPH (50 mM), and hemin (10 µM) were mixed in
208   phosphate-buffered saline (PBS; pH 7.4, containing 0.1% ethanol). After sealing the vial, the
209   reaction mixture was incubated at 37 °C for 2 h. The samples were analyzed using SPME-
210   GC/MS analysis.
211

Collection and analysis of cellular VOLs using SPME-GC/MS

HepG2 cells were seeded (5×10^5 cells/well) in 6-well culture plates and allowed to adhere for 24 h before treatment. Ferroptosis was induced using RSL3 (3.0 µM for HepG2, 0.5 µM for Calu-1, and HEK293), FINO2 (25 µM), and ML162 (3.0 µM). Apoptosis was induced by STS (10 µM) and etoposide (100 µM). Cells were also treated with the ferroptosis inhibitors, Lip-1 (1.0 µM) and DFO (100 µM). Subsequently, the cells and culture medium were transferred to an SPME vial and analyzed using SPME-GC/MS.

Lipid measurements using LC/HRMS

To extract lipids, 1000 µL of a 1:1 (v/v) 1-butanol/methanol solution containing 100 µM butylated hydroxytoluene (BHT) and 100 µM ethylenediaminetetraacetic acid (EDTA) was added to the cell pellets. The mixture was vortexed for 10 s, sonicated for 15 min in an ice-cooled sonic bath, and centrifuged at 16 000 × *g* for 10 min at 20 °C. The supernatant was transferred into a 0.2 mL glass insert with a Teflon-lined cap for liquid chromatography electrospray ionization mass spectrometry (LC-ESI-MS) analysis.

For lipid analysis, a Q-Exactive Focus mass spectrometer (Thermo Fisher Scientific) was connected to an Ultimate 3000 HPLC system (Thermo Fisher Scientific). Samples were separated using a Thermo Scientific Accucore C18 column (2.1 × 150 mm, 2.6 µm) with mobile phase A comprising 10 mM ammonium formate in 50% acetonitrile with 0.1% formic acid and mobile phase B comprising 2 mM ammonium formate in a mixture of acetonitrile, isopropyl alcohol, and water (10:88:2, v/v/v) with 0.02% formic acid. The step gradient was as follows: 65:35 (0 min), 40:60 (0–4 min), 15:85 (4–12 min), 0:100 (12–21 min), 0:100 (21–24 min), 65:35 (24–24.1 min), and 100:0 (24.1–28 min), at a flow rate of 0.4 mL/min, and a column temperature of 35 °C.

The Q-Exactive Focus mass spectrometer was operated in the ESI positive and negative ion modes. The ion source parameters comprised a spray voltage of 3 kV, transfer tube temperature of 285 °C, S-Lens level of 45, heater temperature of 370 °C, sheath gas flow rate of 60, and auxiliary gas flow rate of 20. LC/HRMS in the positive and negative ion modes was employed for the relative quantification of oxidized lipids based on previously published methods[38,39]. The experimental conditions for full-scan MS were as follows: resolving power = 70,000, automatic gain control target = 1×10^6, trap fill time = 100 ms, and scan range = m/z 250–1500. The precursor and product ions of the d-labeled phospholipids analyzed by PRM are listed in **Supplementary Table 3**. The experimental conditions for PRM were as follows: resolving power = 17,500, automatic gain control target = 1×10^6, and trap fill time = 100 ms. The acquired data were analyzed using Qual Browser (Thermo Fisher Scientific).

Apoptosis assay

249   HepG2 cells (5×10^5 cells/well) were seeded in a six-well plate and cultured for 24 h before

250   treatment. After treating the cells with 10 µM STS or 100 µM etoposide for 6 h, the cells were

251   washed with PBS. Accutase (Nacalai Tesque, Inc. Kyoto, Japan) was used to detach cells,

252   which were then washed with 1 mL HBSS. To detect apoptosis, the Annexin V-633 Apoptosis

253   Detection Kit (Nacalai Tesque, Kyoto, Japan) was used according to the manufacturer's

254   protocol. Cells were passed through a cell strainer and assayed using a flow cytometer (BD

255   FACSCanto™ II, BD Biosciences, Franklin Lakes, NJ, USA) with 488 nm/633 nm laser for

256   excitation. Fluorescence intensity was measured using a 660/20 band-pass filter for Annexin

257   V-APC and a 575/26 band-pass filter for propidium iodide. At least 10,000 events were

258   collected for each sample. Data were analyzed using FlowJo software (BD Biosciences).

259

260   ***Senescence-associated β-galactosidase staining***

261   HepG2 cells (1×10^5 cells/well) were seeded in 35-mm dishes and cultured for 7 days in

262   DMEM supplemented with 2.0 µM etoposide. Senescence-associated β-galactosidase (SA-β-

263   Gal) activity was detected using a Senescence β-Galactosidase Staining Kit (Cell Signaling

264   Technology) according to the manufacturer's instructions. Stained cells were observed using a

265   fluorescence microscope (BZ-8000; Keyence, Osaka, Japan).

266

267   ***Oil Red O staining***

268   HepG2 cells (1×10^5 cells/well) were seeded in 35-mm dishes and cultured for 2 days in

269   DMEM supplemented with a fatty acid–albumin complex (linoleic acid and oleic acid in a 2:1

270   molar ratio, Sigma-Aldrich). Lipid accumulation was quantified using an Oil Red O solution

271   (0.5% v/v). Cell images were captured using a BZ-8000 fluorescence microscope (BZ-8000).

272

273   ***Animal model***

274   Male C57BL/6J mice were acquired from CLEA Japan, Inc. (Tokyo, Japan). All mice were

275   housed under sterile conditions at a temperature of 24 ± 1 °C, humidity of $60 \pm 10\%$, and a 12-

276   h light-dark cycle. Drinking water and food were provided *ad libitum* during the study period.

277   The animal study was approved by the Animal Studies Committee of Kyoto University.

278   APAP-acute liver failure (ALF) model mice: A single dose of APAP (300 mg/kg body

279   weight) was intraperitoneally administered to overnight-fasted mice aged eight weeks. Six

280   mice were used in each of the experimental groups. To suppress APAP-induced LPO, NAC

281   (300 mg/kg body weight) in saline was intraperitoneally injected 1 h after APAP administration.

282   MASH model mice: Male mice aged 6 weeks were randomly divided into three groups fed

283   a standard diet (SD), a choline-deficient, L-amino acid-defined, high-fat diet with 60 kcal% fat

284   and 0.1% methionine without added choline (CDAHFD; Research Diets A06071302, Research

285   Diets, Inc. New Brunswick, NJ, USA) or CDHFD with vitamin E (1.0 g/kg diet) for six weeks

286   to establish a MASH mouse model. Six mice were used in each experimental group.

287

VOL analysis using TD-GC/MS

289   A Shimadzu GCMS-QP2020NX thermal desorption system (Shimadzu TD-30R, Shimadzu
290 Corporation, Kyoto, Japan) was used for the GC/MS analysis. The separation column
291 employed was an HP-5ms (60 m × 0.25 mm, I.D. × 0.25 μm). Helium was used as the carrier
292 gas at a flow rate of 1.5 mL/min. The injector temperature was maintained at 250 °C. The
293 temperature program for the column oven started at 35 °C, was held for 4 min, increased to
294 150 °C at 5 °C/min, increased to 280 °C at 10 °C/min, and was held for 10 min. EI was
295 employed for ionization, and selected ion monitoring (SIM) analysis was performed. Volatile
296 oxidized lipids were analyzed using the Shimadzu LabSolutions software (Shimadzu
297 Corporation, Kyoto, Japan). TD-GC/MS was employed for VOL analysis, as shown in **Figures**
298 **4g, o, 5**.

299

Collection of VOLs in mouse-derived gas

301   Mice were placed in a sealed chamber into which artificial air (N_2:78%, O_2:22%) generated
302 using CUBE GM-X3 was introduced at a flow rate of 50 mL/min. An inert-coated
303 biomonitoring tube was connected to the outlet of the gas line, and gas sampling was performed
304 for 60 min. The collected gas samples were analyzed using TD-GC/MS.

305

Extraction of VOLs and oxidized lipids from mouse tissues

307   VOLs were extracted from mouse tissues using an extraction solution (methanol containing
308 100 μM dibutyl hydroxytoluene [BHT] and 100 μM EDTA). Briefly, 1 mL of extraction
309 solution was added to 50 mg of frozen tissue and homogenized using a Bertin Minilys
310 homogenizer (Bertin Technologies SAS, Montigny-le-Bretonneux, France). The homogenate
311 was sonicated for 5 min on ice, allowed to stand for 5 min, and centrifuged (6000 × g for 10
312 min at 4 °C). The supernatant (~700 μL) was collected, and the extracted samples were stored
313 at −80 °C until measurement.

314

Histopathology

316   Mouse liver was freshly sampled, snap-frozen in liquid nitrogen, embedded in SCEM
317 compound (Leica Microsystems, Tokyo, Japan), and stored at −80 °C until analysis. After
318 freezing, 6-μm-thick cryostat sections were obtained at −12 °C. Frozen sections were stained
319 with hematoxylin and eosin (H&E). Images were obtained using a slide scanner (Fino CLARO;
320 DRC, Tokyo, Japan).

321

Human Exhaled Breath Sample Collection

323   The trial protocol was approved by the ethics committees of Keio University Hospital
324 (approval no. 20221057) and Kitano Hospital (approval no. P221100201). All procedures were

325 conducted according to the Declaration of Helsinki and the Ethical Guidelines for Medical and

326 Health Research Involving Human Subjects (Japanese Ministry of Health, Labour and Welfare).

327 Informed consent was obtained from all the participants.

328   Forty-eight healthy participants and 67 patients diagnosed with MASLD or liver cirrhosis

329 (LC) were enrolled between February 2023 and February 2025 at Keio University Hospital and

330 Kitano Hospital (**Supplementary Table 2**). Healthy individuals had no comorbidities, such as

331 hypertension, diabetes, or active lung disease. Patients who had difficulty using a mask for 15

332 min or required oxygen therapy were excluded from the study. All enrolled patients fulfilled

333 the diagnostic criteria for nonalcoholic fatty liver disease (NAFLD) [40] and MASLD[41], with

334 MASH diagnosed according to the Delphi consensus statement[41], and LC diagnosed according

335 to the most recent Japanese Guideline[42]. The MASLD group included 56 patients (including

336 those with MASH but without cirrhosis), and the LC group included 11 patients. The median

337 ages were 58.0 years (interquartile range [IQR] 32–87) and 69.6 years (IQR 58–81),

338 respectively, with female proportions of 60.7% and 54.5%. All LC cases were categorized as

339 Child–Pugh class A. The albumin-bilirubin (ALBI) scores were −2.98 (IQR −3.67 to −2.33)

340 for MASLD and −2.43 (IQR −3.33 to −1.65) for LC; the FIB-4 indices were 1.73 (IQR 0.30–

341 6.61) for MASLD and 4.78 (IQR 1.77–10.86) for LC; and M2BPGi levels were 0.99 (IQR

342 0.21–3.3) and 2.85 (IQR 0.54–5.73), respectively.

343   Breath samples were collected at each institution using breath biopsy collection stations and

344 stored in thermal desorption tubes (**Fig. 5a**). The tubes were transported to Kyoto University

345 for TD-GC/MS analysis (**Fig. 5b**). This multicenter approach enabled the consistent collection

346 and measurement of exhaled breath from geographically distinct sites, ensuring broad

347 applicability.

348

349 ***Breath collection using the breath biopsy collection station***

350   Breath samples were obtained using inert-coated biomonitoring tubes connected to a

351 ReCIVA Breath Sampler (Owlstone Medical, Cambridge, UK). Prior to sample collection, the

352 tubes were prepared in a TD-30R (Shimadzu) under a 20-psi nitrogen gas flow at 280 °C for

353 30 min. Before sampling, the system was calibrated and adjusted according to each individual's

354 breathing pattern. Each tube collected approximately 1.0 L of breath at a flow rate of 200

355 mL/min. A CASPER Portable Air Supply (Owlstone Medical, Cambridge, UK) was used to

356 minimize ambient contamination. The tubes were stored at −20 °C, and VOL analysis was

357 performed within one week of collection.

358

359 ***Liver biopsy***

360   Among the patients enrolled in this study, those scheduled to undergo liver resection for liver

361 tumors underwent sharp dissection biopsy of non-tumor background liver tissue from the

362 planned resection area during surgery. Liver biopsy specimens were immediately snap-frozen
363 in liquid nitrogen and stored at −80 °C.
364

365 *Statistical analyses*
366   Statistical analyses were conducted using GraphPad Prism (version 9.2.0, GraphPad
367 Software, Inc., San Diego, CA, USA), and data were analyzed using unpaired Student's *t*-test,
368 one-way analysis of variance (ANOVA) with Tukey's multiple comparisons test, two-sided
369 Pearson's correlation, or two-way ANOVA with Tukey's multiple comparisons tests. The
370 results are expressed as mean ± standard deviation (SD). Statistical significance was set at $p <$
371 0.05. LASSO regression analysis was performed using the Glmnet R package (v4.1–8) and R
372 v4.3.2. Multivariate ROC curves and AUCs were derived using the "glm" function in R
373 (version 4.3.2).
374

Results

Ferroptosis-derived volatile markers: 1-octen-3-ol and 2-pentylfuran

LPO during ferroptosis is characterized by (1) the formation of lipid hydroperoxides following the addition of molecular oxygen (O_2) to polyunsaturated fatty acid-containing phospholipids (PUFA-PLs) and (2) the subsequent generation of highly oxidized lipids—including truncated short-chain molecules—via Fenton reactions between lipid hydroperoxides and labile iron. We hypothesized that comprehensive profiling of VOLs released from peroxidized PUFAs would facilitate the identification of ferroptosis-associated VOL markers.

To test this hypothesis and minimize confounding factors from non-cellular sources (e.g., culture medium or environmental contaminants), two complementary isotope-tracing approaches were employed using $^{18}O_2$ and deuterium-labeled arachidonic acid (FA20:4-d_5). Briefly, HepG2 cells were treated separately with $^{18}O_2$ (Screening-1) or FA20:4-d_5 (Screening-2) before ferroptosis induction. Oxidative volatolomics was then used to profile the isotope-labeled VOLs released into the gas phase (**Fig. 1a**).

In Screening-1, ferroptosis was induced in HepG2 cells cultured in an $^{18}O_2$ air/H_2^{18}O-medium environment, and ^{18}O-labeled volatile organic compounds (VOCs) were identified via thermal desorption-gas chromatography/high-resolution mass spectrometry (TD-GC/HRMS), yielding 21 labeled candidates (**Fig. 1b, Extended Data Fig. 1a**). In Screening-2, HepG2 cells were incubated overnight with FA20:4-d_5—labeled at the ω-terminal structure. Subsequently, ferroptosis was induced using RAS-selective lethal small molecule 3 (RSL3). Five d_5-labeled VOCs were detected (**Fig. 1c, Extended Data Fig. 1b, Extended Data Fig. 2**). Isotope-labeled VOLs were detected exclusively under conditions in which both isotope labeling and RSL3 treatment were applied. Two VOCs, 1-octen-3-ol and 2-pentylfuran, were identified using both screening methods (**Fig. 1d**). The production of these markers increased in response to ferroptosis inducers (RSL3 and imidazole ketone erastin [IKE]) and decreased with liproxstatin-1 (Lip-1) co-treatment (**Fig. 1e**).

The chemical pathways responsible for the formation of 1-octen-3-ol and 2-pentylfuran from FA20:4 were also investigated. While 2-pentylfuran is presumed to form via intramolecular cyclization of 4-hydroxy-2-nonenal (a product of iron-dependent LPO of ω-6 PUFAs)[8], the precise mechanism of 1-octen-3-ol generation has not been experimentally established (**Fig. 1f, g**). Accordingly, *in vitro* reactions were performed using PUFA peroxides and hemin. Among the tested substrates (15-HpETE, 12-HpETE, 9-HpODE, and 13-HpODE), only 12-HpETE, which bears a ω-9 peroxide structure—yielded 1-octen-3-ol (**Fig. 1h, i**). This was accompanied by the accumulation of key intermediates, including fragmented fatty acids ($C_{12}H_{16}O_3$) and octenyl radicals (•C_8H_{15}) (**Fig. 1j**).

These findings reveal a previously uncharacterized mechanism for 1-octen-3-ol formation involving sequential Fenton reactions and β-scission from FA20:4-derived 12-HpETE. This

412 establishes a direct mechanistic link between ω-9 lipid peroxidation and the generation of 1-
413 octen-3-ol, a ferroptosis-associated volatile marker.

414

Ferroptotic VOLs generated via iron-driven oxidative cleavage of polyunsaturated fatty acids

416 To further elucidate the chemical sources of 1-octen-3-ol and 2-pentylfuran beyond
417 arachidonic acid, a series of *in vitro* oxidation experiments were performed on several PUFAs
418 under iron-dependent conditions.

419 By systematically analyzing VOLs produced in these reactions with solid-phase
420 microextraction (SPME)-GC/MS (**Fig. 2a**), a reference library was constructed that mapped
421 each PUFA to its characteristic iron-dependent oxidation products. This defines the broader
422 chemical landscape of iron-dependent VOL formation (**Fig. 2b**). This approach revealed that
423 1-octen-3-ol and 2-pentylfuran were consistently produced via the iron-driven cleavage of ω-
424 6 PUFAs (**Fig. 2c**). Meanwhile, ω-3 PUFAs generated 2-ethylfuran via the same iron-
425 dependent oxidation pathway (**Fig. 2d; Extended Data Fig. 3**).

426 These findings were further validated in ferroptosis-induced cells using deuterium-labeled
427 PUFAs (**Extended Data Fig. 4**). After incorporating each labeled PUFA into the cell
428 membranes, ferroptosis was induced, and SPME-GC/MS was performed to detect the resulting
429 d-labeled VOLs (**Fig. 2e**). The quantitative GC/MS ions for these labeled VOLs were defined
430 based on the *in vitro* oxidation of the corresponding d-PUFAs (**Extended Data Fig. 5**). The
431 incorporation of FA18:2-d$_4$ (another ω-6 PUFA) significantly increased the levels of d-labeled
432 1-octen-3-ol and 2-pentylfuran levels (**Fig. 2f**). Similarly, the incorporation of DHA-d$_5$ (an ω-
433 3 PUFA) led to the production of d-labeled 2-ethylfuran (**Fig. 2g**).

434 Overall, 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran were identified as key ferroptosis-
435 associated VOL markers derived from distinct PUFA classes. This comprehensive library-
436 based strategy established a clear link between specific PUFAs, iron-dependent oxidative
437 pathways, and the generation of signature VOLs during ferroptosis.

438

Gas-phase accumulation of specific VOLs reflects ferroptosis progression

440 Next, the ability of the three identified VOLs—1-octen-3-ol, 2-pentylfuran, and 2-
441 ethylfuran—to accumulate in the gas phase alongside ferroptosis progression was evaluated.
442 Cells treated with the ferroptosis inducer RSL3 were analyzed at multiple times to measure
443 lactate dehydrogenase (LDH) release, intracellular oxidized lipid levels, and VOLs.
444 Considering that these compounds arise via iron-dependent LPO, the cells were co-treated with
445 Lip-1 or the iron chelator deferoxamine (DFO). LDH release—a classic marker of cell death—
446 increased significantly 4 h after RSL3 treatment (**Fig. 3a**). Highly oxidized phospholipids (e.g.,
447 PE38:4; O3, **Extended Data Fig. 6**) appeared as early as 2 h (**Fig. 3b–c**); however, their
448 formation was blocked by Lip-1 or DFO. Notably, the three VOLs appeared in the gas phase
449 within 2 h of RSL3 treatment (**Fig. 3d**), coinciding with the onset of iron-dependent LPO

450 activity. This effect was suppressed by Lip-1 and DFO (**Fig. 3f**). Thus, these VOLs reliably
451 reflect the progression of ferroptotic cell death as gas-phase markers.

452   The three VOLs were elevated by various ferroptosis inducers (FINO2 and ML162) in
453 multiple cell lines (Calu-1 and HEK293; **Fig. 3f, 3g; Extended Data Fig. 7a, b**). In contrast,
454 apoptosis triggered by staurosporine or etoposide (**Fig. 3h**) failed to increase the abundance of
455 these proteins (**Fig. 3i**). Furthermore, conditions such as etoposide-induced senescence or
456 oleic/linoleic acid–induced lipid accumulation—where mild oxidative stress and LPO can
457 occur without cell death [9,10]—did not elevate the VOLs' levels (**Fig. 3j–k, Extended Data Fig.
458 8**).

459   These results demonstrate that 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran accumulate in
460 the gas phase only when iron-dependent LPO drives ferroptosis and remain unchanged under
461 other types of cell death or non-lethal oxidative stress. Hence, these VOLs are potentially robust
462 gas-phase markers for real-time monitoring of ferroptosis.

463

464 ***Hepatic ferroptosis induces 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran production and***
465 ***release as specific volatile markers***

466   To validate whether the ferroptotic VOL signature previously established in *in vitro* cell
467 experiments is also released as gas from living organisms, we performed *in vivo* animal
468 experiments using a mouse model of liver disease (**Fig. 4**). The liver is prone to ferroptosis
469 owing to its high iron content, and ferroptosis is implicated in the progression of drug-induced
470 acute hepatitis[11]. Therefore, mouse models of liver disease were employed to determine
471 whether the ferroptotic VOL signature discovered *in vitro* translates into ferroptosis-related
472 VOLs produced in the liver *in vivo* and detected in the emitted animal-derived gas.

473   First, an acetaminophen (APAP)-induced acute liver failure (ALF) model was used. High-
474 dose APAP depletes glutathione (GSH), leading to extensive hepatocellular ferroptosis[12]. Male
475 C57BL/6j mice were fasted and treated with APAP, and gas sampling and biochemical analyses
476 were performed 4 h later (**Fig. 4a**). The effect of N-acetylcysteine (NAC)—a GSH precursor—
477 was also evaluated. APAP significantly elevated plasma alanine aminotransferase (ALT)
478 activity and increased oxidized lipids in the liver tissue, indicating liver injury and ferroptosis
479 (**Fig. 4b–d**). However, NAC administration attenuated these effects. Notably, 1-octen-3-ol, 2-
480 pentylfuran, and 2-ethylfuran levels increased in APAP-treated liver tissue and decreased upon
481 NAC administration (**Fig. 4e**).

482   Subsequently, the mice were housed in sealed cages perfused with a VOC-free gas mixture
483 (N_2:78%, O_2:22%, 50 mL/min/cage), and the effluent gas was collected for 1 h using bio-
484 monitoring inert-coated tubes for TD-GC/MS analysis (**Fig. 4f**). Elevated levels of 1-octen-3-
485 ol, 2-pentylfuran, and 2-ethylfuran were detected in the gas released from APAP-treated mice,
486 which decreased after NAC treatment (**Fig. 4g**). Thus, ferroptosis in the APAP-ALF model can
487 be monitored noninvasively via VOL markers in the gas derived from mice.

Ferroptosis also contributes to metabolic dysfunction-associated steatohepatitis (MASH)[13]. Hence, the production of the three ferroptosis VOLs was assessed in a MASH mouse model that fed a choline-deficient, L-amino acid-defined, high-fat diet (CDAHFD) for 6 weeks (**Fig. 4h**). Vitamin E (VE), a known ferroptosis inhibitor, was administered to a subset of mice. CDAHFD-fed mice showed marked hepatic fat accumulation, elevated plasma ALT levels, and increased oxidized phospholipids, indicative of progressive liver injury and ferroptosis. VE supplementation ameliorated these effects. Additionally, the ferroptosis VOL signature was significantly higher in the liver tissue of the CDAHFD group and lower in the VE-supplemented group (**Fig. 4i–k, and Extended Data Fig. 9**). The levels of these markers remained low in non-liver tissues, underscoring their specific origin in ferroptotic hepatocytes (Fig. 4l–n). Furthermore, gas analysis confirmed elevated levels of 1-octen-3-ol and 2-pentylfuran in CDAHFD-fed mice and their reduction following VE intervention. This suggests that ferroptosis in the liver can be detected based on the levels of these VOL markers (**Fig. 4o**).

Collectively, these findings confirm that the ferroptosis-associated VOLs identified *in vitro*—1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran—are also emitted *in vivo* via hepatic ferroptosis. Thus, these VOL markers enable the noninvasive detection of ferroptosis-driven liver injury in acute and chronic disease models.

Noninvasive detection of ferroptosis in patients with MASLD using breath analysis

Finally, we investigated whether the identified VOLs were elevated in the exhaled breath of patients with MASLD, including metabolic dysfunction-associated steatohepatitis (MASH). Breath VOL analysis was conducted on healthy individuals ($n = 48$), patients with MASLD ($n = 56$, comprising MASH without cirrhosis [LC]), and MASLD-related LC patients ($n = 11$) following the collection of breath samples in thermal desorption tubes (**Fig. 5a**) and TD-GC/MS analysis (**Fig. 5b**). The ferroptosis VOL markers showed marked variations among the disease groups (**Fig. 5c**). Patients with MASLD had significantly higher ferroptosis-related VOLs levels (**Fig. 5d**). Specifically, 1-octen-3-ol levels were elevated in patients with MASLD compared to those in healthy individuals ($p = 0.0143$); however, this increase was suppressed in patients with LC. Meanwhile, 2-pentylfuran levels were elevated in MASLD and LC ($p = 0.0022$ for healthy vs. MASLD, $p < 0.0001$ for healthy vs. LC, and $p < 0.0001$ for MASLD vs. LC). In addition, 2-ethylfuran was significantly elevated in the LC group ($p < 0.0001$ for healthy vs. LC and $p = 0.0008$ for MASLD vs. LC). Combining these markers enabled discrimination between healthy individuals and those with MASLD or LC with area under the curve (AUC) values of 0.824 and 0.931, respectively (**Fig. 5e**). Furthermore, integrating ferroptosis VOL markers with standard clinical markers (using least absolute shrinkage and selection operator [LASSO] analysis) distinguished MASLD from LC with high confidence (AUC > 0.9; **Fig. 5f and Extended Data Fig. 10**).

526   Correlations between these VOLs and pathophysiological indices were also evaluated,
527   revealing that 2-pentylfuran showed a moderate correlation (r > 0.4) with fibrosis markers
528   (FIB-4, MAC-2 binding protein glycosylation isomer [M2BPGi]) and the albumin-bilirubin
529   (ALBI) score (r > 0.6) (**Fig. 5g**). Finally, to assess whether human breath VOLs reflect hepatic
530   oxidized lipid levels, oxidized phosphatidylcholine (oxPC), and oxidized triglyceride (oxTG),
531   which were measured in liver biopsy samples, revealed a strong correlation with breath 2-
532   pentylfuran levels (**Fig. 5h**).

533   Collectively, these findings indicate that noninvasive breath analysis of ferroptosis VOLs
534   can identify advanced MASLD stages. Hence, these ferroptosis-derived VOL markers could
535   serve as real-time indicators of disease progression and offer a noninvasive screening strategy
536   for patients with MASLD and LC.

537

538

Discussion

In this study, we identified small fragments of lipid peroxides released as gas during ferroptosis and defined their molecular characteristics. PUFA-derived VOLs, namely 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran, were identified as markers of *in vivo* ferroptosis induced via iron-dependent LPO, which may have clinical relevance (**Fig. 6**).

VOCs are ubiquitous in ambient air, which complicates the accurate detection and identification of cellular-derived VOCs at trace levels owing to background contamination. To address this, we employed stable isotope-labeled molecules, enabling the differentiation of cellular-derived volatile metabolites from environmental contaminants and achieving highly precise analytical results. However, VOCs other than the identified VOL markers, such as ketones and alcohols, may reflect various intracellular metabolic pathways or cellular metabolic conditions. Thus, further elucidation of the biosynthetic pathways and underlying mechanisms responsible for generating these VOCs will significantly advance our understanding of cellular metabolism and improve the specificity of the detection methods.

Standard lipids were oxidized *in vitro*, and 48 VOLs with different chemical structures were detected (**Fig. 2** and **Supplementary Table 1**). The types and amounts of identified VOLs differed significantly with hemin addition (**Fig. 2b**). Nonenal and hexenal-like alkenals were more abundant when only AAPH was added (**Extended Data Fig. 11**), presumably as the hock cleavage products of lipid peroxides[14]. For example, hock cleavage initiated by the protonation of 11-HpETE produces 3-nonenal and its isomerized form, 2-nonenal (**Extended Data Fig. 12**). In contrast, samples treated with both AAPH and hemin showed increased production of vinyl alcohol and 2-alkylfuran (**Fig. 2c, d**). 2-alkylfuran is a cyclized molecule derived from 4-hydroxyalkenal. The formation of 4-hydroxyalkenal is facilitated by the Fenton reaction of 4-hydroperoxyalkenal and PUFA hydroperoxides (**Fig. 1g**). The addition of hemin, which facilitates this reaction, may have increased the amount of 4-hydroxyalkenal produced.

The VOLs 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran were produced by the iron-dependent oxidation of PUFA, and confirmed as markers of ferroptosis induction *in vitro*, *in vivo*, and human breath. Specifically, 1-octen-3-ol is biosynthesized from FA18:2 by certain fungi and plants using 10-lipoxygenase and hydroperoxide lyase[15], respectively. However, whether homologous enzymes or enzymatic activities exist in mammals and whether a nonenzymatic formation reaction occurs remains unknown. Herein, we identified a reaction scheme in which iron plays a significant role in the nonenzymatic formation of 1-octen-3-ol. Furthermore, these VOL markers were derived from membrane lipids in HepG2 cells during ferroptotic progression (**Fig. 2f, g**). Notably, 1-octen-3-ol was produced in greater amounts from FA20:4 than from 2-pentylfuran (**Fig. 2f**). The generation of 1-octen-3-ol preferentially requires a ω-9 hydroperoxide structure, such as 12-HpETE (**Fig. 1f**). In contrast, FA18:2 is less prone to forming 10-HpODE with a hydroperoxide moiety at the C9 position via radical-mediated pathways[16]. In contrast, the formation of 2-pentylfuran is dependent on the presence

577 of a ω-10 hydroperoxide structure, which can be generated through radical-mediated oxidation
578 of both FA20:4 and FA18:2, corresponding to 11-HpETE and 9-HpODE, respectively. This
579 mechanistic difference likely accounts for the distinct production profiles of 1-octen-3-ol and
580 2-pentylfuran from each PUFA substrate.

581   During ferroptosis, labile iron stored in lysosomes is released into the cytosol, resulting in
582 lethal iron-dependent LPO[17]. In addition, the liver is a major organ with iron stores[18], and an
583 increase in free iron levels in the liver tissue may occur during APAP-ALF and MASH
584 induction[19,20]. As these VOLs are selectively produced in ferroptotic cells, particularly
585 hepatocytes, they may be effective markers for differentiating between healthy and diseased
586 states of the liver.

587   The levels of ferroptosis VOL markers in the breath of patients with MASLD or LC were
588 higher than those in healthy individuals. Previous fundamental research using mouse models
589 has indicated that hepatocyte ferroptosis triggers increased hepatic inflammation in the early
590 stages of MASH development[13]. This study confirmed that ferroptosis progresses during the
591 early stages of chronic hepatitis in human. In contrast, the levels of 2-pentylfuran and 2-
592 ethylfuran were higher in patients with LC, demonstrating their utility in distinguishing LC
593 from other conditions. This could be attributed to their high lipophilicity and accumulation in
594 liver tissue due to the reduced degradative capacity associated with impaired liver function in
595 patients with LC. The differences in the properties of these markers could be attributed to their
596 retention times in the body; for example, lipid accumulation in lipid droplets may increase
597 retention time. Furthermore, 2-pentylfuran is degraded by P450[21], and as P450 activity is
598 compromised in patients with LC[22], the accumulation of 2-pentylfuran and 2-ethylfuran
599 produced by ferroptosis in liver tissue may be more pronounced in these patients. Although
600 further detailed investigation of the pharmacokinetics of VOLs is necessary, these results
601 suggest that combining the three VOL markers could be useful for diagnosing MASLD/LC
602 stages and identifying patients (**Extended Data Fig. 13**).

603   MASLD, with an estimated global prevalence of 30%, is the most rapidly growing
604 contributor to liver-related morbidity and mortality[23,24]. Due to its prevalence and close
605 association with the global epidemics of obesity and metabolic syndrome, an easily obtained,
606 noninvasive, and liver-specific biomarker capable of identifying individuals with high-risk
607 MASLD or MASH is among the most challenging unmet clinical needs[25]. In this study, we
608 demonstrated that ferroptosis-related VOLs may be useful as liver-specific mechanistic
609 biomarkers for MASLD. Although hepatic histopathological improvements are necessary for
610 drug development for MASH, the invasive nature of percutaneous liver biopsies, along with
611 their high inter- and intra-investigator variability, are considerable obstacles[26]. Although further
612 investigations are warranted, our technology for analyzing ferroptosis-related VOLs in exhaled
613 breath may prove effective for mass screening, risk stratification, and assessing therapeutic
614 efficacy.

615   However, the bioactivities of these VOLs remain unknown and require further investigation.

616   Intracellular molecules released from ferroptotic cells induce ferroptosis in surrounding cells[27].

617   Low-molecular-weight oxidized lipids, including VOLs, are implicated in ferroptosis

618   propagation. Interestingly, 1-octen-3-one (the oxidized form of 1-octen-3-ol) was highly

619   cytotoxic (**Extended Data Fig. 14**). Similarly, α,β-unsaturated carbonyls, including 1-octen-3-

620   one, are highly electrophilic and cause modifications of nucleophilic amino acid residues,

621   leading to ferroptosis[7]. In addition, compounds containing a furan ring structure, such as 2-

622   pentylfuran and 2-ethylfuran, have also been reported to contribute to the induction of

623   ferroptosis[28]. In addition, their precursor, 4-hydroxynonenal (4-HNE), promotes ferroptosis by

624   downregulating the expression of GPx4[29,30]. Overall, the VOLs produced during ferroptosis

625   and diffused into the environment may contribute to ferroptosis in neighboring cells. Therefore,

626   the bioactivity of the VOLs identified in this study, which are selective for ferroptotic cells,

627   warrants further investigation.

628   A major limitation of using VOL markers in exhaled breath is the unclear origin of VOLs.

629   In this study, we confirmed that the three marker VOLs were specifically produced in the liver

630   of a mouse APAP-ALF/MASH model. However, the organ specificity of VOL sources in

631   human MASLD/LC requires further investigation. In addition, changes in VOL production

632   patterns in the liver due to comorbidities should be characterized. Lipid peroxidation is

633   enhanced in patients with obesity and diabetes[31,32], and alterations in the levels of certain

634   exhaled VOC have also been observed in these populations[33,34]. Additionally, lifestyle factors

635   such as smoking have been reported to influence the profile of exhaled VOCs[35]. Therefore,

636   changes in VOL production associated with lifestyle-related comorbidities such as obesity and

637   diabetes, as well as behavioral and extrahepatic factors such as smoking, alcohol consumption,

638   and pulmonary diseases, must be assessed in large-scale studies. In this study, we analyzed the

639   correlation between the identified VOL markers and smoking habits and the presence of

640   pulmonary diseases; however, no significant associations were observed with either factor

641   (**Extended Data Fig. 15**). In addition, further investigation is warranted to evaluate the *in vivo*

642   specificity of this marker for ferroptosis using models in which ferroptosis is the exclusive

643   mode of cell death, such as Gpx4-knockout mice.

644   The potential influence of dietary lipids and pharmacological interventions on exhaled VOLs

645   warrants further investigation. Since VOL profiles vary depending on the type of precursor

646   lipids (e.g., ω-3 PUFAs vs. ω-6 PUFAs) (**Fig. 2**), the specific VOLs to be targeted may differ

647   according to the subject's dietary habits, which in turn shape their internal lipid profile.

648   Therefore, to establish a more robust detection system, it is important to identify combinations

649   of VOL markers or correction strategies that minimize these confounding effects. In addition

650   to nonenzymatic reactions, lipid fragmentation products can be generated enzymatically via

651   lipid-oxidizing enzymes, such as lipoxygenases (LOX) and cyclooxygenases (COX)[36,37].

652 Accordingly, the impact of medications targeting these enzymes, such as non-steroidal anti-
653 inflammatory drugs (NSAIDs), on the VOL profile should be carefully examined.

654 Detection technologies and clinical diagnostic methods targeting exhaled breath allow
655 noninvasive examination of the physiological status of an individual. However, their
656 development has lagged significantly behind that of liquid biopsy technologies that utilize
657 blood and other bodily fluids. The rate-limiting step in their development is the identification
658 of gas marker molecules that are specific to biological phenomena or diseases. Therefore, using
659 this study as a model, gas markers can be explored as new indicators of metabolic reactions.
660 This can facilitate the development of detection technologies and clinical diagnostics.

661

662 **Conclusions**

663 We employed an innovative analytical platform, oxidative volatolomics, to identify VOLs
664 (1-octen-3-ol, 2-ethylfuran, and 2-pentylfuran) specifically released during ferroptosis. Using
665 these markers, we developed a technology for the noninvasive detection of ferroptosis *in vivo*.
666 We anticipate that the markers and detection technologies developed in this study will be
667 utilized and further developed, contributing to the noninvasive and early detection of
668 ferroptosis-related diseases.

669

670 **Supplementary Material**

671 Supplementary Information is available in the online version of this paper.

672

682

683 **Author Contributions**

684 Y.M. designed and performed the experiments, analyzed the data, and drafted and edited the

685 manuscript. Y.K., P.C., R.M., N.N., K.I., K.T., T.K., R.I., K.T., G. H., S.Y., H.T., G.I., Y.M., and

686 K.S. collected the samples for the clinical study. R.M. M.T. performed animal experiments.

687 K.H., N.T., and M.Y. analyzed the data. Y.K., N.T., S.F., and K.Y. drafted the manuscript. Y.S.

688 designed the experiments, conceived and supervised the project, and wrote and edited the

689 manuscript.

690

691 **Competing Interests**

692 The authors declare no competing financial conflicts of interest.

693

694 **Data Availability Statement**

695 All data generated in this study are provided in the main figures or Supplementary data. The

696 raw and processed MS data generated in this study are publicly available in the Metabolomics

697 Workbench repository under accession **doi: 10.21228/M8JK13.**

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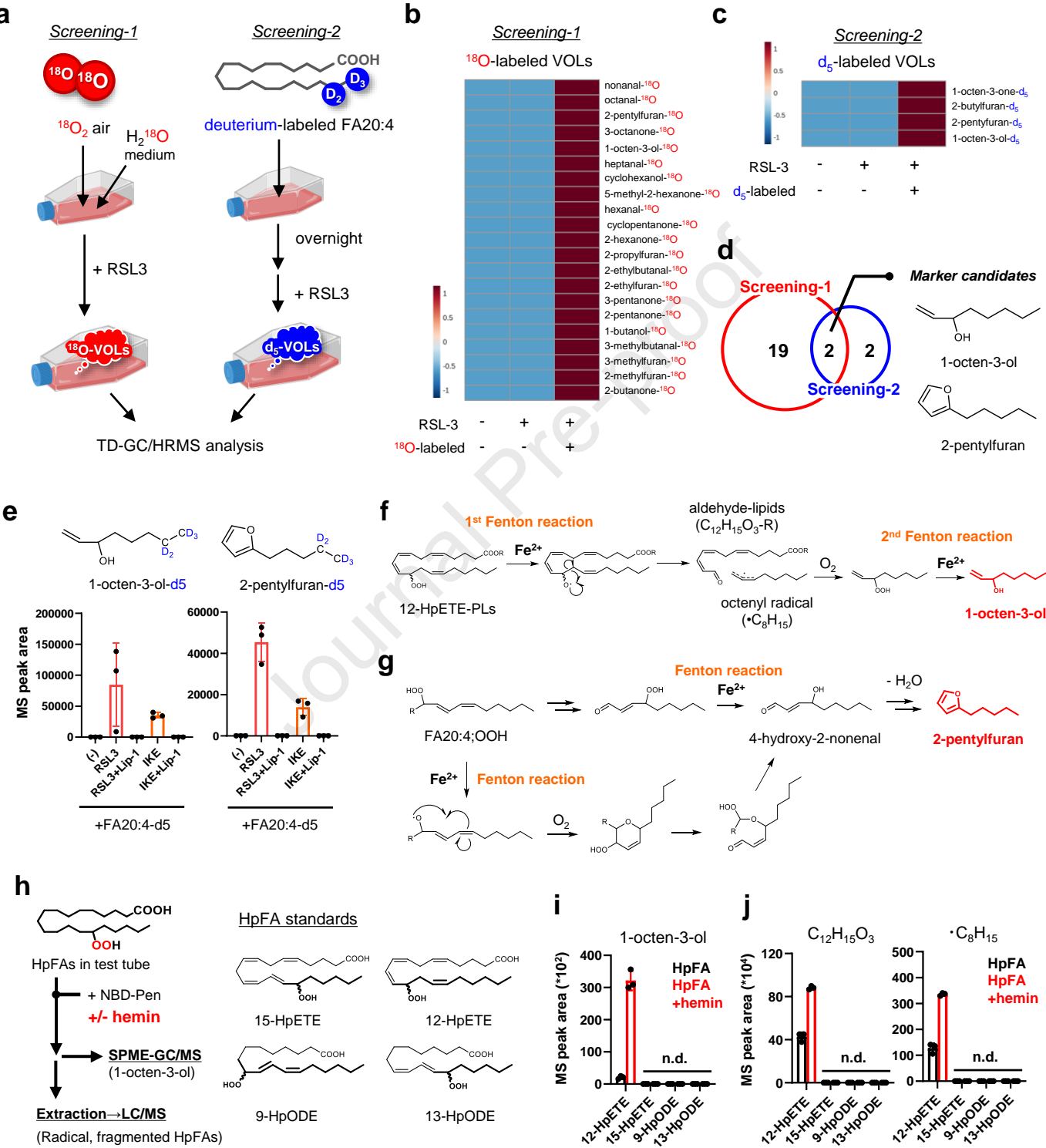863

Figure 1

Figure 1

Fig. 1: Identification of volatile oxidized lipids (VOLs) released during ferroptosis using volatile oxylipidomics

(a) Overview of volatile oxylipidomics. (b, c) Hierarchical cluster analysis of VOLs generated in each screening system. (d) Number of volatile organic compounds (VOC) hits identified in each screening and chemical structures of the commonly detected molecules (1-octen-3-ol and 2-pentylfuran). (e) HepG2 cells were supplemented with FA20:4-d5 to incorporate labeled fatty acids into cell membrane lipids and subsequently treated with 3.0 µM RSL3 or 20 µM IKE in the presence or absence of 1.0 µM Lip-1. After treatment, the amount of d5-labeled VOLs was quantified using thermal desorption–gas chromatography/high-resolution mass spectrometry (TD-GC/HRMS). (f, g) Plausible generation mechanisms of 1-octen-3-ol (f) and 2-pentylfuran (g). (h) Protocol for structural analysis of lipid fragments derived from hydroperoxy fatty acids (HpFAs) after addition of hemin and the chemical structures of the HpFAs used. (i, j) 1-octen-3-ol (i) and $C_{12}H_{15}O_3/{\bullet}C_8H_{15}$. black; HpFA only, red; HpFA+hemin. Data are presented as the mean ± standard deviation of three repeated experiments.

Figure 2

a

(1) *in vitro* LPO

Unsaturated FAs

• R   Fe

LPO initiators

(2) SPME-GC/MS analysis

VOLs

Peak annotation / Quantification

b

ω-3 PUFA derived VOLs

2-hexenal
3-hexenal
2,6-nonadienal
VOL-12
VOL-8
heptadienal-isomer-2
heptadienal-isomer-1
2-pentenal-isomer-2
2-pentenal-isomer-1
1-penten-3-one
VOL-10
2,3-pentaedione
VOL-7
2-ethylfuran
1-penten-3-ol
VOL-13
4-heptenal
VOL-6
VOL-4
VOL-3
VOL-5
VOL-11
VOL-9

ω-6 PUFA derived VOLs

2-nonenal
3-nonenal
nonadien-2-one_isomer-1
nonadien-2-one_isomer-2
2,6-dodecadien-1-al
2-octen-1-ol
2-octenal
2-butylfuran
2,4-decadienal
1-octen-3-one
2,3-octadiene
1-octen-3-ol
2,4-dodecadienal
2,4,7-tridecetrienal
4-decenal
heptanal
hexanoic acid methyl ester
pentanal
hexanal
VOL-2
2-pentylfuran
cis-4,5-epoxy-(E)-2-decenal
2,4-nonadienal
2-heptenal
VOL-1

FA18:2 FA20:4 FA18:2 FA20:4 FA20:5 FA22:6 FA20:5 FA22:6

AAPH   AAPH+hemin   AAPH   AAPH+hemin

ω-6 PUFAs   ω-3 PUFAs

c

OH

1-octen-3-ol

(*10^4)

MS peak area

1000
800
600
400
200
0

FA18:2 FA20:4 FA20:5 FA22:6

ω-6   ω-3

2-pentylfuran

(*10^3)

400
300
200
100
0

FA18:2 FA20:4 FA20:5 FA22:6

ω-6   ω-3

control
AAPH
AAPH +hemin

d

2-ethylfuran

(*10^3)

MS peak area

400
300
200
100
0

FA18:2 FA20:4 FA20:5 FA22:6

ω-6   ω-3

e

HepG2 (in 6-well plate)

D-labeled PUFA

Incubation (o.n.)

RSL3

SPME-GC/MS
(d-labeled VOLs)

SPME-GC/MS

d-VOLs

D-labeled PUFAs

FA18:2-d4

FA20:4-d8

FA22:6-d5

f

1-octen-3-ol

2H: *m/z* 57
2D: *m/z* 59
● = H or D

OH

FA18:2-d4
FA18:2
FA20:4-d8
FA20:4

0 20 40 60 80 100

Ratio of each isomer, %

2-pentylfuran

2H: *m/z* 138
2D: *m/z* 140
● = H or D

FA18:2-d4
FA18:2
FA20:4-d8
FA20:4

0 20 40 60 80 100

Ratio of each isomer, %

g

2-ethylfuran

5H: *m/z* 96
5D: *m/z* 101
● = H or D

FA22:6-d5
FA22:6

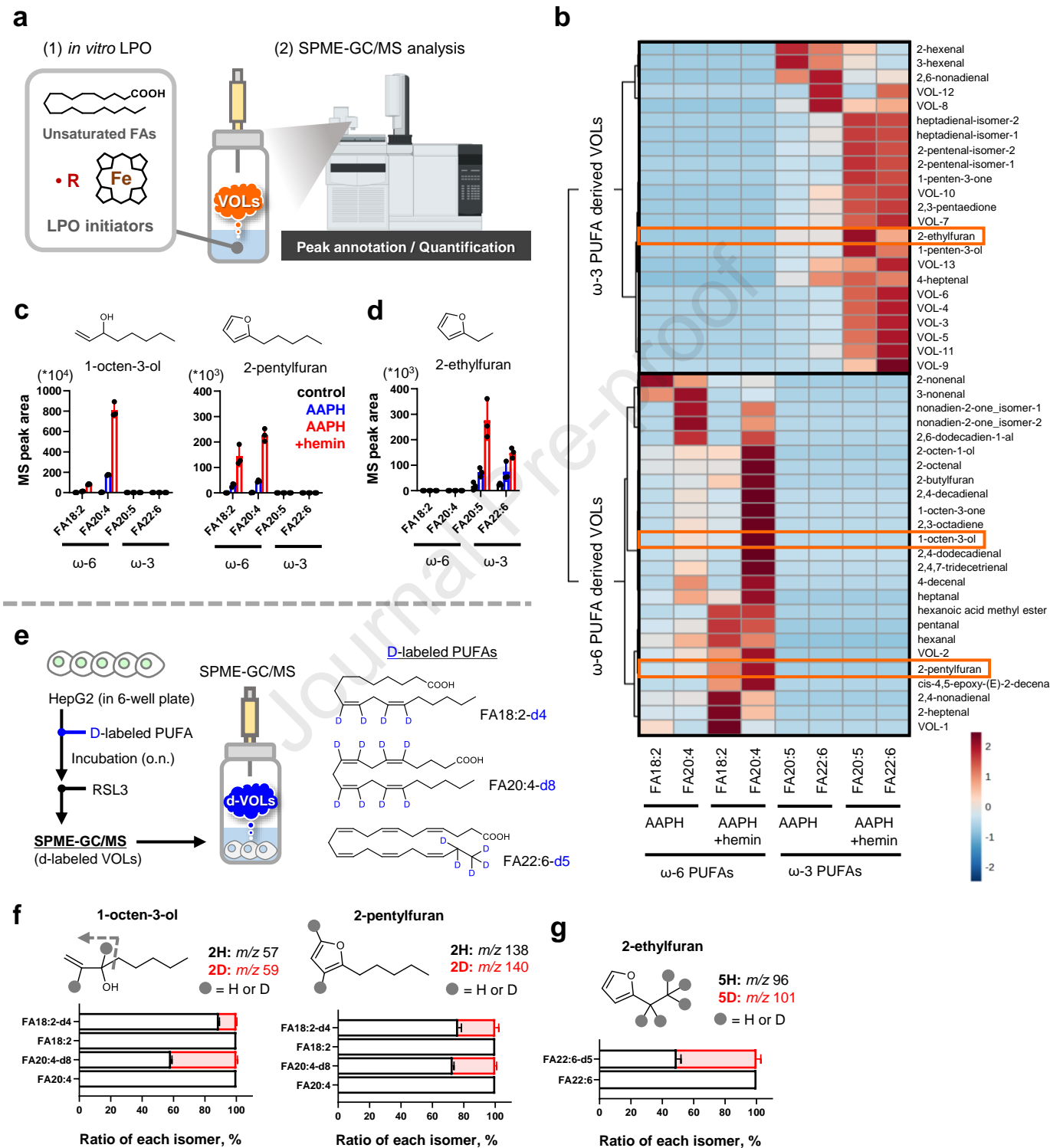0 20 40 60 80 100

Ratio of each isomer, %

Figure 2

Fig. 2: 1-octen-3-ol and 2-pentylfuran are produced from ω-6 PUFAs, and 2-ethylfuran from ω-3 PUFAs, through iron-dependent lipid peroxidation (LPO) during ferroptosis

(a) Outline of the SPME-GC/MS-based identification of PUFA-derived VOLs generated by AAPH or AAPH+hemin stimulation. (b) Hierarchical cluster analysis of VOLs generated in each peroxidation system. Original data are shown in **Supplementary Table 1**. (c, d) MS peak areas of typical ω-6/ω-3-PUFA-derived VOLs observed in each LPO system. (c) 1-octen-3-ol/2-pentylfuran and (d) 2-ethylfuran. (e) Strategy for deuterium (d)-labeling of intracellular PUFA-containing phospholipids and VOLs and the chemical structures of the d-labeled PUFA used. HepG2 cells were treated with 20 µM d-labeled PUFAs for 24 h, followed by the addition of 3.0 µM RSL3. (f, g) Isotopomer abundances of 1-octen-3-ol, 2-pentylfuran (f), and 2-ethylfuran (g) after (d-labeled) PUFA and RSL3 treatment. The MS peak area of each VOL sample was calculated from the corresponding ion chromatogram. Data are presented as the mean ± standard deviation of three repeated experiments.
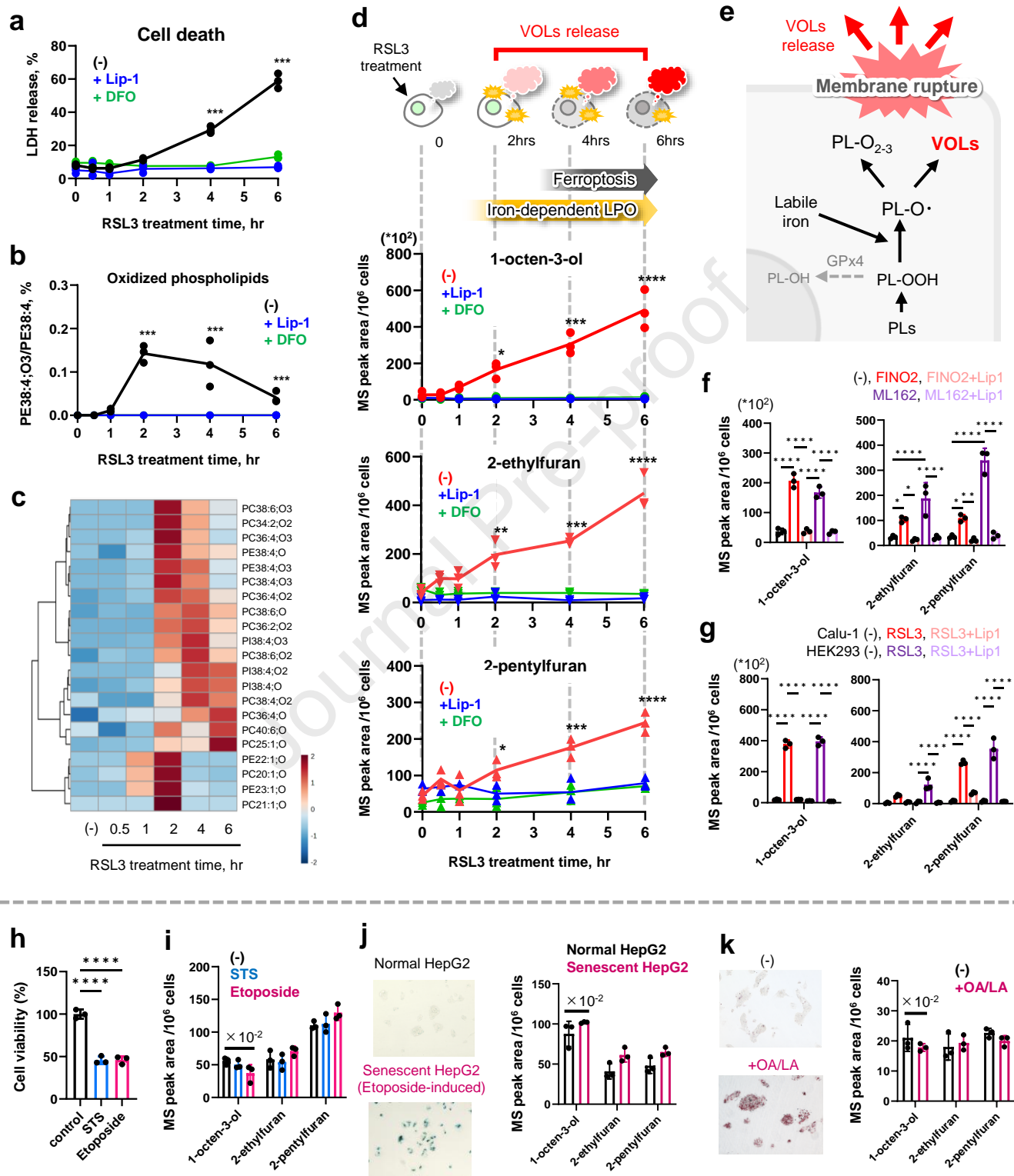
Figure 3

31

Figure 3

Journal Pre-proof

Fig. 3: Ferroptosis progression specifically leads to the release of 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran

(a–d) Time-course analysis of LDH release (a), oxidized lipid production (b, c), and the levels of VOLs in the cellular headspace (d) after treating HepG2 cells with 3.0 µM RSL3. (e) Plausible mechanisms for VOL formation and release during ferroptosis. (f) Levels of 1-octen-3-ol, 2-pentylfuran and 2-ethylfuran after treating HepG2 cells for 4 h with other ferroptosis inducers (25 µM FINO2, 3.0 µM ML162). (g) Levels of 1-octen-3-ol, 2-pentylfuran. and 2-ethylfuran following ferroptosis induction in Calu-1 and HEK293 cells using 0.5 µM RSL3. (h, i) Levels of 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran after treating cells with apoptosis inducers (10 µM staurosporine [STS], 100 µM etoposide) for 6 h. (j, k) Release of VOLs from etoposide-treated HepG2 cells (j) or HepG2 cells with lipid accumulation induced by oleic acid/linoleic acid (k). Data are presented as the mean ± standard deviation of six independent experiments. Statistical significance was assessed using a one-way analysis of variance with Tukey's multiple comparison test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$.
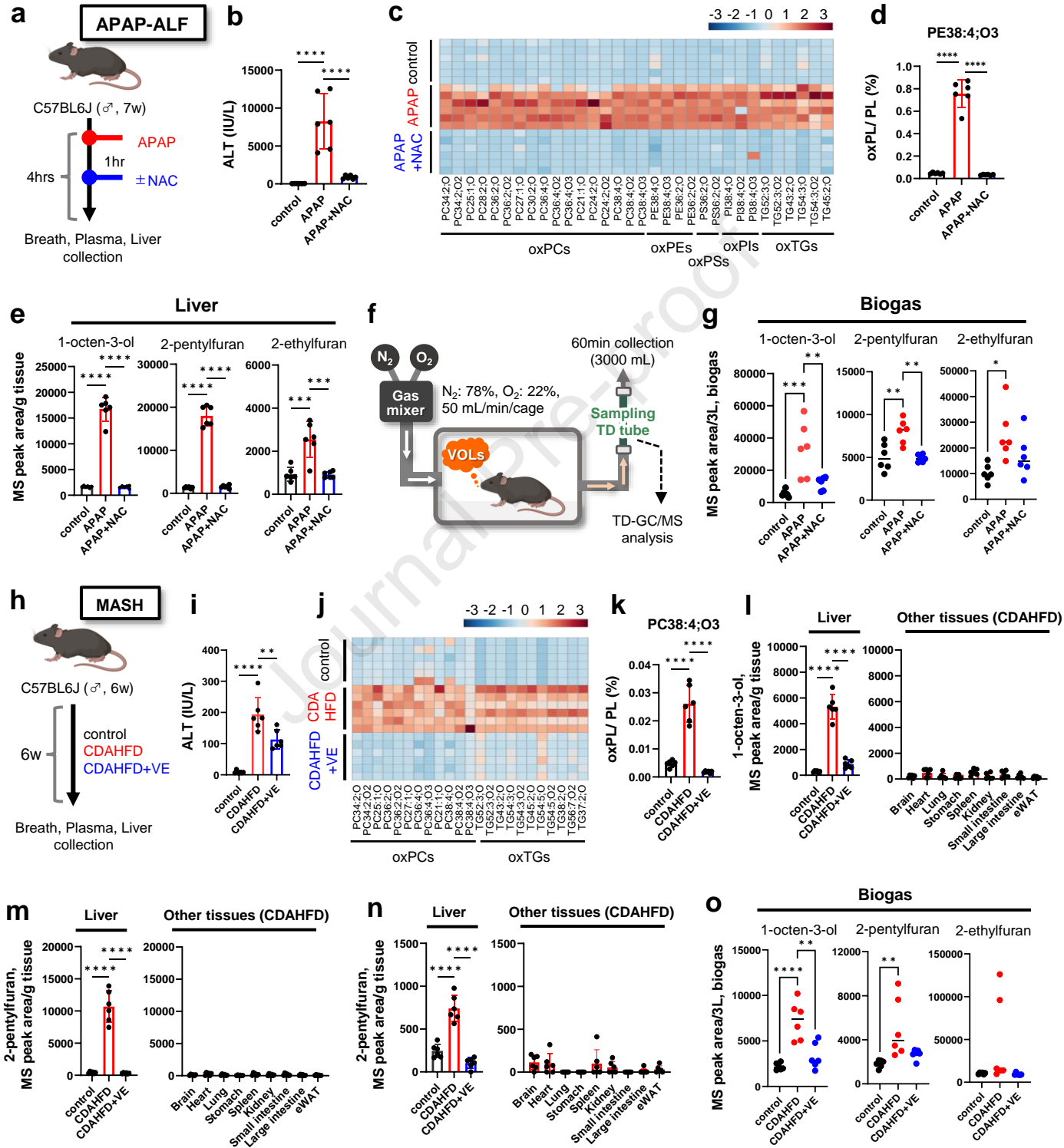
Figure 4

Figure 4

Fig. 4: Emission of ferroptosis VOL markers as biogas from acute or chronic liver disease model mice

(a) Experimental procedure for VOL analysis in APAP-ALF model mice. (b–e) Plasma alanine aminotransferase (ALT) levels (b), hepatic oxidized lipids (c, d), and hepatic VOL levels (e) in control and 300 mg/kg APAP ($\pm$ NAC)-treated groups. (f) Experimental setup for gas collection from mice. (g) Levels of 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran in biogas released from APAP-ALF model mice. (h) Experimental procedure for VOL analysis in choline-deficient, L-amino acid-defined, high-fat diet (CDAHFD)-fed metabolic dysfunction-associated steatohepatitis (MASH) model mice. (I–k) Plasma ALT levels (i) and hepatic oxidized lipid levels (j, k) in control and CDAHFD ($\pm$ VE)-fed groups. (l–o) Levels of 1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran in tissues (l–n) and biogas (o). Data are presented as the mean $\pm$ standard deviation of six independent experiments. Statistical significance was assessed using a one-way analysis of variance with Tukey's multiple comparison test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$.
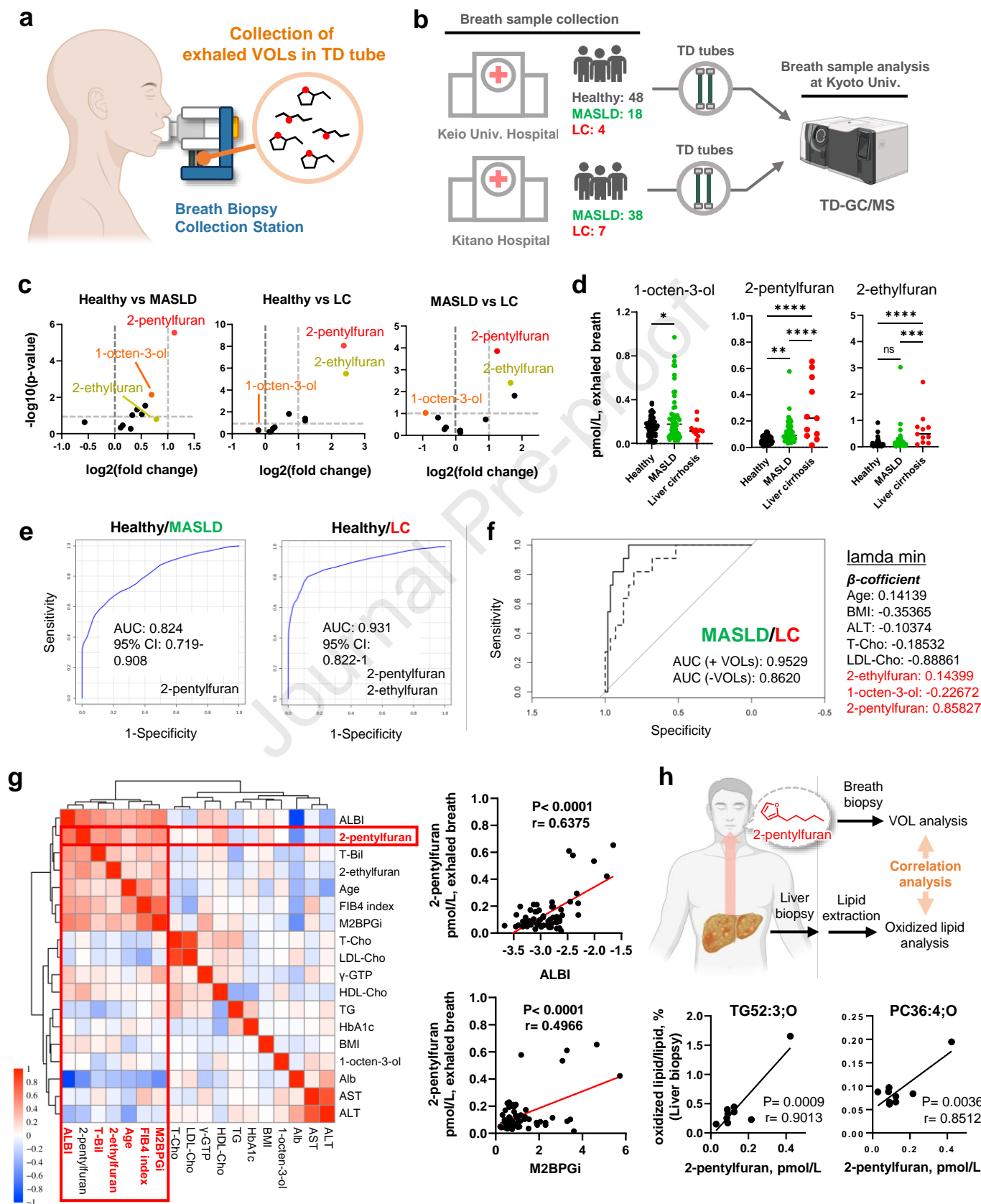
Figure 5

Figure 5

Fig. 5: Elevated exhaled levels of ferroptosis VOL markers in patients with metabolic dysfunction-associated steatotic liver disease/liver cirrhosis (MASLD/LC) compared with healthy individuals

(a) Collection of exhaled VOLs using the Breath Biopsy Collection Station. (b) Implementation framework for clinical trials. (c) Volcano plots depicting the distribution of VOLs in breath samples from patients with each disease condition. (d) Semi-quantification of ferroptosis VOL markers in the exhaled breath of each group. (e) ROC curve analysis of the VOLs of healthy vs MASLD (2-pentylfuran) and healthy vs LC (2-pentylfuran and 2-ethylfuran) comparisons. (f) ROC curve analysis using a combination of VOLs and clinical markers of the MASLD vs LC comparison. Solid line: ROC curve analysis for combined clinical markers and VOLs (AUC; 0.9529); dashed line: ROC curve analysis for clinical markers only (AUC; 0.8620). (g) Correlation analysis between ferroptosis VOL markers and clinical markers. ALBI; Albumin-bilirubin score, T-Bil; Total bilirubin, M2BPGi; Mac-2 binding protein glycosylation isomer, T-Cho; Total cholesterol, LDL-Cho; Low-density lipoprotein cholesterol, HDL-Cho; High-density lipoprotein cholesterol, TG; Triglyceride, Alb; Albumin. For the 2-pentylfuran and M2BPGi and ALBI correlation, two-sided Pearson's correlation (r) and p-values are indicated. (h) Correlation between oxidized lipid levels in liver biopsy samples and 2-pentylfuran level in exhaled breath. Statistical significance (d) was assessed using one-way analysis of variance with Tukey's multiple comparison test. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$.
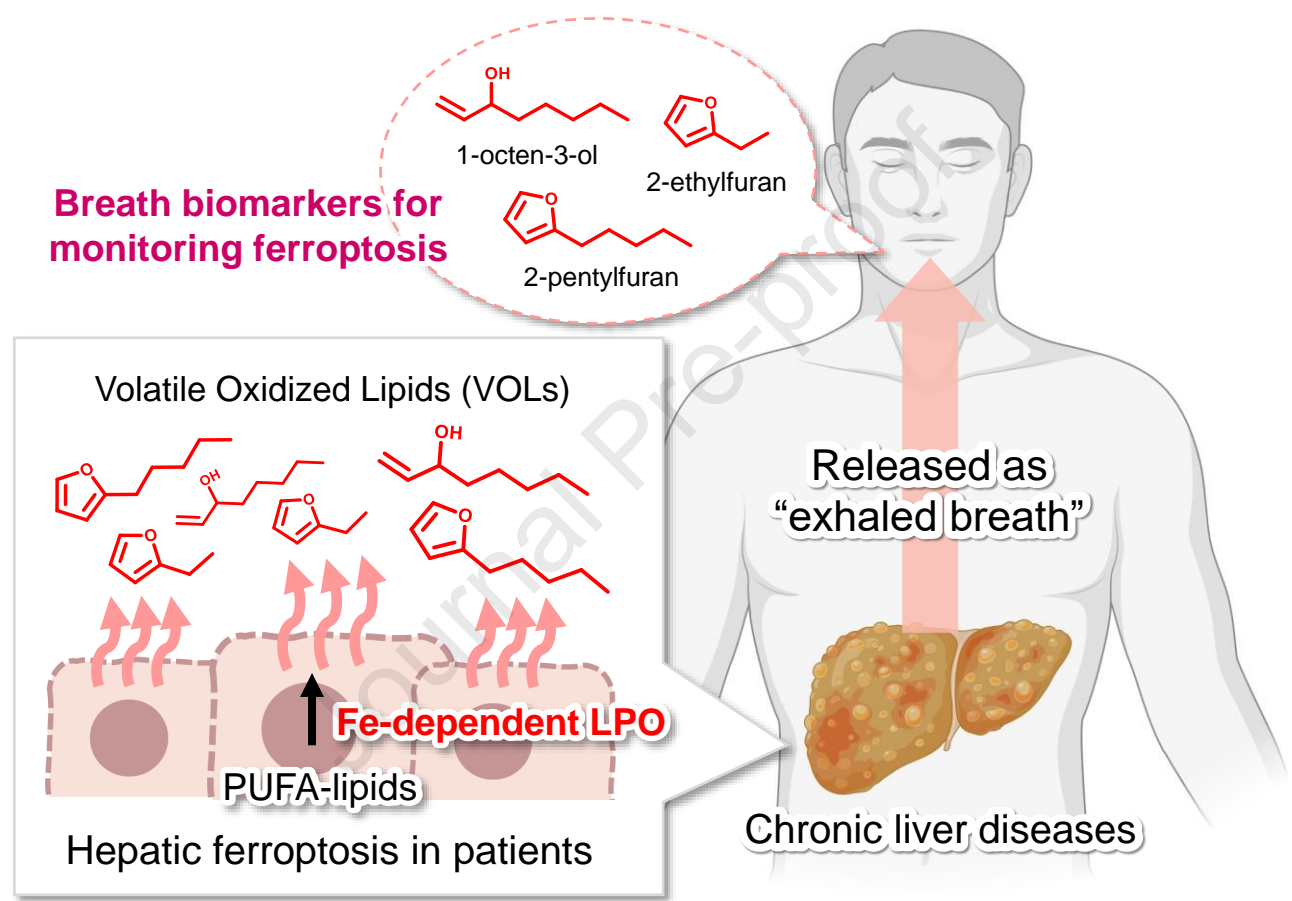
Figure 6

Fig. 6: Schematic illustration of the present study. VOL markers—1-octen-3-ol, 2-pentylfuran, and 2-ethylfuran—generated via iron-dependent lipid peroxidation and detectable in exhaled breath could serve as real-time, non-invasive indicators of in vivo ferroptosis and the progression of chronic liver diseases. The figure was created with BioRender (biorender.com).

Declaration of Interest Statement

☒ The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

☐ The author is an Editorial Board Member/Editor-in-Chief/Associate Editor/Guest Editor for this journal and was not involved in the editorial review or the decision to publish this article.

☐ The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: