

Detection of Volatile Organic Compounds from Preclinical Lung Cancer Mouse Models

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To cite this version:

Flora Gouzerh, Laurent Dormont, Bruno Buatois, Maxime Hervé, Maicol Mancini, et al.. Detection of Volatile Organic Compounds from Preclinical Lung Cancer Mouse Models. SSRN Electronic Journal, 2022, 10.2139/ssrn.4091348. hal-04399814

HAL Id: hal-04399814 <https://hal.science/hal-04399814v1>

Submitted on 20 Sep 2024

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Abstract

- Volatile organic compounds (VOCs) may help detect cancer tumour. This study addressed this question, as well
- as two methodological issues: 1) repeatability, comparing VOCs profiles obtained with two Solid Phase Micro
- Extraction (SPME) fibers used simultaneously; 2) detectability of cancer VOCs biomarkers, comparing profiles
- obtained following 1h versus 24h exposure of SPME fibers. We analyzed VOCs composition of soiled bedding
- obtained from a lung adenocarcinoma mouse model in which cancer was induced by doxycycline ingestion. We
- compared the VOCs profile of soiled bedding of cancerous (CC) and non-cancerous (NC) mice, before (T0),
- after two-weeks (T2) and after twelve weeks (T12) doxycycline ingestion. The results indicate : 1) qualitative
- and quantitative consistency in VOCs detection by two distinct SPME fibers ; 2) although more VOCs were
- detected following a 24h compared to 1h SPME exposure, none of the former molecules were related to cancer;
- 3) doxycycline impacted VOCs emissions in both CC and NC mice; 4) cancer impacted four VOCs at T12 only
- : the benzaldehyde which showed higher levels in CC mice and the hexan-1-ol and two mice pheromones, the 2-
- *sec*-butyl-4,5-dihydrothiazole and the 3,4-dehydro-*exo*-brevicomine, which showed lower levels in CC mice.
- Our study points out that the use of two SPME fibers and an extraction duration of 1h may be considered a good
- compromise allowing detection of cancer biomarkers while easing bench constraints.

- **Key Worlds** SPME, repeatability , odour signature, cancer biomarkers, EFGR cancer, *Mus musculus.*
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1. Introduction

 Volatile organic compounds (VOCs) are carbon-based small molecules, which are characterized by their volatility at ambient temperature. During the last decades, increasing attention has been devoted to exploring the relevance of VOCs emitted by the body as potential biomarkers of pathologies. VOCs are emitted and can be analysed from breath, skin, saliva, sweat, blood, urine and faeces. Nowadays, VOCs are commonly used as biomarkers of various diseases [1,2] such as fibrosis [3], asthma [4], Alzheimer disease [5], diabetes and tuberculosis [6,7], and an increasing attention has been given at using them for cancer detection [8].

 VOCs extraction involves the use of effective trapping methods, of which Solid-Phase Micro- Extraction (SPME) is the most commonly used [9]. For instance, this technique has been successfully used to detect VOC biomarkers in the context of diseases such as gastrointestinal disorders [10], cholera [11], tuberculosis [12] and cancers [13–15]. Accuracy and repeatability of VOC extraction strongly rely on the characteristics of the SPME fiber used [16] and on extraction conditions such as temperature and duration [17–19]. Some journals recommend using a single SPME in comparative studies, arguing that the use of more than one fiber in such studies may result in biased interpretations. However, when a study includes a large number of samples, it may be unpractical to use a single fiber for all experiment, and multiple successive use of the same fiber may also induce measurements biases due, for example, to wear. Another methodological issue is sampling duration. Emission of VOCs from biological substrate can be dynamic [20] and, in order to account for emission variations, comparative studies need to homogenize sampling duration. A short sampling duration may not allow capturing the diversity of VOCs emitted by the substrate nor detecting components present in very low concentrations, partly because competition between molecules present in the headspace may disadvantage the adsorption of components present in relatively small proportions [21–23].

 In the context of investigations of variation in VOCs composition induced by lung cancer in mice, the present study addressed repeatability of results obtained when sampling involved two different SPME fibers, and the impact of sampling duration on the detection of changes in the VOCs composition of mice soiled bedding during the course of tumour development. We also assessed whether VOCs induced by lung cancer were better detected after a 24h as compared to a 1h SPME exposure, and sought to identify murine lung adenocarcinoma VOC biomarkers. The odour sources were soiled beddings obtained from a genetically modified mouse model suffering from lung cancer, and considered an early and a late stage of tumour development.

- 2. Material and methods
- 2.1 Ethical clearance
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 All the precautions for animal welfare were followed during our experiments, and our study received ethic clearance from the Ethical Committee for Animal Experimentation (French Ministry of Higher Education, Research and Innovation) number 1645-22123.

2.2 Animals

100 Two genotype lines were involved in this study: WT and CCSP-rtTA / $EGFR^{T790M/L858R}$ [24,25]. The

101 WT mice, lacking both CCSP-rtTA and EGFR^{T790M/L858R} transgenes, do not develop any tumour upon

doxycycline (antibiotic) induction, hereafter non-cancerous (NC) mice, and the CCSP-rtTA /

103 EGFR^{T790M/L858R} mice, carrying both transgenes, develop lung adenocarcinoma upon doxycycline

104 induction, hereafter cancerous (CC) mice. To induce the EGFR^{T790M/L858R} in a lung specific manner,

 the mice received a doxycycline supplemented food (1mg/kg). The treatment lasted 12 weeks starting when the mice were 13 weeks old.

NC and CC male mice were obtained from the IRCM (Montpellier Cancer Research Institute) at the

age of 6 weeks and maintained at the breeding facilities of the IRD (Institute of Research and

Development) in Montpellier. Before the start of the experiment they were maintained in groups of 2

to 4 mice in transparent plastic cages (26.8cmW*21.5cmL*14.1cmH). Each cage contained sawdust, a

cellulose square, hay and a cardboard tunnel. The mice were observed and weighted once a week

during the entire experimental period to monitor their health. All mice were euthanized at the end of

the experiment at the IRCM at 25 weeks-of age. The non-cancerous status of all NC and the cancerous

 status of all CC mice were confirmed by histopathological analysis of haematoxylin & eosin stain of whole lung sections obtained after necropsy.

2.3 Odour source sampling

 Odour sources were two-week old soiled bedding of CC and NC mice. Each mouse was isolated at the age of 10 weeks in a cage containing 130g sawdust and a cellulose square. The housing conditions of all donors were homogenized as much as possible, and all mice were given the same quantity of food. Every two weeks, all the soiled bedding of every cage was collected in a plastic bag and kept at - 20 ° C and replaced by cleaned bedding (i.e. 130g sawdust). Soiled bedding analyzed in this study were pools of bedding obtained from 3 to 4 different mice. A sample of clean sawdust was also collected in a plastic bag and frozen to serve as a control for the VOCs analyses.

2.4 VOCs extraction

PMDS-DVD (polydimethylsiloxane and divinylbenzene) SPME fibers, recently purchased, and which

had never been used before, were used in this study (thickness 65μm, Stable Flex 24Ga, Sigma-

Aldrich, Bellefonte, USA). The odour samples were processed as follows. A frozen soiled bedding

was thawed while being maintained on ice during the entire procedure. Then a 3g aliquot was

transferred into a 125 ml glass vial, which was then sealed with a rubber septum. The VOCs sampling

phase took place in an oven maintained at 22°C. The SPME fiber was introduced into the vial after

piercing the septum with a needle after a 3min equilibration time, its position was always the same,

~2cm above the stimulus **(Fig. 1)**.

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2.5 VOCs characterisation by Gas Chromatography- Mass Spectrometry

 The SPME extract was injected into a Gas Chromatography - Mass Spectrometry (GC-MS), quadrupole mass spectrometer Shimadzu QP2010-SE (Shimadzu, Kyoto, 282 Japan) for further identification and relative quantification of its chemical content. Desorption was achieved by inserting the SPME fiber into the Ultra Inlet liner (for SPME/Purge and trap, 0.75mmID, Agilent CrossLab, Santa-Clara, USA) placed in Split/Splitless GC-MS injector heated at 250°C. The injection was made using a split ratio of 1:4 to allow a Gaussian form of low boiling point compounds and so, a better separation and further integration. The GC was equipped with an Optima 5-MS fused silica capillary column (30 m x 0.25 mm x 0.25 µm film thickness, Macherey-Nagel, Düren, Germany). Helium was used as the carrier gas (1 146 ml.min⁻¹). The GC temperature was maintained at 40 °C for 2 min, after which the temperature increased 147 by 5°C every minute till it reached 175 °C, and then by 12°C/min. till it reached 220°C.

2.6 Identification of the VOCs present in the extracts

 Spectra were analysed with the resident software (GCMS Solution, Shimadzu, Kyoto, Japan). Identification of compounds was based on chromatograms analysis of peak retention time (RT) and examination of their mass spectra. Retention indices (RI) were calculated using as reference the RI of a series of n-alkanes injected in the same apparatus (Alkanes standard solution, 04070, Sigma-Aldrich). Identification of compounds was made by both spectral and RI comparison with reference databases (NIST 2011, Wiley 292 Registry Ninth; e.g. Adams, 2007, Pubchem, 293 https://pubchem.ncbi.nlm.nih.gov/). Synthetic compounds were also used as a reference to confirm the identity of some VOCs. The peak surface area of each compound was based on the Total Ion Current Chromatogram (TICC). To be conservative, we excluded from the analyses compounds that were present in both our controls (clean bedding) and our biological samples, even when they were in higher proportion in the biological samples.

 Compounds identified with their mass spectra but present in the form of traces could not be quantified with the above described method and were given an arbitrary value corresponding to 10% of the surface area calculated for the smallest peak identified in the entire dataset. Peak areas for all compounds were then summed per chromatogram, and their relative proportion calculated as the ratio of the compound surface area divided by the sum of the surface areas of all other compounds identified in that chromatogram.

- 2.7 Testing the repeatability of odour sampling
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 We tested the repeatability of odour sampling among two different SPME fibers (F1 and F2) pertaining to the same new pack of 3 fibers (Sigma-Aldrich). The fibers were conditioned following 170 the manufacturer's recommendations, i.e. 30 minutes at 250°C. They were simultaneously introduced into a glass bottle containing the odour sample (3g of soiled bedding of 13-weeks old NC mice, Fig.1) and were exposed at the same time to the bottle headspace for 1h. The distance from the odour sample was controlled to be the same for the two fibers (~2cm), and the distance between the two fibers was ~1.5 cm. The same procedure was repeated at seven occasions with the same fibers and the aliquots 175 extracted were obtained from the same odour source, maintained at -20° between samplings. The 176 seven occasions corresponded to the first, $8th$, $15th$, $22th$, $30th$, $39th$ and $45th$ use of each fiber.

2.8 Testing the influence of sampling duration

 We analyzed the qualitative and quantitative variations of VOCs obtained from 18 samples of soiled bedding sampled with two fibers simultaneously, one exposed to the headspace for 1h the other for 181 24h. The odour sources were obtained from CC and NC mice at three experimental conditions: T0, i.e. before ingestion of doxycycline, T2, i.e. after two weeks' doxycycline treatment, and T12, i.e. after 12 weeks' doxycycline treatment. (Fig**. 1)**. We proceeded as follows, two SPME fibers were inserted into the same glass bottle containing soiled bedding and were exposed simultaneously to an odour source. The two fibers were placed at the same distance from this source and at a 1cm-distance from each other. After 1h exposure, one of the fibers was removed, while the other was left for another 23h. The procedure for the 18 samples lasted three days, and each day an additional glass bottle containing clean bedding was also sampled following the same protocol and served as a control.

2.9 Identifying VOCs biomarkers of murine lung cancer

 The same 18 odour sources as the above were processed. However, here each of the sources was represented by three replicates to control for sampling bias (i.e. collection of 3g aliquots for VOCs extraction). All together we analyzed 54 biological samples, In addition, a sample of clean bedding was extracted every day to serve as a control. Two SPME fibers were used alternatively for the purpose of this analysis, randomizing their use among NC and CC.

2.10 Statistical analyses

 All statistical analyses were performed with Rstudio version 3.4.4 [26]. We used the following packages: ade4 [27], vegan [28], mixOmics [29], ggplot2 [30], RVAideMemoire [31], Hotelling [32]. Variation of the relative proportions of VOCs was assessed with a multivariate approach using redundancy analyses (RDAs) followed by permutation F tests [33] and the R package vegan [28]. Relative proportions were CLR-transformed prior to RDAs, and as our data included zeroes, a small constant of an order of magnitude smaller than the smallest non-zero value was added to all values prior to the transformation (i.e. 0.01 if the smallest non-zero value is 0.1). Factors included in RDAs depended on the questions tested. The first model questioned the consistency of extraction by the two SPME fibers used simultaneously at 7 sampling occasions, and included "fiber identity" and "sampling occasions" as factors. The second RDA model addressed the impact of sampling/extraction duration on detection of cancer biomarker VOCs. Two models were considered. Because the number of VOCs detected after 24h extraction was higher than that after 1h extraction, the first model analysed only the data obtained after 210 24h and sought to identify presence of VOCs induced by cancer tumour, not detected after 1h extraction. 211 The model included experimental conditions (T0, T2, T12), health status (CC, NC) and their interactions as factors. The second model considered the VOCs identified both after 1h and 24h sampling and addressed whether quantitative differences existed. This model included duration (1h/24h), experimental conditions (T0, T2, T12), and health status (CC, NC) and their interactions as factors.

 Finally, the entire data set (56 samples corresponding to 3 replicates of each of 18 odour sources) obtained after 1h extraction was analysed using four RDA models to identify candidate cancer VOCs. In the two first models we addressed the impact of experimental condition (T0, T2 and T12) on VOCs composition for each health status separately. This model included experimental conditions as a fixed factor and replicates (3 per sample) and fiber identity (2 fibers used) as random factors (condition). The third model asked whether the VOCs composition of NC and CC at T0 was similar and included health status (NC/CC) as a fixed factor and replicates (3 per sample), and fiber identity (2 fibers used) as random factors (condition). The last model excluded the VOCs identified as different between NC and CC before the doxycycline treatment (at T0) and addressed whether the rest of the VOCs differed between health status, experimental conditions and their interactions (with replicate and fiber identity as random factors).

- 226 When relevant, we identified and listed the VOCs showing an absolute correlation coefficient > 0.8 with the RDA constrained axis associated with significant effects in a given model.
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- 3. Results
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- 3.1 Consistency of VOCs sampling across fibers and impact of multiple samplings

 We detected 89 compounds **(Table.S1)** in the technical controls (clean bedding). When compounds from this list were also detected in the biological samples at similar amounts (soiled bedding), they were not considered in the final list of compounds. The composition of the volatile profiles specific to the biological samples is given in **Table.1**. We identified the same 17 compounds with the two fibers (**Table.1**). Quantitative variation was assessed with a RDA analysis including the 17 compounds with "fiber identity" (2 modalities) and "sampling occasions" (7 modalities) as factors (model 1: Table 3). 239 VOCs composition did not vary significantly with fiber identity ($F = 0.957$, $P = 0.395$), while it did between sampling occasions (F= 34.185, P < 0.001) (**model 1 : Table.2**). Almost all compounds, except two, showed variations in proportions over time (Figure S1). However, these variations were 242 not linear and did not seem to relate to fiber wear. Although the same quantity of soiled bedding from the same odour source was analysed each time, variation in the quantity and quality of urine or faeces present in each aliquot might be the source of VOCs variation. These results further suggested that it might be important to use multiple replicates of every sample when comparing relatively small 246 quantities (here $3g$) of a complex substrate such as soiled bedding.

3.2 Influence of sampling duration on VOCs' detection: 1h versus 24h

 The same 17 compounds as in the first experiment were detected after 1h extraction and the 17 VOCs plus 11 others after 24h extraction (**Table.S2, Fig. 3).** A RDA was first performed to address if among 252 the 11 VOCs only found after 24h extraction any or all were related to cancer tumour development. We designed a model considering as fixed factors experimental conditions (T0, T2, T12), health status (CC, NC) and their interaction (**model 2: Table.2**). VOCs detected only after a 24-h extraction did not differ between NC and CC nor interactively with experimental conditions. Variation in VOCs composition 256 differed significantly only between experimental conditions $(F=3.910, P < 0.001)$, pointing out significant differences in the VOCs profiles at T0 compared to T2 and T12 **(model 2 : Table.2).** We then analysed the 17 VOCs common to the 1h and 24h extractions, and tested the same model with an additional fixed factor, sampling duration, and its interactions with the other two factors mentioned above (**model 3 : Table 2**). The results indicated a significant influence of the three main factors, but no interaction between sampling duration and health status (**Fig.4)** further suggesting that 24h extraction was not necessary to address the impact of cancer on the mice VOCs profile.

- 263 3.3 Cancer influence on VOCs production
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 We did not detect qualitative differences in VOCs composition between CC and NC mice (**Table.3**). To address possible quantitative differences, we proceeded in several steps. First we tested the impact of experimental conditions within the CC and within the NC samples. The two RDA analyses comprising replicates (3 per sample) and fiber identity (2 fibers used) as random factors, and experimental conditions (T0, T2 and T12) as fixed factor, indicated that the random factors explained a relatively negligible part of the total variation (9.60 % and 2.60% respectively for NC and for CC). The experimental conditions explained 24.87 % (NC) and 57.85% (CC) of the total variance. The VOCs composition varied significantly between the three experimental conditions, both for NC and CC mice (NC: F = 4.174 ; CC : F = 16.085 ; P < 0.001; post hoc tests all P< 0.001; **model 4 & 5 : Table.2),** further indicating the influence of experimental conditions on the VOCs profiles of both NC and CC mice.

 The second step consisted in evaluating the impact of cancer. To address this question, we first verified if the VOCs composition of CC and NC mice differed significantly before the start of the doxycycline diet, at T0, which was the case (**model 6 : Table.2)** indicating that the two genotypes had a different VOCs profile at the start of the experiment**.** Seven compounds had an absolute correlation > 0.8 with the RDA main axes, and hence were identified to be involved in the differences between CC and NC at T0 (**Table. 3)**. To further address the impact of cancer we removed these molecules from the data set and performed a RDA analysis including only the 10 VOCs that did not differ between CC and NC mice at T0 **(model 7 : Table.2 and Table.4).** The two random factors, replicates and fiber identity, accounted for a negligible part of the total variance, respectively, 3.32% and 1.05%. Both experimental conditions 285 (T0, T2 and T12), health status (NC and CC) and their interaction were found to be significant (all $P <$ 0.005; **Table.2**). As expected, we did not detect a significant difference in VOCs' composition between CC and NC mice at T0 (P=0.622)**(Table.4)**. At T2 the two mice types (CC, NC) were also characterised 288 by the same VOCs profile (P=0.495) but different from T0 (P<0.001), suggesting that at T2 only the doxycycline diet had altered the VOCs composition and that it influenced similarly CC and NC mice. The VOCs profiles of NC and CC were significantly different at T12 (P< 0.001; **Table.4**), and the CC profile at T12 was significantly different from that of CC at T2 and from that of NC both at T2 and T12, indicating that lung adenocarcinoma development influenced the VOCs' composition of CC mice **(Fig. 5**). The doxycycline treatment did not seem to have induced a change between T2 and T12 as far as NC 294 mice were concerned (T2-T12 NC, $P = 0.259$). However, our analyses pointed out 4 compounds differing between the VOCs profiles of CC at T2 and at T12. Three of these compounds, 3,4-dehydro- *exo*-brevicomine, 2-sec-butyl-4,5-dihydrothiazole and hexan-1-ol, were present in lower relative proportions in CC mice at T12 as compared to both NC and CC individuals at T2. The fourth compound, benzaldehyde, was present in higher relative proportions in CC at T12 (**Table.3**). Our analyses pointed out 4 other compounds impacted by the doxycycline treatment: the hexan-1-ol, 3-methyl-butan-1-ol, *(x)*-2,4,4-trimethyl-pent-2-enal and the fenchone. The hexan-1-ol and the 3-methyl-butan-1-ol were either absent or present as a track before the start of the doxycycline diet (**Table.3**), while *(x)*-2,4,4- trimethyl-pent-2-enal disappeared from the VOCs profiles after tumour induction by doxycycline. Concerning the fenchone, it decreased over time in NC mice, while in CC mice it decreased at T2 and then increased again at T12 (**Table.3)**.

4. Discussion

 Characterization of VOCs composition can quickly become tedious when one needs to analyze a large number of samples with a single SPME fiber, and when fiber exposure duration for each sample is long. However, when several fibers are used and extraction duration per sample is reduced, the repeatability of the results and the detectability of the molecules may be questioned. This paper addressed these two methodological issues and sought to identify VOCs present in mice soiled bedding that could be considered as cancer biomarkers.

Repeatability when using multiple SPME fibers.

 The use of several SPME fibers (with the same adsorption phase) to analyse a large number of samples might induce a bias due to inherent heterogeneity between fibers [34]. However, our results showed that, when all other conditions were controlled (duration, temperature etc.), we did not detect significant qualitative and/or quantitative difference in the VOCs profiles obtained by two distinct SPME fibers exposed simultaneously to the same source of odour. Our results also indicated that when two different fibers were used in a comparative study and fiber identity was included in the model as a random factor, the percentage of variation caused by this factor was marginal (**models 4 and 5 : Table.2**).

 Unexpectedly, our results also indicated strong variation in the VOCs profile of the same source of odour sampled at several occasions. The variation in the relative proportions of the different VOCs was not linear and did not suggest an effect of fiber wear (between the first and the 45th use). A possible explanation of our results might be sample heterogeneity. Soiled bedding is a complex substrate, containing saw dust, food leftover, urine, faeces and other body secretions, difficult to homogenise. For the purpose of our study, we used pools of two-weeks old soiled bedding that we mixed as much as possible before sampling our aliquots (3g). Our results suggest that some of the aliquots used in this part of our study were not similar, further suggesting that to avoid such bias larger volumes than 3g should be used, and if not possible several replicates should be used to control for sampling heterogeneity, which we did to address cancer impact on VOCs (see below).

Influence of sampling duration on detection of VOCs related to cancer.

 The release and adsorption of volatile compounds is dynamic [35–37]. Some molecules will be released immediately after exposure while others will not, this may be due to their chemical property and/or their state in the odour source (e.g. bound to urinary proteins [38,39] Low molecular weight VOCs will be quickly adsorbed by the SPME fiber while other molecules will take longer to be emitted and hence their detection could be delayed [20]. Increasing sampling duration can increase the number of molecules likely to be adsorbed. By increasing the duration of VOCs extraction to 24h, we were able to detect 11 additional molecules compared to the 1h sampling. Compounds detected only after a 24h sampling were mostly relatively heavy compounds that came out at the end of the chromatogram profile. The lower volatility of such molecules may explain why they were only observed in the 24h samples [40,41]. However, the increase in sampling duration did not necessarily result in an increase of the amount of compound trapped on the fiber for all compounds. For example, the relative proportions of some compounds such as 3,4 -dehydro-*exo*-brevicomine did not change between the 1h and 24h profiles while the proportion of thiazoline was higher in the 24h profiles compared to the 1h ones. This could be explained by the fact that while 3,4 -dehydro-*exo*-brevicomine is released during the first 20 minutes of sample exposure, thiazoline is emitted continuously for 24h [20]. Among the 11 molecules exclusively detected after 24h extraction hexanoic acid and *(E)-* nerolidol were described in mice urine that were heated at different temperatures between 45°C and 65°C, enhancing their volatility [42–45]. Some of the other compounds such as heptanoic acid and p- cymen -8 -ol are derived from microbial communities [46] and their emission was facilitated by heating or longer exposure to oxidative degradation as it is the case when extraction time lasts 24h. Nonetheless, among the 11 compounds detected exclusively after a 24h extraction, none differed

 between CC and NC mice, further indicating that 1h sampling should suffice to address cancer impact on mice VOC's profile (at least for such mouse model and such cancer type). Further, considering the cancer literature, none of the 11 molecules detected only in the 24h profiles was referenced as cancer biomarkers in mice. One compound was mentioned as a possible cancer biomarker in humans, the hexanoic acid, in gastric [47,48] and gastroesophageal cancer [49].

Cancer influence on VOCs production.

 The use of soiled bedding allows to sample VOCs emitted by mice in their everyday activities in their cages, making the sampling non-invasive and non-stressful for the mice. It also allows collection of VOCs from a variety of sources simultaneously such as faeces, urine, saliva, preputial gland etc. Among the 17 compounds identified in this study, seven had not been reported in another publication on mice to our knowledge **(Table.3).** The other compounds, such as propanoic acid, hexan-1-ol or butanoic acid, were described before [44]. The propanoic acid was reported to be emitted from the body or from faeces,

 while other compounds such as benzaldehyde was reported to be present in mice urine [43,44,50]. Some compounds that we found both in the control and in the biological samples, that we decided not to consider here, were reported as mouse compounds in the literature, such as acetic acid, pentan-1-ol, heptanal [44], nonanal [44,51] or again p-cymene [51].

 Our results revealed significant quantitative differences in the VOCs profiles of the two mice lines (NC, CC) before doxycycline treatment (at T0). As these mice were kept in exactly the same conditions, and were all same age males and litter mates, the only difference that we see that could explain their different VOCs profile is their relatively slight different genetic background: presence versus lack of CCSP-rtTA 377 and EGFR^{T790M/L858R}. Small differences in genetic background were already shown to impact the VOCs profile of mice, [52–54]. Seven compounds differed between NC and CC at T0. Except for the 3- methylbutan-1-ol, these compounds were present in greater relative proportion in NC than in CC mice. Among the molecules described in the mouse VOCs literature, only the propanoic acid and the 3- methylbutan-1-ol were reported [55], the other four compounds, nopinone, camphor, camphene hydrate and *(E)*-3-pinocamphone are terpenes also found in plants.

- Our study revealed the effect of both doxycycline and cancer tumour development on the mice VOCs profile. Doxycyline diet seems to influence VOCs emissions both in NC and CC mice. One compounds disappeared from the profile (e.g. (x)-2,4,4-trimethyl-pent-2-enal) while two other compounds were present in higher proportions after the addition of doxycycline, i.e. hexan-1-ol and 3-methylbutanol, which are compounds notably emitted by bacteria [56], indicating that the doxycycline treatment, by eliminating some bacteria, might have favoured others, impacting the mice gut microbiome composition and thus inducing a change in the VOCs emitted by the mice [57].
- Our results indicated that the impact of cancer on VOCs composition was only detectable at T12, i.e. after 12 weeks of treatment when all mice harboured lung adenocarcinoma [25], and that this impact involved four compounds. Benzaldehyde was found in higher proportions in CC than NC mice at T12. It is a compound commonly found in mice urine [44,58–60], and was already reported in studies using mice to study cancer [61–63]. Benzaldehyde was also found in human cancers and might be considered as a relatively general biomarker of lung [64–67], colorectal [68] or breast cancer [69]. Most of the time this compound was studied in cell cultures and was found to increase in cancerous cells, here we show that cancer impact the whole body emission. Benzaldehyde was also found to be discriminating in studies on Alzheimer's disease [70]. The three other candidate VOCs identified in this study to be impacted by cancer development were found in smaller proportions in CC as compared to NC mice, namely, hexan-1-ol, 2-*sec*-butyl-4,5-dihydrothiazole (Thiazoline) and 3,4-dehydro-*exo*-brevicomine (Brevicomine). Of these three compounds, two are known mice pheromones described in male mice, brevicomine and thiazoline. These pheromones were shown to be involved in male dominance and male attractiveness to female mice [71]. Thiazoline was pointed as a candidate in another mouse lung cancer

 study [72]. The onset of cancer therefore seems to impact pheromone emissions in sick mice, and hence is expected to impact their social life. The third candidate molecule is the hexan-1-ol also found in mice urine [44]. Like in our study, hexan-1-ol was found in smaller proportions in cancerous as compared to non-cancerous individuals in investigations of human lung cancer [63], melanoma [73] and head and neck cancer [74]Finally, among the compounds identified in similar proportions in both NC and CC sources in our study, the camphor was found to discriminate against colorectal cancer in mice [75], and the 3 methylbutanoic acid was found to discriminate melanoma [73].

5. Conclusion

 We were able to show that when used under strict controlled conditions, the use of different SPME fibers may have a marginal impact on the variance observed between samples. The use of two fibers allowed us to perform a larger number of samples in a relatively short time. In addition, our study validated the use of mouse soiled bedding to extract VOCs from mice without being invasive. A longer sampling period may make it easier to identify compounds present in small amounts during a shorter sampling period. However, in our study, 1h extraction was sufficient to isolate the compounds that differentiated cancerous from non-cancerous mice. We did not detect cancer induced VOCs variation at an early stage of tumour development (T2), although behavioural investigations showed that mice discriminated between CC and NC soiled bedding at T2 [76]. The latter results suggest either, that our study method was not sensitive enough (compared to a mouse nose) to detect informative VOCs at T2, or most probably that other molecules, e.g. proteins, might be impacted by tumour development [77]. We identified four candidate compounds that discriminated CC from NC soiled bedding at T12: the benzaldehyde and the hexan-1-ol confirming earlier studies on mice and humans cancer, and two mice pheromones the Thiazoline and the Brevicomine, indicating that cancer could potentially impact social and sexual interactions in mice natural populations and, as such, might be selected against.

REFERENCES

- [1] M. Shirasu, K. Touhara, JB Review The scent of disease : volatile organic compounds of the human body related to disease and disorder, 150 (2011) 257–266. https://doi.org/10.1093/jb/mvr090.
- [2] S. Sethi, R. Nanda, T. Chakraborty, Clinical application of volatile organic compound analysis for detecting infectious diseases, Clinical Microbiology Reviews. 26 (2013) 462–475. https://doi.org/10.1128/CMR.00020-13/ASSET/BCFA621B-3481-44BA-9028- 04A98288403B/ASSETS/GRAPHIC/ZCM9990924260004.JPEG.
- [3] M. Barker, M. Hengst, J. Schmid, H.J. Buers, B. Mittermaier, D. Klemp, R. Koppmann, Volatile organic compounds in the exhaled breath of young patients with cystic fibrosis, European Respiratory Journal. 27 (2006) 929–936. https://doi.org/10.1183/09031936.06.00085105.
- [4] B. Ibrahim, M. Basanta, P. Cadden, D. Singh, D. Douce, A. Woodcock, S.J. Fowler, Non-invasive phenotyping using exhaled volatile organic compounds in asthma, Thorax. 66 (2011) 804–809. https://doi.org/10.1136/thx.2010.156695.
- [5] A. Mazzatenta, M. Pokorski, F. Sartucci, L. Domenici, C. di Giulio, Volatile organic compounds (VOCs) fingerprint of Alzheimer's disease, Respiratory Physiology & Neurobiology. 209 (2015) 81–84. https://doi.org/10.1016/j.resp.2014.10.001.
- [6] M. Phillips, R.N. Cataneo, T. Cheema, J. Greenberg, Increased breath biomarkers of oxidative stress in diabetes mellitus, Clinica Chimica Acta. 344 (2004) 189–194. https://doi.org/10.1016/j.cccn.2004.02.025.
- [7] M. Phillips, R.N. Cataneo, R. Condos, G.A. Ring Erickson, J. Greenberg, V. la Bombardi, M.I. Munawar, O. Tietje, Volatile biomarkers of pulmonary tuberculosis in the breath, Tuberculosis. 87 (2007) 44–52. https://doi.org/10.1016/j.tube.2006.03.004.
- [8] F. Gouzerh, J.-M. Bessière, B. Ujvari, F. Thomas, A.M. Dujon, L. Dormont, Odors and cancer: Current status and future directions, Biochimica et Biophysica Acta (BBA) - Reviews on Cancer. (2021) 188644. https://doi.org/10.1016/J.BBCAN.2021.188644.
- [9] Z. Zhang, J. Pawliszyn,) Zhang, Z.; Pawliszyn, J. /. High Resolut. Chromatogr. 67 (1995) 1477– 1483. https://pubs.acs.org/sharingguidelines (accessed November 16, 2021).
- [10] C.E. Garner, S. Smith, B. Lacy Costello, P. White, R. Spencer, C.S.J. Probert, N.M. Ratcliffem, Volatile organic compounds from feces and their potential for diagnosis of gastrointestinal disease, The FASEB Journal. 21 (2007) 1675–1688. https://doi.org/10.1096/fj.06-6927com.
- [11] C.E. Garner, S. Smith, P.K. Bardhan, N.M. Ratcliffe, C.S.J. Probert, A pilot study of faecal volatile organic compounds in faeces from cholera patients in Bangladesh to determine their utility in disease diagnosis, Trans R Soc Trop Med Hyg. 103 (2009) 1171–1173. https://doi.org/10.1016/J.TRSTMH.2009.02.004/2/M_103-11-1171-FIG001.GIF.
- [12] M. Syhre, S.T. Chambers, The scent of Mycobacterium tuberculosis, Tuberculosis. 88 (2008) 317–323. https://doi.org/10.1016/J.TUBE.2008.01.002.
- [13] H. Liu, H. Wang, C. Li, L. Wang, Z. Pan, L. Wang, Investigation of volatile organic metabolites in lung cancer pleural effusions by solid-phase microextraction and gas chromatography/mass spectrometry, Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences. 945–946 (2014) 53–59. https://doi.org/10.1016/j.jchromb.2013.11.038.
- [14] D. Poli, M. Goldoni, M. Corradi, O. Acampa, P. Carbognani, E. Internullo, A. Casalini, A. Mutti, Determination of aldehydes in exhaled breath of patients with lung cancer by means of on- fiber-derivatisation SPME-GC/MS, Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences. 878 (2010) 2643–2651. https://doi.org/10.1016/j.jchromb.2010.01.022.
- [15] J. Rudnicka, T. Kowalkowski, T. Ligor, B. Buszewski, Determination of volatile organic compounds as biomarkers of lung cancer by SPME-GC-TOF/MS and chemometrics, Journal of

 [16] E. Dixon, C. Clubb, S. Pittman, L. Ammann, Z. Rasheed, Solid-Phase Microextraction and the Human Fecal VOC Metabolome, PLoS ONE. 6 (2011) 18471. https://doi.org/10.1371/journal.pone.0018471. [17] H. Lord, J. Pawliszyn, Evolution of solid-phase microextraction technology, Journal of Chromatography A. 885 (2000) 153–193. https://doi.org/10.1016/S0021-9673(00)00535-5. [18] W. Miekisch, J.K. Schubert, D.A. Vagts, K. Geiger, Analysis of Volatile Disease Markers in Blood, Clinical Chemistry. 47 (2001) 1053–1060. https://doi.org/10.1093/CLINCHEM/47.6.1053. [19] T. Živković Semren, I. Brčić Karačonji, T. Safner, N. Brajenović, B. Tariba Lovaković, A. Pizent, Gas chromatographic-mass spectrometric analysis of urinary volatile organic metabolites: Optimization of the HS-SPME procedure and sample storage conditions, Talanta. 176 (2018) 537–543. https://doi.org/10.1016/J.TALANTA.2017.08.064. 491 [20] A. Cavaggioni, C. Mucignat-Caretta, M. Redaelli, G. Zagotto, The scent of urine spots of male mice, Mus musculus: changes in chemical composition over time, Rapid Communications in Mass Spectrometry. 20 (2006) 3741–3746. https://doi.org/10.1002/RCM.2789. [21] J. Song, B.D. Gardner, J.F. Holland, R.M. Beaudry, Rapid Analysis of Volatile Flavor Compounds in Apple Fruit Using SPME and GC/Time-of-Flight Mass Spectrometry, (1997). https://pubs.acs.org/sharingguidelines (accessed January 25, 2022). [22] T. Gó, P. Martos, J. Pawliszyn, Strategies for the Analysis of Polar Solvents in Liquid Matrixes, (1997). https://pubs.acs.org/sharingguidelines (accessed January 25, 2022). [23] E.E. Stashenko, J.R. Martínez, Sampling volatile compounds from natural products with headspace/solid-phase micro-extraction, Journal of Biochemical and Biophysical Methods. 70 (2007) 235–242. https://doi.org/10.1016/J.JBBM.2006.08.011. [24] D. Li, T. Shimamura, H. Ji, L. Chen, H.J. Haringsma, K. McNamara, M.C. Liang, S.A. Perera, S. Zaghlul, C.L. Borgman, S. Kubo, M. Takahashi, Y. Sun, L.R. Chirieac, R.F. Padera, N.I. Lindeman, P.A. Jänne, R.K. Thomas, M.L. Meyerson, M.J. Eck, J.A. Engelman, G.I. Shapiro, K.K. Wong, Bronchial and Peripheral Murine Lung Carcinomas Induced by T790M-L858R Mutant EGFR Respond to HKI-272 and Rapamycin Combination Therapy, Cancer Cell. 12 (2007) 81–93. https://doi.org/10.1016/j.ccr.2007.06.005. [25] E.B. Mur, S. Bernardo, L. Papon, M. Mancini, E. Fabbrizio, M. Goussard, I. Ferrer, A. Giry, X. Quantin, J.L. Pujol, O. Calvayrac, H.P. Moll, Y. Glasson, N. Pirot, A. Turtoi, M. Cañamero, K.K. Wong, Y. Yarden, E. Casanova, J.C. Soria, J. Colinge, C.W. Siebel, J. Mazieres, G. Favre, L. Paz- Ares, A. Maraver, Notch inhibition overcomes resistance to tyrosine kinase inhibitors in EGFR-driven lung adenocarcinoma, J Clin Invest. 130 (2020) 612–624.

Chromatography B: Analytical Technologies in the Biomedical and Life Sciences. 879 (2011)

3360–3366. https://doi.org/10.1016/j.jchromb.2011.09.001.

- https://doi.org/10.1172/JCI126896.
- [26] R Core Team, A language and environmentfor statistical computing. R Foundation forStatistical Computing, Vienna, Austria., (2020). lhttps://www.r-project.org/.
- [27] J. Thioulouse, A.B. Dufour, T. Jombart, S. Dray, A. Siberchicot, S. Pavoine, Multivariate analysis of ecological data with ade4, Multivariate Analysis of Ecological Data with Ade4. (2018) 1–294. https://doi.org/10.1007/978-1-4939-8850-1.
- [28] J. Oksanen, F.G. Blanchet, M. Friendly, R. Kindt, P. Legendre, D. Mcglinn, P.R. Minchin, R.B. O'hara, G.L. Simpson, P. Solymos, M. Henry, H. Stevens, E. Szoecs, H.W. Maintainer, Package "vegan" Title Community Ecology Package Version 2.5-7, (2020).
- [29] F. Rohart, B. Gautier, A. Singh, K.A. Lê Cao, mixOmics: An R package for 'omics feature selection and multiple data integration, PLOS Computational Biology. 13 (2017) e1005752. https://doi.org/10.1371/JOURNAL.PCBI.1005752.
- [30] M.H. Wickham, Package "ggplot2" Type Package Title An implementation of the Grammar of Graphics, (2014).
- [31] M. Hervé, Package "RVAideMemoire" Encoding latin1 Type Package Title Testing and Plotting Procedures for Biostatistics, (2022).
- [32] J. Curran, Package "Hotelling" Title Hotelling's T^2 Test and Variants, (2021).
- [33] M.R. Hervé, F. Nicolè, K.-A. Lê Cao, Multivariate Analysis of Multiple Datasets: a Practical Guide for Chemical Ecology, Journal of Chemical Ecology 2018 44:3. 44 (2018) 215–234. https://doi.org/10.1007/S10886-018-0932-6.
- [34] G.A. Gõmez-Ríos, N. Reyes-Garcés, J. Pawliszyn, Evaluation of a multi-fiber exchange solid- phase microextraction system and its application to on-site sampling, Journal of Separation Science. 38 (2015) 3560–3567. https://doi.org/10.1002/JSSC.201500158.
- [35] R.E. Shirey, Optimization of Extraction Conditions for Low-Molecular-Weight Analytes Using Solid-Phase Microextraction, Journal of Chromatographic Science. 38 (2000) 109–116. https://doi.org/10.1093/CHROMSCI/38.3.109.
- [36] J. Pawliszyn, Theory of Solid-Phase Microextraction, Handbook of Solid Phase Microextraction. (2012) 13–59. https://doi.org/10.1016/B978-0-12-416017-0.00002-4.
- [37] C. Liaud, M. Brucher, C. Schummer, C. Coscollà, H. Wolff, J.J. Schwartz, V. Yusà, M. Millet, Utilization of long duration high-volume sampling coupled to SPME-GC-MS/MS for the assessment of airborne pesticides variability in an urban area (Strasbourg, France) during agricultural application, Http://Dx.Doi.Org/10.1080/03601234.2016.1191916. 51 (2016) 703– 714. https://doi.org/10.1080/03601234.2016.1191916.
- [38] R.J. Beynon, J.L. Hurst, Urinary proteins and the modulation of chemical scents in mice and rats, Peptides (N.Y.). 25 (2004) 1553–1563. https://doi.org/10.1016/J.PEPTIDES.2003.12.025.
- [39] J. Kwak, C.C. Grigsby, M.M. Rizki, G. Preti, M. Köksal, J. Josue, K. Yamazaki, G.K. Beauchamp, Differential binding between volatile ligands and major urinary proteins due to genetic variation in mice, Physiology & Behavior. 107 (2012) 112–120. https://doi.org/10.1016/J.PHYSBEH.2012.06.008.
- [40] M.P. Gianelli, M. Flores, F. Toldrá, Optimisation of solid phase microextraction (SPME) for the analysis of volatile compounds in dry-cured ham, Journal of the Science of Food and Agriculture. 82 (2002) 1703–1709. https://doi.org/10.1002/JSFA.1249.
- [41] A. Marco, J.L. Navarro, M. Flores, Volatile compounds of dry-fermented sausages as affected by solid-phase microextraction (SPME), Food Chemistry. 84 (2004) 633–641. https://doi.org/10.1016/S0308-8146(03)00288-7.
- [42] C. Mucignat-Caretta, M. Redaelli, A. Orsetti, M. Perriat-Sanguinet, G. Zagotto, G. Ganem, Urinary Volatile Molecules Vary in Males of the 2 European Subspecies of the House Mouse and Their Hybrids, Chemical Senses. 35 (2010) 647–654. https://doi.org/10.1093/CHEMSE/BJQ049.
- [43] M.L. Schaefer, K. Wongravee, M.E. Holmboe, N.M. Heinrich, S.J. Dixon, J.E. Zeskind, H.M. Kulaga, R.G. Brereton, R.R. Reed, J.M. Trevejo, Mouse Urinary Biomarkers Provide Signatures of Maturation, Diet, Stress Level, and Diurnal Rhythm, Chemical Senses. 35 (2010) 459–471. https://doi.org/10.1093/CHEMSE/BJQ032.
- [44] F. Röck, S. Mueller, U. Weimar, H.-G. Rammensee, P. Overath, Comparative Analysis of Volatile Constituents from Mice and their Urine, Journal of Chemical Ecology. 32 (2006) 1333– 1346. https://doi.org/10.1007/s10886-006-9091-2.
- [45] B.S. Goodrich, S. Gambale, P.R. Pennycuik, T.D. Redhead, Volatiles from feces of wild male house mice, Journal of Chemical Ecology 1990 16:7. 16 (1990) 2091–2106. https://doi.org/10.1007/BF01026922.
- [46] Z.R. Huang, W.L. Guo, W. bin Zhou, L. Li, J.X. Xu, J.L. Hong, H.P. Liu, F. Zeng, W.D. Bai, B. Liu, L. Ni, P.F. Rao, X.C. Lv, Microbial communities and volatile metabolites in different traditional fermentation starters used for Hong Qu glutinous rice wine, Food Research International. 121 (2019) 593–603. https://doi.org/10.1016/J.FOODRES.2018.12.024.
- [47] S. Kumar, J. Huang, N. Abbassi-Ghadi, P. Španěl, D. Smith, G.B. Hanna, Selected ion flow tube mass spectrometry analysis of exhaled breath for volatile organic compound profiling of esophago-gastric cancer, Analytical Chemistry. 85 (2013) 6121–6128. https://doi.org/10.1021/ac4010309.
- [48] S. Kumar, J. Huang, N. Abbassi-Ghadi, H.A. MacKenzie, K.A. Veselkov, J.M. Hoare, L.B. Lovat, P. Spanel, D. Smith, G.B. Hanna, Mass spectrometric analysis of exhaled breath for the identification of volatile organic compound biomarkers in esophageal and gastric adenocarcinoma, Annals of Surgery. 262 (2015) 981–990. https://doi.org/10.1097/SLA.0000000000001101.
- [49] J. Huang, S. Kumar, N. Abbassi-Ghadi, P. Španěl, D. Smith, G.B. Hanna, Selected ion flow tube mass spectrometry analysis of volatile metabolites in urine headspace for the profiling of gastro-esophageal cancer, Analytical Chemistry. 85 (2013) 3409–3416. https://doi.org/10.1021/ac4000656.
- [50] F.J. Schwende, D. Wiesler, J.W. Jorgenson, M. Carmack, M. Novotny, Urinary volatile constituents of the house mouse,Mus musculus, and their endocrine dependency, Journal of Chemical Ecology 1986 12:1. 12 (1986) 277–296. https://doi.org/10.1007/BF01045611.
- [51] H.A. Soini, D. Wiesler, S. Koyama, C. Féron, C. Baudoin, M. v. Novotny, Comparison of urinary scents of two related mouse species, mus spicilegus and mus domesticus, Journal of Chemical Ecology. 35 (2009) 580–589. https://doi.org/10.1007/S10886-009-9628-2/FIGURES/3.
- [52] B. Jemiolo, T.M. Xie, F. Andreolini, A.E.M. Baker, M. Novotny, The t complex of the mouse: Chemical characterization by urinary volatile profiles, Journal of Chemical Ecology 1991 17:2. 17 (1991) 353–367. https://doi.org/10.1007/BF00994338.

 [53] M.L. Schaefer, K. Yamazaki, K. Osada, D. Restrepo, G.K. Beauchamp, Olfactory Fingerprints for Major Histocompatibility Complex-Determined Body Odors II: Relationship among Odor Maps, Genetics, Odor Composition, and Behavior, (2002). [54] K. Yamazaki, G.K. Beauchamp, J. Bard, E.A. Boyse, Chemosensory Identity and the Y Chromosome, Behavior Genetics. 20 (1990) 990. [55] B.S. Goodrich, S. Gambale, P.R. Pennycuik, T.D. Redhead, Volatiles from feces of wild male house mice, Journal of Chemical Ecology 1990 16:7. 16 (1990) 2091–2106. https://doi.org/10.1007/BF01026922. [56] A. Korpi, J. Järnberg, A.L. Pasanen, Microbial volatile organic compounds, Critical Reviews in Toxicology. 39 (2009) 139–193. https://doi.org/10.1080/10408440802291497. [57] S. Zomer, S.J. Dixon, Y. Xu, S.P. Jensen, H. Wang, C. v. Lanyon, A.G. O'Donnell, A.S. Clare, L.M. Gosling, D.J. Penn, R.G. Brereton, Consensus multivariate methods in gas chromatography mass spectrometry and denaturing gradient gel electrophoresis : MHC-congenic and other strains of mice can be classified according to the profiles of volatiles and microflora in their scent-marks, Analyst. 134 (2008) 114–123. https://doi.org/10.1039/B807061J. [58] A. Fujita, T. Okuno, M. Oda, K. Kato, Urinary volatilome analysis in a mouse model of anxiety and depression, PLOS ONE. 15 (2020) e0229269. https://doi.org/10.1371/JOURNAL.PONE.0229269. [59] J. Kwak, A. Willse, K. Matsumura, M.C. Opiekun, W. Yi, G. Preti, K. Yamazaki, G.K. Beauchamp, Genetically-Based Olfactory Signatures Persist Despite Dietary Variation, PLOS ONE. 3 (2008) e3591. https://doi.org/10.1371/JOURNAL.PONE.0003591. [60] E. Arnáiz, D. Moreno, R. Quesada, Determination of Volatiles in Mouse Urine by Headspace Solid Phase Microextraction and Gas Chromatography-Mass Spectrometry, Analytical Letters . 47 (2014) 721–729. https://doi.org/https://doi.org/10.1080/00032719.2013.853182. [61] M. Woollam, M. Teli, P. Angarita-Rivera, S. Liu, A.P. Siegel, H. Yokota, M. Agarwal, Detection of Volatile Organic Compounds (VOCs) in Urine via Gas Chromatography-Mass Spectrometry QTOF to Differentiate Between Localized and Metastatic Models of Breast Cancer, Scientific Reports 2019 9:1. 9 (2019) 1–12. https://doi.org/10.1038/s41598-019-38920-0. [62] A. Sever, A. Abd, Y. Matana, J. Gopas, Y. Zeiri, Biomarkers for Detection and Monitoring of B16 Melanoma in Mouse Urine and Feces, Journal of Biomarkers. 2015 (2015) 9. [63] Y. Hanai, K. Shimono, H. Oka, Y. Baba, K. Yamazaki, G.K. Beauchamp, Analysis of volatile organic compounds released from human lung cancer cells and from the urine of tumor- bearing mice, Cancer Cell International. 12 (2012) 7. https://doi.org/10.1186/1475-2867-12-7. [64] O. Barash, N. Peled, U. Tisch, P.A. Bunn, F.R. Hirsch, H. Haick, Classification of lung cancer histology by gold nanoparticle sensors, Nanomedicine: Nanotechnology, Biology, and Medicine. 8 (2012) 580–589. https://doi.org/10.1016/j.nano.2011.10.001. [65] M.P. Davies, O. Barash, R. Jeries, N. Peled, M. Ilouze, R. Hyde, M.W. Marcus, J.K. Field, H. Haick, Unique volatolomic signatures of TP53 and KRAS in lung cells., British Journal of Cancer. 111 (2014) 1213–1221. https://doi.org/10.1038/bjc.2014.411. [66] Z. Jia, H. Zhang, C.N. Ong, A. Patra, Y. Lu, C.T. Lim, T. Venkatesan, Detection of Lung Cancer: Concomitant Volatile Organic Compounds and Metabolomic Profiling of Six Cancer Cell Lines

- of Different Histological Origins, ACS Omega. 3 (2018) 5131–5140.
- https://doi.org/10.1021/acsomega.7b02035.
- [67] M. Ligor, T. Ligor, A. Bajtarevic, C. Ager, M. Pienz, M. Klieber, H. Denz, M. Fiegl, W. Hilbe, W. Weiss, P. Lukas, H. Jamnig, M. Hackl, B. Buszewski, W. Miekisch, J. Schubert, A. Amann, Determination of volatile organic compounds in exhaled breath of patients with lung cancer using solid phase microextraction and gas chromatography mass spectrometry, Clinical Chemistry and Laboratory Medicine. 47 (2009) 550–560.
- https://doi.org/10.1515/CCLM.2009.133.
- [68] D.F. Altomare, A. Picciariello, M.T. Rotelli, M. de Fazio, A. Aresta, C.G. Zambonin, L. Vincenti, P. Trerotoli, N. de Vietro, Chemical signature of colorectal cancer: case–control study for profiling the breath print, BJS Open. 4 (2020) 1189–1199. https://doi.org/10.1002/bjs5.50354.
- [69] C.L. Silva, R. Perestrelo, P. Silva, H. Tomás, J.S. Câmara, Volatile metabolomic signature of human breast cancer cell lines, Scientific Reports. 7 (2017) 1–8. https://doi.org/10.1038/srep43969.
- [70] H. Tian, H. Wen, X. Yang, S. Li, J. Li, Exploring the effects of anthocyanins on volatile organic metabolites of alzheimer's disease model mice based on HS-GC-IMS and HS-SPME-GC–MS, Microchemical Journal. 162 (2021) 105848. https://doi.org/10.1016/J.MICROC.2020.105848.
- [71] M. v Novotny, D.- Jena, R.J. Beynon, J.L. Hurst, M.M. Bradford, J.E. Bruce, G.A. Anderson, J. Wen, R. Harkewicz, R.D. Smith, J.L. Hurst, R.J. Beynon, K. Laemmli, C. Lautenschlager, W.S. Leal, J. Clardy, A.M. Chen, M.L. Erickson, E. Mortz, T.N. Krogh, H. Vorum, A. Görg, J. Peng, S.P. Gygi, Q. Li, C.L. Schultz, M. Moini, M. Muzio, A.M. Chinnaiyan, F.C. Kischkel, K.O. Rourke, A. Shevchenko, J. Ni, C. Scaffidi, J.D. Bretz, M. Zhang, R. Gentz, M. Mann, P.H. Krammer, M.E. Peter, V.M. Dixit, N.C. van de Merbel, U.A. Th, M.P. Washburn, D.A. Wolters, M.P. Washburn, J.R. Yates, I.I.I.N. Biotechnol, J. Zhou, G. Robertson, X. He, S. Dufour, A.M. Hooper, J.A. Pickett, N.H. Keep, L.M. Field, B. Applications, E.S.P.B. V, U.D.N. Lipidique, R. Henry, L.D.C. Organique, F. Couderc, N. Demont, Pheromones, binding proteins and receptor responses in rodents., Biochem Soc Trans. 31 (2003) 117–122. https://doi.org/10.1042/BST0310117.
- [72] K. Matsumura, M. Opiekun, H. Oka, A. Vachani, S.M. Albelda, K. Yamazaki, G.K. Beauchamp, Urinary volatile compounds as biomarkers for lung cancer: A proof of principle study using odor signatures in mouse models of lung cancer, PLoS ONE. 5 (2010). https://doi.org/10.1371/journal.pone.0008819.
- [73] J. Kwak, M. Gallagher, M.H. Ozdener, C.J. Wysocki, B.R. Goldsmith, A. Isamah, A. Faranda, S.S. Fakharzadeh, M. Herlyn, A.T.C. Johnson, G. Preti, Volatile biomarkers from human melanoma cells, Journal of Chromatography B. 931 (2013) 90–96. https://doi.org/10.1016/J.JCHROMB.2013.05.007.
- [74] H. Shigeyama, T. Wang, M. Ichinose, T. Ansai, S.W. Lee, Identification of volatile metabolites in human saliva from patients with oral squamous cell carcinoma via zeolite-based thin-film microextraction coupled with GC–MS, Journal of Chromatography B. 1104 (2019) 49–58. https://doi.org/10.1016/J.JCHROMB.2018.11.002.
- [75] D. Liu, L. Ji, M. Li, D. Li, L. Guo, M. Nie, D. Wang, Y. Lv, Y. Bai, M. Liu, G. Wang, Y. Li, P. Yu, E. Li, C. Wang, Analysis of volatile organic compounds released from SW480 colorectal cancer cells and the blood of tumor-bearing mice, Translational Cancer Research. 8 (2019) 2736. https://doi.org/10.21037/TCR.2019.10.21.
- [76] F. Gouzerh, B. Buatois, M.R. Hervé, M. Mancini, A. Maraver, L. Dormont, F. Thomas, G. Ganem, Odours of cancerous mouse congeners: detection and attractiveness., Biology Open . (2022).
- [77] H. Zhang, J. Cao, L. Li, Y. Liu, H. Zhao, N. Li, B. Li, A. Zhang, H. Huang, S. Chen, M. Dong, L. Yu, J. Zhang, L. Chen, Identification of urine protein biomarkers with the potential for early detection of lung cancer, Scientific Reports 2015 5:1. 5 (2015) 1–13. https://doi.org/10.1038/srep11805.
- [78] A. Fujita, M. Ota, K. Kato, Urinary volatile metabolites of amygdala-kindled mice reveal novel biomarkers associated with temporal lobe epilepsy, Scientific Reports 2019 9:1. 9 (2019) 1–13. https://doi.org/10.1038/s41598-019-46373-8.
- [79] B. Jemiolo, F. Andreolini, D. Wiesler, M. Novotny, Variations in mouse (Mus musculus) urinary volatiles during different periods of pregnancy and lactation, Journal of Chemical Ecology 1987 13:9. 13 (1987) 1941–1956. https://doi.org/10.1007/BF01014677.
- [80] K. Miyashita, A.B. Robinson, Identification of compounds in mouse urine vapor by gas chromatography and mass spectrometry, Mechanisms of Ageing and Development. 13 (1980) 177–184. https://doi.org/10.1016/0047-6374(80)90060-3.
- [81] K. Osada, T. Tashiro, K. Mori, H. Izumi, The Identification of Attractive Volatiles in Aged Male Mouse Urine, Chemical Senses. 33 (2008) 815–823. https://doi.org/10.1093/CHEMSE/BJN045.
- [82] M. v. Novotny, H.A. Soini, S. Koyama, D. Wiesler, K.E. Bruce, D.J. Penn, Chemical identification of MHC-influenced volatile compounds in mouse urine. I: Quantitative proportions of major chemosignals, Journal of Chemical Ecology. 33 (2007) 417–434. https://doi.org/10.1007/s10886-006-9230-9.
- [83] S. Harvey, B. Jemiolo, M. Novotny, Pattern of volatile compounds in dominant and subordinate male mouse urine, Journal of Chemical Ecology 1989 15:7. 15 (1989) 2061–2072. https://doi.org/10.1007/BF01207438.
- [84] M. Novotny, F.J. Schwende, D. Wiesler, J.W. Jorgenson, M. Carmack, Identification of a testosterone-dependent unique volatile constituent of male mouse urine: 7-exo-ethyl-5- methyl-6,8-dioxabicyclo[3.2.1]-3-octene, Experientia 1984 40:2. 40 (1984) 217–219. https://doi.org/10.1007/BF01963608.
- [85] J.X. Zhang, X.P. Rao, L. Sun, C.H. Zhao, X.W. Qin, Putative Chemical Signals about Sex, Individuality, and Genetic Background in the Preputial Gland and Urine of the House Mouse (Mus musculus), Chemical Senses. 32 (2007) 293–303. https://doi.org/10.1093/CHEMSE/BJL058.
- [86] M.N. Kayali-Sayadi, J.M. Bautista, L.M. Polo-Díez, I. Salazar, Identification of pheromones in mouse urine by head-space solid phase microextraction followed by gas chromatography– mass spectrometry, Journal of Chromatography B. 796 (2003) 55–62. https://doi.org/10.1016/J.JCHROMB.2003.08.001.
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ACKNOWLEDGEMENTS

- We are grateful to Pascal Boutinaud, Nathalie Barougier, Camille Rosa, Anais Pichevin, Mélanie
- Moreno and Gwenaelle Vigo for support in the animal husbandry and lab work.

FUNDING

- FG was supported by the doctoral fellowship of the University of Montpellier. FT was supported by
- 727 the MAVA Foundation, and by the ANR TRANSCAN (ANR-18-CE35-0009).

COMPETING INTERESTS

The authors declare no competing financial interests.

AUTHOR CONTRIBUTIONS

- FG, GG, FT, LD developed the concept, FG realized the experiments, FG and GG performed the
- analytical measurements and analyzed the date with MH. BB supervised the chemical ecology aspects.
- MM and AM performed euthanasia of EGFR mice and verification of their cancer status. FG and GG
- prepared and edited the manuscript and all authors have reviewed the manuscript.
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FIGURE

 Fig. 1 : A schematic of the experimental protocol. Each bag contained a mixture of bedding obtained from 3 to 4 mice, at different experimental conditions (T0: before treatment, T2: after 2-weeks doxycycline diet; and T12: after 12 weeks doxycycline diet). Three different types of tests were performed: - 1h VOCs sampling using two different SPME fibers exposed simultaneously to the head space and involving T0 non-cancerous (NC) soiled bedding (testing "Fiber effect"); - Two fibers exposed simultaneously, one for 1-h the other for 24h and involving soiled bedding of NC and CC (cancerous) soiled bedding sampled at T0, T2 and T12 (testing "Effect of sampling time"); and – A single fiber exposed for 1h and involving 3 replicates of the same source of NC and CC soiled bedding 749 at T0, T2 and T12 ("Cancer influence").

Fig. 2 : Pic surface areas (a), and relative proportions (b) of compounds identified in mice soiled

bedding samples after 1h versus 24h SPME extractions.

Constrained RDA Dim 1:45, 21%

Fig. 3 : Score plots of the redundancy analysis (RDA) comparing variation of 17 VOCs identified in

the 1h extraction profiles (T1) versus the 24h extraction profiles (T24) of non-cancerous (NC) and

cancerous (CC) mice soiled bedding.

 Fig. 4 : Score plots of the redundancy analyses (RDAs) comparing variation of 17 VOCs identified following 1h SPME extraction of non-cancerous soiled bedding (A) and cancerous soiled bedding (B) at three experimental condition (T0: before treatment, T2: after 2 weeks doxycycline diet; and T12: after 12 weeks doxycycline diet). Correlation circles represent the position of VOCs showing an absolute correlation coefficient > 0.8.

 Fig. 5 : Score plots of the redundancy analysis (RDA) addressing variation of 10 VOCs present in the profiles obtained after1h extraction of NC (non-cancerous) and CC (cancerous) soiled bedding at three experimental conditions; (T0: before treatment, T2: after 2 weeks doxycycline diet; and T12: after 12 weeks doxycycline diet). The correlation circle identifies VOCs showing an absolute 775 correlation coefficient > 0.8 .

- 778 **Table 1** : List of the 17 VOCs identified after simultaneous exposure of two SPME fibers to soiled 779 bedding of non-cancerous mice at T0. Only the components not present in the technical controls (clean
- 780 bedding) were retained in this list (Table S1).
- 781

783 **: attempt of identification when the exact configuration could not be determined.*

784 ^a retention time

- 785 b calculated retention indices
- 786 ^cMaximum number of occurences.
- 787 $\frac{d}{dx}$ relative proportion of the compounds (%, mean +/- SE)
- 788
- 789

790 **Table 2** : Results of the permutation F tests based on redundancy analyses linking VOCs composition to health status (NC non-cancerous mice *versus* CC

791 potentially cancerous mice).

793 **Table 3 :** Relative proportions (%, mean ± se) of 17 VOCs identified in mice soiled bedding following 1-h SPME sampling, with reference to health status 794 (non-cancerous, NC, cancerous, CC) at three experimental conditions (T0, T2 and T12). VOCs present both in the biological and the control samples are not 795 listed here.

797 **: attempt of identification; in some cases the configuration could not be determined.*

798 +*VOCs differing between CC and NC mice at T0*

	T ₀ _CC		T ₂ CC T ₁₂ CC	T _{2_NC}
T0_NC	0,622			< 0,001
T ₂ NC		0,4948		
T12_NC			< 0,001	0,2593
T ₂ CC	< 0,001			
T12 CC		< 0,001		

Table 4 : Post-hoc tests following RDA model 7.

SUPPLEMENTARY MATERIALS :

 Fig.S1 : Representation of a) Score plots of the redundancy analysis (RDA) comparing variation of 806 relative proportion of 17 volatile organic components sampled with fibers F1 and F2 at seven occasions;

807 b) A correlation circle identifying compounds showing an absolute correlation coefficient > 0.8 and c)

Variation of the relative proportion of the compounds pinpointed in the RDA analysis.

815 **Table S1 :** List of VOCs identified both in the biological samples and the controls. In blue compounds

816 found only in soiled bedding and retained for our analyses. In pink: the n-alkanes used as standards for

817 the identification of the compounds.

^a 818 *Calculated retention indices*

^b 819 *Retention time*

820

821 **Table S2** : VOCs identified following 24h versus 1h extraction of soiled bedding. Only the components

822 not present in the technical controls (clean bedding) are listed here.

825 a retention time

826 b calculated retention indices

827 ^c frequency of occurrence of the compounds.

828 d relative proportion of the compounds (%, mean +/- SE)

829 • relative quantity of the compounds (mean $+/-$ SE)

830