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Licenciada em Bioquímica

**Theoretical and experimental  
approaches towards the non-invasive  
and selective detection of microbial  
pathogens**

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# Abstract

Non-invasive diagnostics of microbial infections based on the volatome is an area of increasing interest. This work set the ground for the development of an artificial nose for the diagnosis of bacterial infections. In a first part, the aim was to find a group of volatile organic compounds (VOCs) able to distinguish between 8 clinically relevant pathogens. The second part aimed to engineer the selectivity of VOC-responding materials (biogels) through the incorporation of VOC-specific peptides.

A systematic review and analysis of available literature data relating the detection of VOCs in human samples with the presence of specific pathogen infections was performed. Statistical classification methods were employed to make a metasearch for potential pathogen VOC biomarkers using those data. A minimal set of VOCs that allows the distinction between *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Aspergillus fumigatus*, *Escherichia coli*, *Helicobacter pylori*, *Proteus mirabilis*, *Klebsiella pneumoniae* and *Mycobacterium tuberculosis* was suggested.

A comparison between the set of VOCs found by the analysis and biomarkers previously reported for the same pathogens was made. As a result, a set of potential biomarkers for pathogen infection is suggested: indole for *E.coli*; 2-pentylfuran for *A.fumigatus*; isobutene for *H.pylori*; cymol for *M.tuberculosis*; hydrogen cyanide and methyl thiocyanate for *P.aeruginosa*; and 3-methylbutanoic acid for *S.aureus*.

The feasibility of engineering biogels VOC-selectivity was assessed by incorporating in the materials a benzene-sensitive peptide previously reported (P1) and two modified versions containing norleucine (P2) or biphenylalanine (P3) at the C-terminal. The optical response of the as-produced materials to several VOCs was tested on an in-house developed electronic-nose (e-nose). The biogels without any peptide responded more sharply to benzene and acetone. The addition of P1 amplifies the response to benzene and toluene. The addition of P2 and P3 amplified the response signal to both acetone and benzene.

**Keywords:** volatile organic compounds; biomarkers; pathogen infections; liquid crystal; ionic liquid; electronic-nose.



# Resumo

O diagnóstico não invasivo de infecções microbianas baseado no volatoma é uma área de crescente interesse. Este trabalho definiu o caminho para o desenvolvimento de um nariz artificial para o diagnóstico de infecções bacterianas. Numa primeira parte, o objetivo era encontrar um grupo de compostos orgânicos voláteis (VOCs) capaz de distinguir entre 8 patógenos clinicamente relevantes. A segunda parte teve como objetivo desenvolver a seletividade de materiais que respondem à presença de VOCs (biogéis) através da incorporação de três péptidos diferentes específicos para VOCs.

Realizou-se uma revisão sistemática e análise dos resultados disponíveis na literatura relativos à detecção de VOCs em amostras humanas em casos de infecções causadas por agentes patogénicos específicos. Utilizaram-se métodos de classificação estatística para realizar uma metapesquisa para identificar potenciais VOC biomarcadores de patógenos, usando esses dados. Foi sugerido um conjunto mínimo de VOCs que permite distinguir entre *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Aspergillus fumigatus*, *Escherichia coli*, *Helicobacter pylori*, *Proteus mirabilis*, *Klebsiella pneumoniae* e *Mycobacterium tuberculosis*.

Realizou-se depois uma comparação entre o conjunto de biomarcadores voláteis encontrados na nossa análise e os biomarcadores reportados anteriormente, para os mesmos patógenos. Como resultado, foi sugerido um conjunto de potenciais biomarcadores para infecções patogénicas: indole para *E.coli*; 2-pentilfurano para *A.fumigatus*; isobuteno para *H.pylori*, cimenos para *M.tuberculosis*; cianeto de hidrogénio e metil-tiocianato para *P.aeruginosa*; e ácido 3-metilbutanóico para *S.aureus*.

A viabilidade de desenvolver a seletividade para VOCs em biogéis foi avaliada pela incorporação de um péptido, anteriormente reportado, sensível ao benzeno (P1) e duas versões modificadas do mesmo, contendo norleucina (P2) ou bifenilalanina (P3) no C-terminal, nos materiais. A resposta ótica dos materiais produzidos a vários VOCs foi testada no nariz eletrónico (e-nose) desenvolvido *in-house*. Os biogéis sem qualquer péptido responderam de forma mais acentuada ao benzeno e acetona. A adição de P1 amplificou a resposta para o benzeno e o tolueno. A adição de P2 e P3, amplificou o sinal de resposta tanto para acetona como para o benzeno.

**Termos-chave:** compostos orgânicos voláteis; biomarcadores; infecções por agentes patogénicos; cristal líquido; líquido iónico; nariz eletrónico.



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# List of Abbreviations

[BMIM][DCA]- 1-Ethyl-3-methylimidazolium dicyanamide

5CB- 4- Cyano- 4'- pentylbiphenyl

AF- *Aspergillus fumigatus*

AR- Allergic reactions

BI- Bone infections

BLI- Blood infections

CA- *Candida albicans*

CAP- Community-acquired pneumonia

CAR- Carboxen

CD- *Clostridium difficile*

CF- Cystic fibrosis

CJ- *Campylobacter jejuni*

CO<sub>2</sub>- Carbon dioxide

DVB- Divinylbenzene

EC- *Escherichia coli*

EF- *Enterococcus faecalis*

EI-MS- Electron ionization- Mass Spectrometry

e-nose- Eletronic-nose

ESI- Electrospray ionization

FID- Flame ionization detector

FITC- Fluorescein isothiocyanate

GC- Gas chromatography

GC-FID- Gas chromatography coupled with flame ionization detector

GC-MS- Gas chromatography Mass Spectrometry

GC-SAW- Gas chromatography surface acoustic wave

GD- *Giardia duodenalis*

GI- Gastric infections

GTI- Gastrointestinal infections

H<sub>2</sub>O - Water

H<sub>3</sub>O<sup>+</sup> - Hydronium ion

HI- *Haemophilus influenzae*

HP- *Helicobacter pylori*

HS-SPME- Headspace solid-phase microextraction

ICU- Intensive care unit

IL- Ionic liquid

IMR-MS- Ion molecule reaction Mass Spectrometry

IMS- Ion mobility spectrometry

KI- Kidney infections

KP- *Klebsiella pneumoniae*

LC- Liquid crystal

LDR- Light Dependent Resistor

LED- Light Emitting Diode

LP- *Legionella pneumophila*

m/z- Mass to charge ratio

MC- *Moraxella catarrhalis*

MCC- Multi- capillary column

MCC-IMS- Multi- capillary column coupled to ion mobility spectrometry

MM- *Morganella morganii*

MRSA- *Methicillin-resistant Staphylococcus aureus*

MS- Mass Spectrometry

MT- *Mycobacterium tuberculosis*



N<sub>2</sub> - Nitrogen

NM- *Neisseria meningitidis*

NO<sup>+</sup>- Nitrosonium ion

NTD- Needle Trap Device

O<sub>2</sub><sup>+</sup> - Oxygen ion

PA- *Pseudomonas aeruginosa*

PCA- Principal component analysis

PCR- Polymerase chain reaction

PDMS- Polydimethylsiloxane

PET- Polyethylene Terephthalate

PF- *Plasmodium falciparum*

PLSDA- Partial least squares Discriminant Analysis

PM- *Proteus mirabilis*

PN- Pneumonia

POM- Polarized Optical Microscopy

ppb- parts per billion

ppm- parts per million

ppt- parts per trillion

PTFE- Polytetrafluoroethylene

PTR-MS- Proton Transfer Reaction Mass Spectrometry

PTR-TOF-MS- Proton Transfer Reaction- time of flight- Mass Spectrometry

PV- *Proteus vulgaris*

SA- *Staphylococcus aureus*

SE- *Staphylococcus epidermidis*

SESI-MS- Secondary Electrospray Ionization Mass Spectrometry

SI- Sinusitis

SIFT-MS- Selected Ion Flow Tube Mass Spectrometry

SKI- Skin infections

SP- *Streptococcus pneumoniae*

SPME- Solid- Phase Microextraction

TB- Tuberculosis

TD-GC-MS- Thermal desorption Gas Chromatography Mass Spectrometry

UTI- Urinary tract infections

VAP- Ventilator- associated pneumonia

VOC- Volatile organic compound

WHO- World health organization

# 1. Introduction

## 1.1 Infectious diseases

Diseases caused by pathogenic microorganisms (bacteria, viruses, parasites or fungi) are called infectious diseases [1]. Worldwide, infectious diseases are the leading cause of death of children and one of the leading causes in adults [2], and an early diagnosis is essential to initiate appropriate antimicrobial therapy for efficient patient management [3].

Tuberculosis, pneumonia and malaria are examples of infectious diseases that affect the population worldwide. Tuberculosis (TB) is caused by the bacterium *Mycobacterium tuberculosis* that usually affect the lungs and occurs in every part of the world [4]. Although it is curable and preventable, an earlier diagnosis is essential. In 2014, 9.6 million people fell ill with TB and 1.5 million died from the disease. Also, globally, an estimated 480 000 people developed multidrug-resistant TB [5]. However, Africa carried the most severe burden, with 281 cases per 100 000 population in 2014 (compared with a global average of 133) [5].

Pneumonia is a form of acute respiratory infection that affects the lungs and it is the largest infectious cause of death in children worldwide, accounting for 15% of all deaths in children under 5 years old, in 2015 [6]. This pulmonary disease can be caused by a number of infectious agents, including viruses, bacteria or fungi. The most common are *Streptococcus pneumoniae* and *Haemophilus influenzae*. *Pseudomonas aeruginosa* is an uncommon cause of community-acquired pneumonia (CAP), but a common cause of hospital-acquired pneumonia [7]. Ventilator-associated pneumonia (VAP) is a common hospital-acquired infection occurring in the intensive care unit (ICU) and it is often caused by *Pseudomonas aeruginosa*. It is a complication of mechanical ventilation with an attributable mortality risk of 13% [8] even among patients receiving appropriate antimicrobial therapy. To date, the diagnosis is based on clinical criteria in combination with bacterial culture results.

Malaria is caused by infection with protozoan parasites belonging to the genus *Plasmodium* [9]. The parasites are transmitted to people through bites of infected female *Anopheles* mosquitoes, called "malaria vectors". There are five parasite species that cause malaria in humans [10], and two of these species – *P. falciparum* (most prevalent on the African continent) and *P. vivax* (predominates in many countries outside Africa) – pose the greatest threat [10]. About 3.2 billion people are at risk of malaria [10]. According to World Health Organization (WHO) estimates, released in December 2015, there were 214 million cases of malaria in 2015 and 438 000 deaths. Accurate diagnosis of malaria is important to provide adequate treatment and help prevent the emergence of resistant strains of malaria parasites [11]. However, diagnosis continues to present challenges. Currently, the majority of diagnoses rely on a combination of clinical presentation and the old approach of visualizing parasites on a stained blood film. There

remains a need for a simple, inexpensive, and reliable diagnostic test for malaria that can be performed *in situ* or in other primary healthcare settings in remote areas [12].

There are two key contributing factors for the highly negative prognosis in infectious diseases [13]. The first is the late diagnosis, usually performed using invasive and expensive procedures. The second is the lack of medical/laboratorial infrastructures in developing countries. Current methods for detecting microorganisms from clinical samples (culturing, polymerase chain reaction (PCR) and immunological methods) have some limitations regarding time, cost and complexity [14][15][16]. Therefore, there is an urgent need to develop fast, cheap, and accurate tests for the diagnosis of infectious diseases, so that it is possible to initiate early pathogen detection and subsequent specific treatment.

### **1.2 Volatile organic compounds (VOC) analysis in clinical samples towards diagnosis**

Humans emit, normally, a broad range of VOCs, which can be both odorous and non-odorous [17]. VOCs can be emitted from different secretions of the human body [18]. Emission varies with many factors such as age, diet, sex, physiological condition and possibly genetic background [19]. Therefore, body odours can be considered as individual ‘odour-fingerprints’ [20]. Pathological processes, such as infection and endogenous disorders, can influence those odour fingerprints by producing new VOCs or by changing the ratio of VOCs that are normally produced [21]. The correlation between VOCs and health is well known since the old clinical practices. For instance, Hippocrates recognized the diagnostic value of body odours and reported several disease-specific odours emanated from two different samples: urine and sputum [22].

There are some advantages associated with identifying specific combinations of VOCs (VOCs profiling) associated with human diseases [22]. The composition of clinical samples headspace gives valuable information about both endogenous and exogenous compounds. The first ones will reflect biochemical processes in the body while the second will be originated by exogenous microorganisms, offering new possibilities for non-invasive clinical diagnostics [23]. Headspace sampling can be used to collect VOCs from liquid (urine, blood) and solid (skin, stool) samples [24].

Clinical sample VOC analysis represents a convenient and simple alternative to the time consuming and expensive traditional methods used in clinical laboratory diagnosis. For that, analysis and identification of compounds that are found to be characteristic of a certain infection in clinical samples (infection VOC biomarkers) has been target of substantial research and is emerging as a promising diagnosis tool in modern analytical chemistry [19].

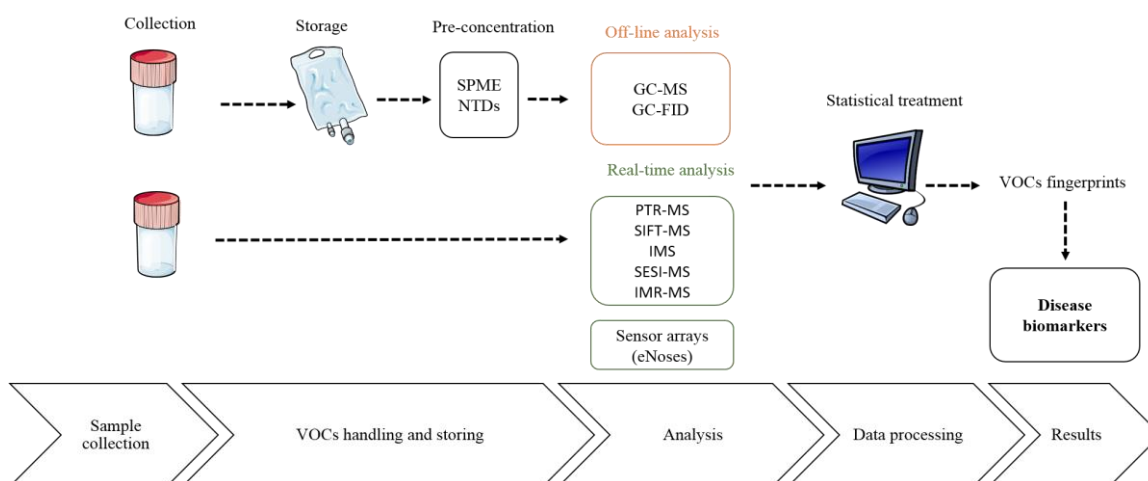
VOC biomarkers for infections are important clinical tools not only due to their disease-detecting potential [24] but also because there are other conceivable applications of VOCs in the field of infectious diseases [25]. Namely:

- Monitoring disease severity and control;
- Predicting prognosis of a disease;
- Evaluating treatment;
- Screening/predicting risk for different diseases in population studies.

VOC analysis from clinical samples has been developing into an attractive proposition because it is non-invasive and the available techniques to measure VOCs (such as Gas Chromatography Mass Spectrometry) are very sensitive (ppt<sub>v</sub>-ppb<sub>v</sub>) to detect compounds [26], and procedures (such as Selected Ion Flow Tube Mass Spectrometry and Ion Mobility Spectrometry) [26] [27] allow real-time measurement of compounds in the body.

### 1.3 VOCs study experimental outline

VOCs study can involve different steps, depending on the analytical method employed [28]. Usually, the main steps involved are: sample collection, VOCs handling and storing, analysis, processing of the data obtained and, finally, result output [29] (Figure 1.1). Depending on the chosen analytical method, VOCs may need to be captured, pre-concentrated and then stored [17].



**Figure 1.1-** Schematic representation of VOCs study experimental outline. SPME- Solid-Phase Microextraction; NTDs- needle trap devices; GC-MS- Gas-Chromatography Mass Spectroscopy; GC-FID- Gas Chromatography coupled to Flame Ionization Detector; PTR-MS- Proton Transfer Mass Spectroscopy; SIFT-MS- Selected Ion Flow Tube Mass Spectroscopy; IMS- Ion Mobility Spectrometry; SESI-MS- Secondary Electrospray Ionization Mass Spectrometry; IMR-MS- Ion Molecule Reaction Mass Spectrometry.

### 1.3.1 Sample collection

Sampling is one of the most relevant steps. Collection of blood and urine samples, for example, has standardized procedures [30]. Briefly, the sample is introduced in a sealed vial and the volatile components will diffuse into the gas phase until an equilibrium is reached and a sample is then taken from the headspace [31]. In the case of breath analysis, sampling is not so trivial. Breath collection is done by exhaling directly into sampling bags [19]. It can be done in two different ways: through a single breath or multiple breaths. Although single breath collection is less time-consuming, multiple breath analysis is more reproducible in terms of sample composition. So, for screening of potential VOCs associated with a given disease and determination of a specific set of biomarkers, multiple breath analysis is required. The risk of contamination with exogenous compounds from the oral cavity and the surrounding environment is always high and may compromise the analysis and the results [26][32]. These problems result in the variation of the number of compounds and their concentration, which may impair the analytical reproducibility and data reliability [32].

Regarding samples handling, there are some parameters that should be carefully considered to avoid wrong conclusions about the origin of the identified VOCs to be taken [29] [27]. Some of these parameters are sample storage and the interference of environmental VOCs.

### 1.3.2 Storage

When real-time analysis is not possible, samples need to be stored. Storage should be at very low temperature to reduce VOCs loss, and the samples should be stored as soon after being taken as possible. In the case of liquid or solid samples, they should immediately be placed in an appropriate container and frozen to -80°C or lower [30]. The container should be clean, produce no VOCs and should not change its characteristics with temperature variation and storage.

Breath sampling can be performed directly or indirectly according to the most suitable analysis to be performed. Direct sampling is preferable because there is no need to store for later analysis, so the decomposition of samples or loss of compounds by diffusion is avoided [19]. However, when direct analysis is not possible, the storage is an important factor to consider [26]. There are several ways to store breath and samples headspace [26]. The most typical examples are:

- Tedlar® bags (PTFE-polytetrafluoroethylene);
- Nalophan bags (PET-polyethylene terephthalate);
- Glass vials (for SPME);
- Thermal desorption tubes (different adsorbents, used in TD-GC-MS).

Currently, Tedlar® bags are the most common materials for VOC storage. However, Nalophan bags are also popular due to its low price, inertness, and relatively good durability [33].

### 1.3.3 Pre-concentration

During these procedures, interfering compounds could affect the analytical results. To minimize the interference, sometimes an intermediate step between sampling and analysis is required to increase the concentration of the target analytes over the interfering compounds [34]. There are several pre-concentration techniques available [35], such as Solid-Phase Microextraction (SPME) and Needle Trap Device (NTD) [36],[37].

Among the VOC sample pre-concentration methods, SPME is the most used one [24][38][39][40][41]. This technique involves the use of a fiber coated with an extracting phase which can extract different kinds of analytes, depending on the chosen fiber. The quantity of extracted analyte is proportional to its concentration in the sample. Its headspace variant (HS-SPME), in which analytes belonging to solid or liquid samples are extracted from the headspace, has gained major importance regarding VOC sampling [41].

The Needle Trap Device is an emergent alternative, consisting on a syringe that allows the combination of both the sampling and the pre-concentration steps in a single device [42]. This device is composed by a needle containing a sorbent material packed inside. The sorbent constitution is variable and includes Carboxen (CAR), Divinylbenzene (DVB), Polydimethylsiloxane (PDMS) [43]. In this method, the sample can be actively drawn in and out by diffusion, gas-tight syringe or automated devices, such as vacuum pumps [44]. Unlike SPME, NTD is an exhaustive methodology, allowing an increase in the concentration of several compounds by using more sample volume [38]. Moreover, sample storage, prior to analysis, is also possible with NTD and has been shown to deliver reproducible results for several days of storage, depending on the target analytes [45].

### 1.3.4 Analysis methods

To identify the different substances within a clinical sample, such as breath or headspace of liquid or solid samples, analytical methods are needed. Since the first reports about exhaled breath composition [46], several methodological improvements and alternatives have been implemented in VOC analysis. Nowadays, VOC analysis is no longer limited to the off-line laboratory approach, as there are many real-time methods available [47]. Table 1.1 summarizes the features of the most commonly used analytical methods for the characterization of gaseous samples towards VOC identification. The real-time analysis alternatives include analytical methods such as proton transfer reaction mass spectrometry (PTR-MS) and its variations

(proton transfer reaction-time-of flight-mass spectrometry (PTR-TOF-MS)), IMS and IMS coupled with multi-capillary columns (MCC-IMS), selected ion flow tube mass spectrometry (SIFT-MS) and secondary electrospray-ionization mass spectrometry (SESI-MS). All these real-time options reduce several experimental steps related with sampling, storage and pre-concentration of the samples, allowing a faster analysis and reducing the loss of information during these steps. More recently electronic noses (e-noses) [48] have been developed and applied to breath analysis with promising results [49]. Many e-nose approaches rely on pattern recognition and perform a qualitative characterization of volatiles unlike the analytical devices that give absolute quantification of volatiles. However, whether the characterization is quantitative or qualitative, the target VOCs always have to be identified by expensive comprehensive methodologies, usually involving mass spectrometry detection [29][38]. Therefore, the search for a reliable tool for VOC analysis assessment is still in progress [32].

#### **1.3.4.1 Off-line analysis methods**

Gas chromatography (GC) was the analytical method used in the initial studies in VOC analysis [46] and until today it is the gold standard method when coupled to mass spectrometry (MS). Most exhaled breath VOCs reported so far have been identified and quantified using MS-based methods [38].

##### **1.3.4.1.1 GC-MS (Gas-chromatography mass spectrometry)**

GC-MS [20] [26] [32] allows the analysis of compounds in the concentration range from ppb to ppt (Table 1.1). In GC-MS, analysis occurs when volatilized samples are separated in a chromatographic column based on parameters, such as the polarity of the GC column. This system ionizes the target ions, separates them by mass to charge ( $m/z$ ) ratio and then uses the fragmentation patterns to quantify the amount of each VOC present in the analyzed sample [32].

##### **1.3.4.1.2 GC-FID (Gas-chromatography coupled to flame ionization detector)**

In GC-FID [50], VOCs are burned in the FID, producing ions and electrons that can conduct the electric current and this information is used for detection and eventually quantification. GC-FID usually exhibits high sensitivity, large linear response range, and low noise. The FID detector is mass sensitive and its response is not altered significantly by changes in mobile-phase flow rate [32] [38].



### 1.3.4.2 Real-time analysis

Real-time analysis [13] obtains immediate results and does not require collection and storage of samples, eliminating a major source of experimental errors [26]. It has some advantages when compared with off-line analysis. However, real-time analysis also has some disadvantages, such as the expensive maintenance of the equipment used, the high cost of data acquisition and the fact that detection limits cannot be improved by pre-concentrating the samples. Therefore, vestigial VOCs will not be detected by this approach [26] [32].

#### 1.3.4.2.1 PTR (Proton Transfer Reaction Mass Spectrometry)

PTR-MS [26][38][51][52] application in VOC biomarkers research is increasing, mainly because it can deliver results in a real-time analysis, with high sensitivities for VOCs detection (Table 1.1) and quantification (up to the ppt<sub>v</sub> range) [51]. In this analysis H<sub>3</sub>O<sup>+</sup> ions are used for proton-transfer reactions with many common VOCs, having almost no reaction with the abundant atmospheric gases (N<sub>2</sub>, CO<sub>2</sub> and H<sub>2</sub>O) [51][52]. However, PTR-MS has some limitations. This methodology does not allow the identification of compounds with the same molecular weight, because the detection relies on the atomic mass of compounds and the resolution of MS instruments is limited [24]. Therefore, a time of flight mass spectrometer can be linked to the PTR (PTR-TOF-MS) [53][54] to overcome this issue. In this technique, the ions are accelerated to a regular energy by an electric field. Then, the ions travel a defined distance without acceleration and the m/z will determine the time of flight of the compound. This methodological improvement makes possible the separation between distinct chemical compounds with the same molecular weight [38] [53]. However, as mentioned above, since pre-concentration is not possible, trace VOCs can hardly be detected using this approach, and this procedure is a much more expensive technique than GC-MS [53].

#### 1.3.4.2.2 SIFT (Selected Ion Flow Tube Mass Spectrometry)

SIFT-MS [26][38][55] is a technique that allows the measurement of trace concentrations of VOCs in humid air, including breath samples. In a general way, VOCs are collected into the flow tube and ionized with precursor ions (usually H<sub>3</sub>O<sup>+</sup>, NO<sup>+</sup>, or O<sub>2</sub><sup>+</sup>), forming the product ions, which are then quantified by MS [56]. This technique, just as PTR-MS, has some disadvantages. Due to the chemical ionization process, not all compounds are detectable (e.g. small hydrocarbons cannot be detected due to their low proton affinity) [26]. The issue of the proper identification of compounds is addressed in SIFT-MS by using different reactant ions (H<sub>3</sub>O<sup>+</sup>, NO<sup>+</sup>, or O<sub>2</sub><sup>+</sup>), which exhibit different ion-molecule reactions. Due to the different precursor ion generation, sensitivity of SIFT-MS detection and quantification (Table 1.1) is lower (ppb<sub>v</sub> range) than PTR-MS (ppt<sub>v</sub> range) [26][38].

#### **1.3.4.2.3 IMS (Ion Mobility Spectrometry)**

IMS [57] was initially developed for the high sensitive detection of illegal drugs and explosives, but then it was adapted to industrial and environmental applications, particularly for process control in food quality analysis and air quality control [58]. In the IMS analysis, ions are separated based on their mobility as they travel through a purified gas, in an electric field at the atmospheric pressure [58][59]. This can be achieved using commercially available IMS, without and with different gas chromatographic columns, as MCC (multi-capillary column) /IMS [60],[61]. The sensitivities that can be accomplished with IMS have made it suitable for breath analysis (ppbv - pptv range) [38]. There are already some successful examples of some IMS strategies applied in the diagnosis of pulmonary diseases (lung cancer, lung infections and asthma) as well as other bacterial infections [60]–[63].

#### **1.3.4.2.4 SESI-MS (Secondary Electrospray Ionization Mass Spectrometry)**

SESI [64] ionization occurs by proton transfer reactions between the electrospray solution and the volatile analyte, and is therefore suitable for the analysis of hetero-organic molecules, just as in traditional electrospray ionization (ESI). However, unlike the standard procedure, the proton transfer process of SESI occurs in the vapor phase rather than in solution [64]. The distinctive advantage that SESI provides over other ionization methods is that it is possible to fragment specific peaks (provided the appropriate type of mass spectrometer has been applied for SESI), which is an important tool for compound identification. SESI-MS has a sensitive detection limit (pptv range) [65] and it has been applied to the detection of explosive gaseous samples, human breath vapor, as well as in the identification of clinically relevant pathogens [66][67].

#### **1.3.4.2.5 IMR (Ion Molecule Reaction Mass Spectrometry)**

Electron impact ionization used in conventional mass spectrometry (EI-MS) leads to dissociative ionization of neutral gaseous compounds, thus creating complex fragmentation patterns. This fact limits the identification and quantification of gas mixtures containing different compounds of the same chemical group [68]. To overcome this limitation the ionization of small molecules via ion-molecule reactions (IMR), can be applied, which allows the reduction of fragmentation caused by high energy electron impact ionization. The IMR-MS technique was initially used to measure absolute gas concentrations of cars emissions, caloric plants (fermentation and catalytic processes) and to medical applications [69]. Nowadays, it is also used to the analysis of microbial headspace VOC composition for bacterial species differentiation [70]. This method uses soft chemical ionization for sample molecule ionization and displays no or only minimal fragmentation. The IMR sensitivity varies (ppm<sub>v</sub>-ppbv range) depending on the components measured, system setup and settings. Also, it has the capability of measuring compounds within milliseconds [71]. In this technique, positively charged atomic

ions interact with neutral sample gas molecules [70]. The two-body collision processes result in the formation of product ions whenever the ionization potential of the sample molecule is lower than the potential energy of the incoming primary ion. Differences in ionization potential between primary and product ions may result in a bond rupture and hence a lower molecular weight fragment ion. However, fragmentation is typically avoided due to the soft ionization process [69].

**Table 1.1-** Comparison of the mode of operation, sensibility, advantages and disadvantages of the different analytical methods.

Analytical method	Mode of operation	Sensibility	Advantages	Disadvantages	Refs
<b>GC-MS</b>	Off-line	ppt <sub>v</sub> -ppb <sub>v</sub>	Reproducible Identification of unknown VOCs and profiling possible	Pre-concentration needed; Slow; Quantification requires known compounds; Real-time measurements not possible; Expensive; Not suitable for clinical use	[26][38]
<b>PTR-MS</b>	Real-time	ppt <sub>v</sub>	No pre-concentration needed Potential for on-line testing	VOC chemical identification and complete profiling not possible	[38][26] [51][52]
<b>SIFT-MS</b>	Real-time	ppb <sub>v</sub>	No pre-concentration needed; measures in real-time; fast; Potential for on-line testing; Measures in headspace possible	VOC chemical identification and profiling not possible	[26][38][55][56]
<b>IMS</b>	Real-time	ppb <sub>v</sub> -ppt <sub>v</sub>	No pre-concentration needed; Low cost; Suitable for clinical use	Identification of unknown compounds is not possible	[38] [57]–[59]
<b>SESI-MS</b>	Real-time	ppt <sub>v</sub>	No pre-concentration needed; Fast	Complex VOC mixtures can cause unreliable results	[64][65]

Analytical method	Mode of operation	Sensibility	Advantages	Disadvantages	Refs
IMR-MS	Real-time	ppm <sub>v</sub> -ppb <sub>v</sub>	No pre-concentration needed; Fast	Sometimes formation of secondary ions that may have the same weight as the primary ions	[68]–[70]

#### 1.4 Data processing

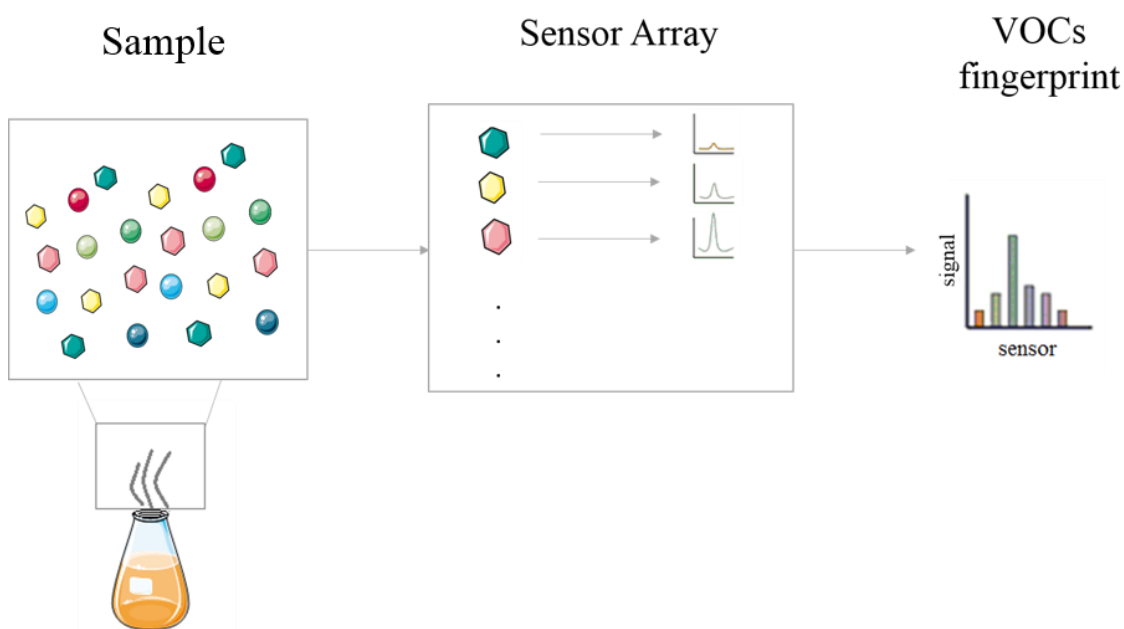
The statistical data treatment that follows the analysis step can be particularly inconvenient [26]. Although there is a full range of tools available to handle data complexity, until now there is no agreement regarding the selection and usage of those tools to discover volatile biomarkers that work with acceptable sensitivity and specificity for clinical applications [38]. In the majority of cases there are complex relationships between the group of compounds found in a clinical sample. For these reasons, volatiles identification and profiling using bioinformatics is a promising approach [72]. Specially, when adopting a strategy of identifying patterns instead of individual VOCs a more elaborate method of data analysis is required, such as Principal Component Analysis (PCA) [73] and Partial Least Squares Discriminant Analysis (PLSDA) [27], which allow a reduction of the dimensionality of data. Di Natale et al [74], for instance, found a set of putative biomarkers for lung cancer, in a group of 42 patients, using e-nose and GC-MS methodologies and recurring to PLSDA to analyze the data. On the other hand, Montuschi *et al.* [75] used PCA analysis to obtain an asthma VOCs pattern based on 27 patients. Data can then be plotted and a visualization of similarities and differences between data sets is possible. Also, it is possible to identify individual components, instead of patterns, that will be responsible for the differences observed between data sets and, finally, identify if some of those compounds are biomarkers [72].

#### 1.5 Electronic noses (e-noses)

In order to measure different VOCs, many applications have combined various sensors and materials into a single array, leading to the development of a device able to detect and distinguish odorous compounds- an e-nose [76]. Electronic noses follow an approach which closely resembles mammalian olfaction, by measuring the whole spectrum of VOCs without identification of the individual components [77]. Although individual VOCs cannot be identified, the output of e-noses represents a signature of the VOC pattern (fingerprint) (Figure 1.2), which can be analyzed by pattern recognition algorithms to discriminate VOC mixtures and potentially to detect diseases [19]. There are several formats for e-nose sensors, [77] [78], which are summarized in Table 1.2. Unlike GC-MS and other analytical techniques, e-noses do not contribute to the discovery of biomarkers that are specific for a disease.

Nonetheless, they can be used to compare samples to see if they have similar VOC profiles. Also, due to the low-cost and implementable nature of this technology, it has great potential for clinical use [79]. Pavlou *et al.* [80] successfully discriminated some bacterial cultures, associated with tuberculosis, such as *M.tuberculosis* and *M.avium*, by using the volatile patterns resulting from an electronic nose based on a 14 sensor conducting- polymer sensor. Wang *et al.* [81] used a colorimetric sensor to analyze 14 breath biomarkers, such as ammonia, acetone and ethane, in a breath analysis study.

Currently, e-nose research is focusing on finding materials with high sensitivity and good selectivity for VOCs detection to improve the sensitivity and specific discrimination between the pathogens producing them [82][83].



**Figure 1.2** – Schematic representation of volatile compounds recognition by an electronic nose device.

**Table 1.2-** Summary of the different e-noses formats and their mode of operation [78][84].

Sensor Format	Mode of operation
<b>Conducting-polymer sensor</b>	VOCs interact and attach to the polymer surface changing the resistance which results in changes in the signal.
<b>Metal oxide sensor</b>	Oxide materials contain chemically adsorbed oxygen species, which can interact with the VOCs, altering the conductivity of the oxide.
<b>Metal oxide silicon field-effect sensor</b>	Related to metal oxide sensors but the output signal is originated from a change in potential when the VOCs react at a catalytic surface.

Sensor Format	Mode of operation
<b>Piezoelectric crystal</b>	Adsorption of VOCs onto the membrane results on a change in the magnitude of the resonance frequency that is related to the mass of the volatile analyte.
<b>Surface acoustic-waves device</b>	Based on waves emitted along the surface of a crystal by the electric field of surface-deposited aluminium electrodes.
<b>Optical sensor</b>	Based on a light source that interacts with the volatile analyte. The signal is measured in absorbance, fluorescence, reflectance or chemiluminescence.
<b>Electrochemical sensor</b>	Responses are dependent on the electrochemical characteristics of the VOCs that are oxidized or reduced at the working electrode and at the counter electrode. Generated voltage of the reactions between the electrodes is measured.

### 1.6 Challenges and future directions

Although VOCs profiling is a potential clinical tool, the technique is still not part of routine analysis. Before it is implemented in clinical analysis there are some steps that need to be validated. The first one is an extensive validation of the current available VOCs profiles. Also, further development of the sample-collection devices and the sampling mechanisms is required in order to facilitate taking reliable, reproducible samples [85]. Numerous factors can influence emitted VOCs [86]. More research is required to further identify microbial specific VOCs and this situation is aggravated by the fact that the specific VOCs produced by a given microbe in its natural environment can be different from what is observed *in vitro* due to the use of different growth medium, incubation conditions and possible presence of other microorganisms [40]. Also, the analytical method should be carefully studied. For instance, when direct analysis is not possible, the samples need to be stored, which can affect the original composition of the collected sample [19] [26]. The physiological meaning and biochemical origin of endogenous VOCs so far are not clear. Although more insight is needed this is not an easy study, because the origin of VOCs can be the result of widely different biochemical pathways [87]. Finally, further refinement of sampling techniques and the development of new tools that combine the strengths of the e-nose (cheap, time efficient) [19][77] IMS (real-time) [57][60]–[63] and GC-MS (sensitive, compound identification) [20] [26] [32] [46] will favour the introduction of VOCs analysis into clinical practice.

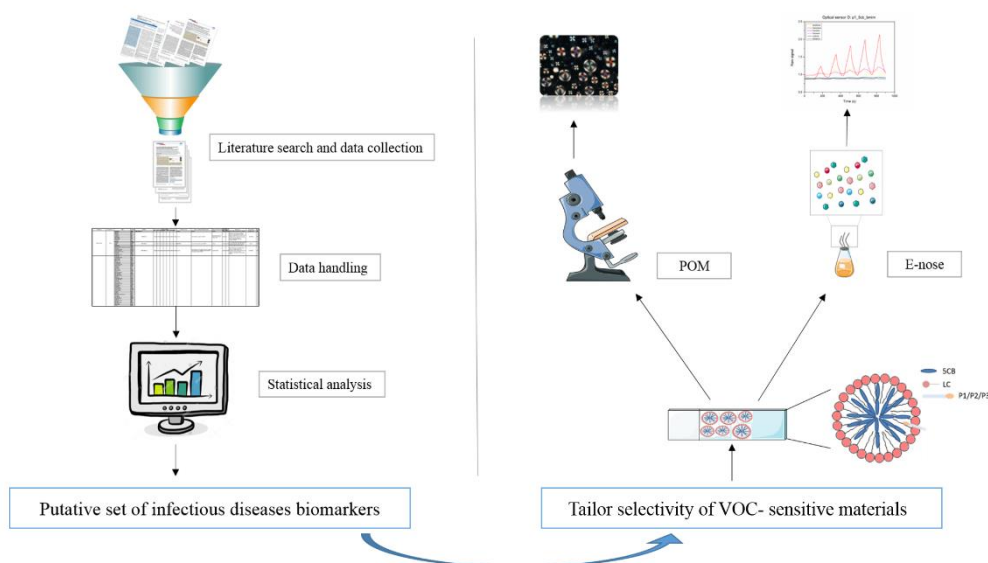
## 2. Aim of the work

Non-invasive diagnostics of microbial infections based on the volatome is an area of increasing interest. E-noses have emerged as a low-cost technology with great potential for clinical use. Also, LC based-devices have been reported as reliable, low-cost and high-sensitive gas sensors for biological and chemical sensing [88][89]. However, there is a need to find sensitive materials with good selectivity for VOCs detection to improve the specific discrimination between the pathogens.

In the first part of this work a search for infectious diseases VOC biomarkers was conducted (Figure 2.1). A literature search was performed and data of interest was collected and organized in a database. A statistical analysis was performed and machine learning algorithms were used to design and develop a model to distinguish pathogens based on the detected VOCs and to identify a possible set of VOC biomarkers for those pathogens.

The second part of this work consisted of a proof-of-concept study in which a proprietary e-nose and a gel-like VOC-sensitive material with optical properties were employed. The work aimed to explore a method for tailoring the selectivity of the material towards a particular set of VOCs (Figure 2.1). As a case-study, three different VOC-specific peptides were incorporated into the materials and the response of the modified material was characterized by Polarized optical microscopy (POM) and further tested on the e-nose.

The long-term goal of this study is to use both the classifier and the combination of responses of the VOC-sensitive biogels to the different solvents in aid to the identification and distinction of volatiles emitted by clinically relevant pathogens.



**Figure 2.1-** Schematic representation of the thesis project.





### 3. Looking for infectious disease VOC biomarkers

#### 3.1 Introduction

Infectious diseases are one of the major causes of death worldwide. The early detection and identification of the causative microorganisms allows a prompt initiation of appropriate antimicrobial therapy essential for an efficient patient management [3]. Simultaneously, the global spread of antimicrobial resistance to antibiotics is a predominant reason why infectious diseases continue to be target of attention [90]. The misuse of antibiotics is one of the factors that contributes to the selection of drug-resistant pathogens, increasing the need for a correct identification of the microorganisms responsible for infections [91]. The traditional methods for bacterial detection and identification rely on culture and colony counting methods [14], polymerase chain reaction (PCR) [14][16] and immunology-based methods [14][15]. However, these suffer from some major drawbacks. First, the majority can only be used for organisms that can be cultivated *in vitro*. Second, they are still time-consuming and require technical expertise and equipment [4][5][7].

Disease-related biomarkers are chemicals which presence/absence in the body differs according to the health status of an individual [24], indicating the presence or severity of a particular disease state. These biomarkers may have endogenous (produced within the body) or exogenous (introduced into an organism) origin.

Pathogenic microbes release unique combinations of metabolites in the body, which represent potential infectious disease biomarkers [13]. Analyses of volatile organic chemicals from different bodily fluids (blood, saliva, urine, faeces, milk, breath and skin) have the potential for bacterial identification and differentiation [92][24][93], since some pathogenic metabolites are VOCs, and therefore offer the possibility of fast diagnosis and disease monitoring, when compared to traditional methods. This approach has only begun to receive attention recently, mainly because VOCs are present in the body in low concentrations (ppt<sub>v</sub>-ppm<sub>v</sub>), making it essential to use analytical methods with high sensitivity ranges [23]. The advances in analytical techniques increased the potential for VOCs detection and GC-MS has become the gold standard instrument for headspace VOC analysis as it offers high sensitivity (ppt<sub>v</sub>-ppb<sub>v</sub>) and extensive compound libraries are available, making compound identification easier [19]. Before VOC analysis can be implemented in clinical diagnosis, possible volatile biomarkers must be known. Currently, the search for VOCs as disease biomarkers has been the focus of many studies that, in some cases achieved different results. Kunze *et al.* [61], for instance, identified 2-ethyl-1-hexanol, acetone, 2-phenylacetaldehyde, ammonia (dimer), dodecane, nonanal and ammonia in the headspace of a *P.aeruginosa* isolate obtained from a blood sample. On the other

hand, Savelev *et al.* [39] found 2-nonanone, 2,4-dimethyl-1-heptene, 1-heptene, isopentanol and limonene in a *P.aeruginosa* isolate obtained from a breath sample. The use of distinct samples (breath, blood, urine, skin, faeces), sampling methods and analytical techniques contributes to a variety of results, making difficult to have firm conclusions about the relevance of each VOC as infection biomarker.

In this chapter, we performed a systematic review of existing literature, and analysis of published results relating the detection of VOCs in human samples with the presence of specific pathogen infections. Hence, we suggest a minimal set of VOCs that allow the distinction between 8 clinically relevant pathogens - *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Aspergillus fumigatus*, *Escherichia coli*, *Helicobacter pylori*, *Proteus mirabilis*, *Klebsiella pneumoniae* and *Mycobacterium tuberculosis*. The set of volatile biomarkers found in our analysis was compared with the biomarkers found in other works, for the same pathogens. As a result, we identified a set of potential biomarkers for pathogen infection: indole for *E.coli*; 2-pentylfuran for *A.fumigatus*; isobutane for *H.pylori*; cymol for *M.tuberculosis*; hydrogen cyanide and methyl thiocyanate for *P.aeruginosa*; and 3-methylbutanoic acid for *S.aureus*.

## 3.2 Materials and methods

### 3.2.1 Literature search and data collection

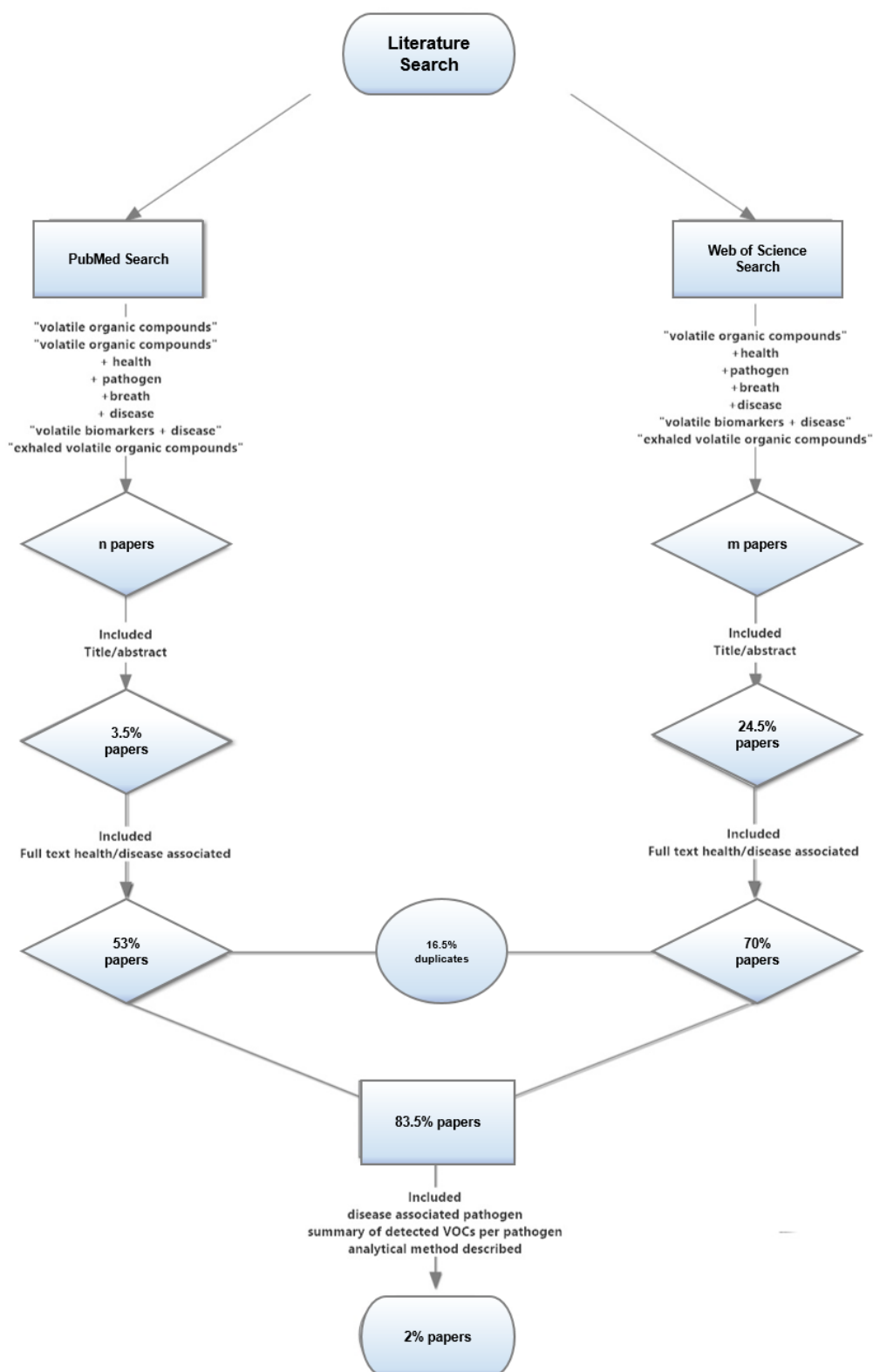
Data was collected from bibliography obtained through a systematic search in the PubMed and the Web of Science on-line libraries, performed in the period between October 12<sup>th</sup>, 2015 and February 10<sup>th</sup>, 2016. The search terms in PubMed were: “volatile organic compounds”; “volatile organic compounds + health/breath/pathogen/disease”, “volatile biomarkers + disease” and “exhaled volatile organic compounds”. For the Web of Science search, the terms were the same except for “volatile organic compounds”, that was not used alone, because the PubMed search with that same term included many articles that were not associated with health (pathogen).

The retrieved articles were selected for examination if the title and/or abstract suggested the investigation of microbial pathogens and the measurement of VOCs regarding human clinical perspective.

Further screening of the selected articles was performed according to the flowchart depicted in Figure 3.1. Relevant articles were selected for the study according to a set of inclusion criteria:

- i. The article’s subject should concern the human clinical research field (plants, soils and animals research fields were not included);
- ii. The article should indicate the name of the disease-associated pathogen (the disease name alone was not informative enough);

- iii. The article should report quantitative or qualitative information regarding the individual VOCs instead of reporting only VOC patterns;
- iv. The analytical method used to detect and quantify the VOCs should be described;
- v. The article should provide a summary of the detected VOCs per pathogen.



**Figure 3.1-** Strategy followed for the selection of papers.

### 3.2.2 Data handling

Data of interest was collected after reading the fulltext of the selected articles and organized in a database. A major table was compiled with information retrieved from the articles, organized in the following columns: pathogen identification (name and classification), bacterial strain (when applicable), VOCs associated with each pathogen and corresponding PubChem ID (for univocal VOC identification), the type of sample (saliva, blood, breath, skin, urine, faeces and milk) where each VOC was identified (when applicable), the analytical method used to detect it, culture conditions/growth medium, incubation time before analysis, detected VOC concentration range (value and unit) and the respective bibliographical reference (Appendix 2).

Since data collection and respective organization in the database was performed by a single subject, table filling errors might occur. To quantify those errors, data validation was performed by a second independent subject. 3 articles were chosen randomly from the set of 44, and the database was filled by the second subject with the information collected from the articles. It was found that in 100 VOCs present in those articles there were 3 table filling errors, resulting in an associated error of 3%.

A new database was created by a transformation on the structure of the previous described database to facilitate further data processing. Data was re-organized and a new parameter was added: the number of experiments. Some articles included results from more than one experimental condition: for example, results obtained with distinct growth media, with different analytical methods, or even with multiple bacterial strains. To account for these situations, for the same article, each dataset obtained in a specific experimental condition was considered as a distinct experiment. Hence, an article may describe several experiments and one pathogen may have a higher number of associated experiments than the corresponding number of papers. The database organization was one entry per experiment related to one pathogen, and the Boolean (true or false) indication of the detection of each of the VOC.

The number of hits was considered as the number of times that each VOC was detected in all of the experiments. In some cases data was normalized to percentage of the total number of experiments to facilitate data visualization and interpretation.

Cytoscape software was used as a visualization tool, to generate pathogen-VOC interaction graphs that allow an easy identification of the evolution of data processing during this work.

### 3.2.3 Statistical analysis

A new filtering criteria was applied to the full dataset (major table modified), in order to reduce the unbalance of data between pathogens. Therefore, a sub-dataset including only pathogens with more than 3 associated experiments was used. The result was 8 pathogens: 2 Gram<sup>+</sup> bacteria, 1 fungus and 5 Gram<sup>-</sup> bacteria.

The main goals of this study were to devise a model to distinguish pathogens based on the detected VOCs and to identify a putative set of VOC biomarkers for those pathogens. For that, machine learning methods based on statistical classification were used to design and develop the algorithms for pathogen classification from the transformed sub-data matrix with 8 pathogens, 269 VOCs and 174 experiments. Each line of the matrix is a features vector, which refers to an experiment where a pathogen was present and contains a binary vector reporting the identification (or not) of a VOC. Each VOC presence is considered a feature in the features vector.

A set of computing steps were executed in order to generate classifiers and estimate the classification error rate (detailed in Figure 3.2). This computational work was performed by Prof. Hugo Filipe Silveira Gamboa (Faculdade de Ciências e Tecnologia- Universidade Nova de Lisboa, Departamento de Física).

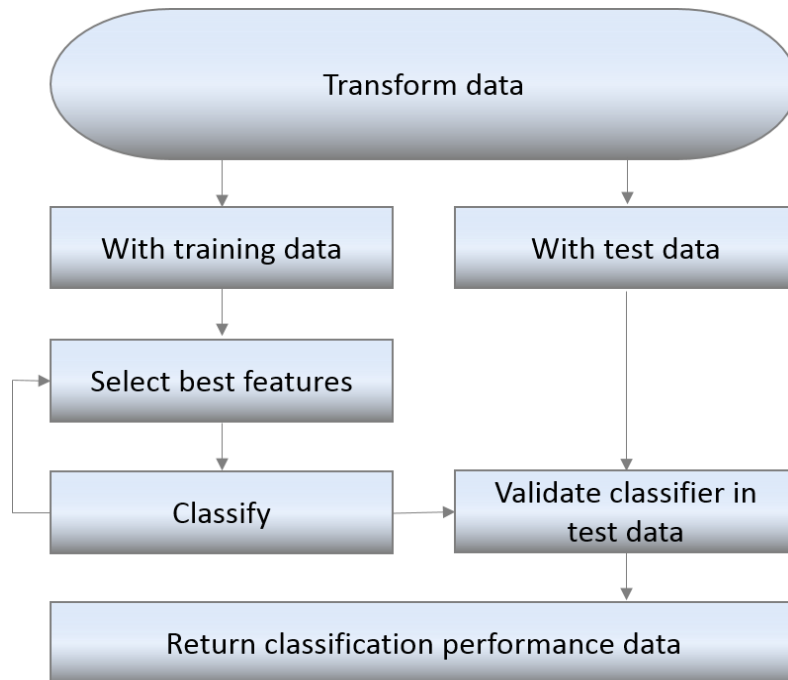
The first computational step consisted in transforming the database with the collected information on the papers to generate the binary vector of features. We separated the data into training and test datasets for validation purposes. Then, we executed a feature selection mechanism to identify a good subset of features that generated low classification error. The classifier was trained in the process to search for the best subset of features. The process ended with a validation step where we computed the final error and classifier behavior by computing the confusion matrix when a dataset not used in the training phase was used.

In the classification process we used statistical based classifiers to be able to separate the pathogens based on the binary VOC input data.

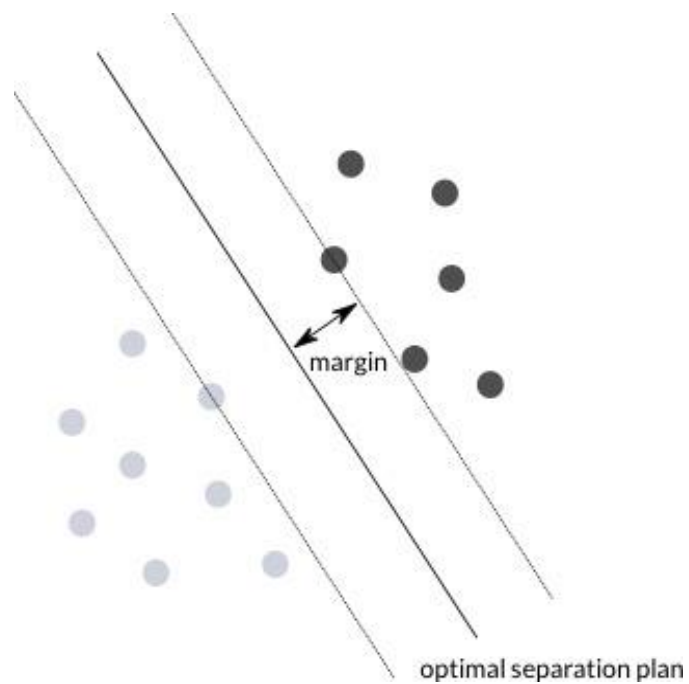
We tested several standard classifiers [94]: decision trees, naive Bayes classifier, nearest neighbour classifier and support vector machines (SVM). In our preliminar tests the results generated by the SVM always outperformed the other classifiers. We selected the SVM with linear kernels as the classifier to execute the feature selection process.

The support vector machine [95] classification method operates a transformation on the data projecting the data to a higher dimension space than the original data structure and applies an optimization technique to find an optimal separation plan in the new transformed space. In

figure 3.3 we show an example of a separation plan selection that maximizes the separation margin between the two classes.



**Figure 3.2-** Classification steps.



**Figure 3.3-** Support Vector Machine optimal separation plan and separation margin.

The base learning process of the SVM optimizes the margin distance by selecting a separation plan of a two class problem. This process is replicated to each pair of pathogens.

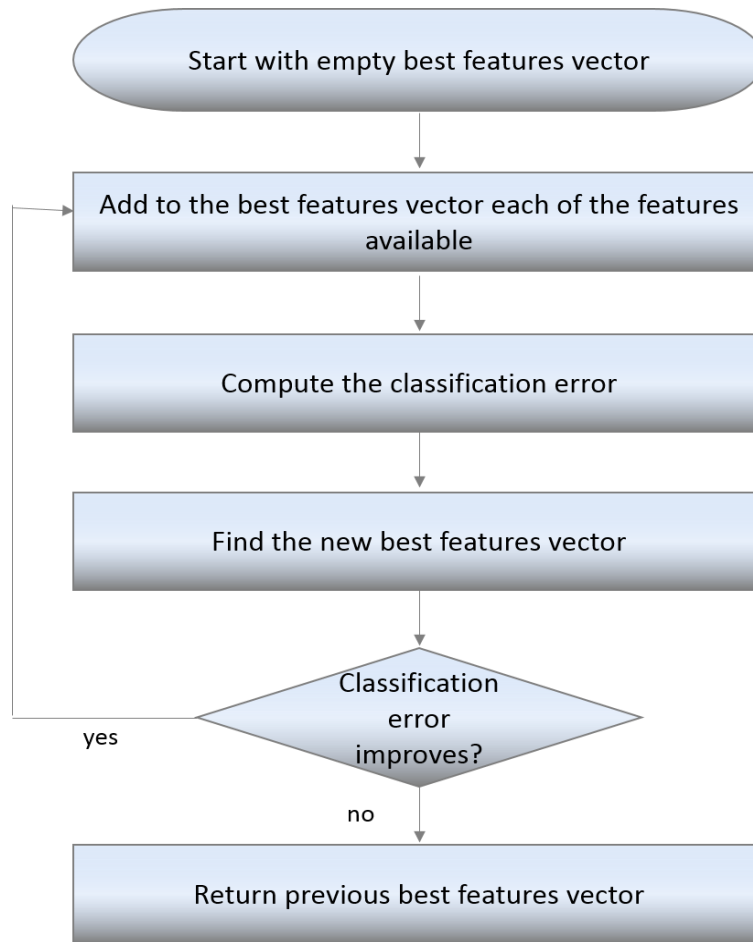
The classification task was performed in two modes [96].

- i. Multi- class: we defined a multi-class problem where we considered each pathogen as a separated class and a set of SVM were adjusted to each pathogen. This mode is also called identification mode, when we have several classes and we want to identify to which class the data belongs (multi-class problem). The question this classifier will address is: “Based on these VOC what is the most probable pathogen (from the set of selected pathogens)”.
- ii. Dual-class: for each pathogen we were interested in verifying if the sample VOCs corresponded to the pathogen or to any other one. This is also called verification, where we are interested in verifying if our assumption of the data belonging to a specific class is true. We are answering to this type of questions: “Does the new data belong to, e.g. *P. aeruginosa*?”

A selection of the best VOCs subset was executed by standard feature selection mechanisms implemented in both modes of classification.

Two non-optimal mechanisms are normally applied: the sequential forward feature selection and the sequential backward feature selection. In the first case we start with an empty vector of features, adding one feature at a time and growing the vector until the classification error stops decreasing. In the backward mechanism we start with the full feature vector and remove one feature at a time, reducing the dimension of the feature vector. We used the later one because it requires a lower computational complexity and the results tend to be similar [97]. The steps executed in the sequential forward selection algorithm are depicted in figure 3.4.





**Figure 3.4-** Sequential forward feature selection.

In the case of multi-class, we executed the feature selection for all the classes, returning a vector of the best features for separating the pathogens.

In the dual class problem, where we verified the possibility of a pathogen against all the other pathogens, we executed the feature selection for each case returning for each pathogen a set of features that better separated the pathogen from all the others.

All the results are reported based on a cross validation mechanism where we use a training dataset to find the best features and train the classifier, and a testing dataset where we report the classification error rate and the confusion matrix. We used the leave-one-out [98] cross-validation method, that removes only one pathogen example from the training data set and tests the classification in this sample that has never been presented to the classifier. The results are the average values of running this process for each pathogen example.

### 3.3 Results and discussion

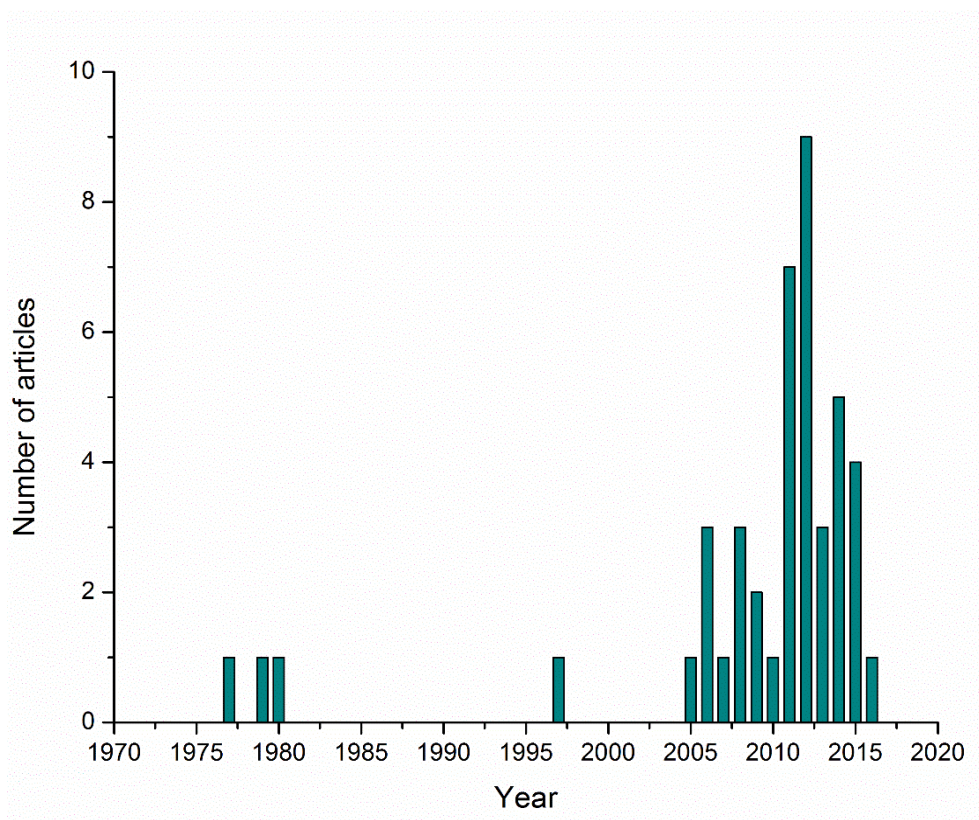
#### 3.3.1 Pathogen-VOC interactions in the bibliography

The PubMed search resulted in 9738 articles of which 341 articles were selected based on title and/or abstract. Full text was read and tested for inclusion criteria, and this step resulted in the inclusion of 180 articles in this study (Figure 3.1). The Web of Science search resulted in 474 articles. 116 articles were selected based on title and/or abstract. After reading the full text, 81 additional publications fitted the criteria. The results obtained using the two online databases were compared and duplicated hits were removed, finally resulting in the inclusion of 44 articles in the study, based on their full text content.

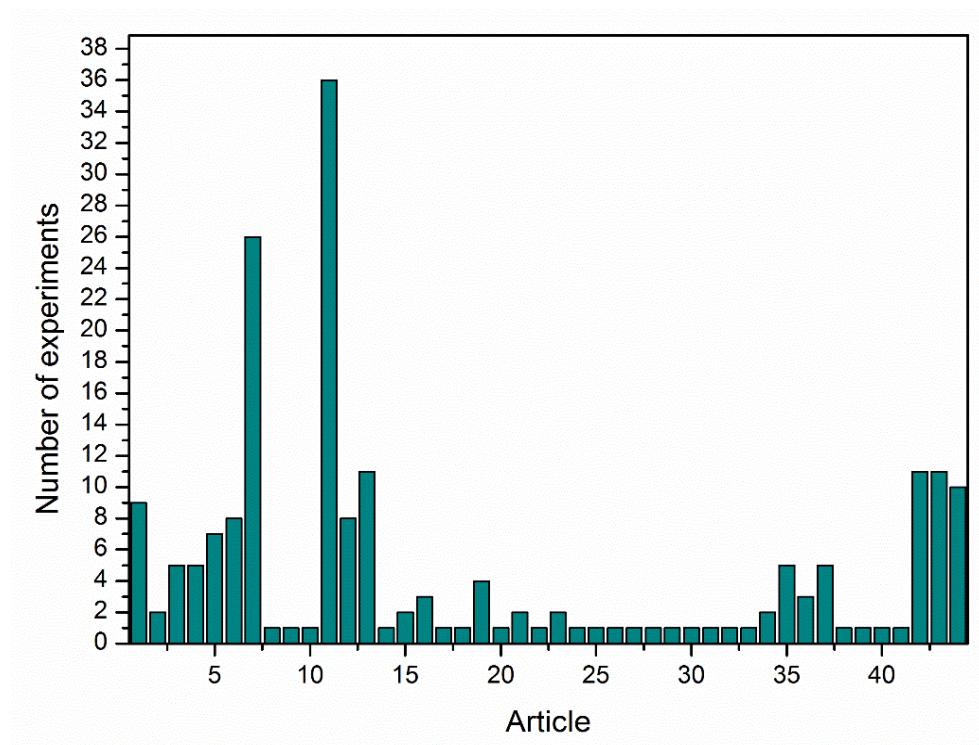
The included articles were published between 1977 and 2016 (Figure 3.5), with an accentuated increase in the number of publications between 2011 and 2012. Most of the publications concerns the last 11 years (2005-2016), corresponding to 88.2% of the total number of collected articles, while the articles concerning the previous years corresponds to 11.8% of the total articles, showing that VOCs have been increasingly studied as potential disease biomarkers over the last 10 years.

In total, the 44 articles report 23 pathogens, 418 VOCs and 199 experiments. The number of experiments present in each articles varies. Papers 11 and 7 are the ones with more associated experiments (36 and 26, respectively) while there are many papers with only 1 reported experiment, such as articles 8, 9 and 10 (Figure 3.6).

The selected articles refer to 23 distinct disease associated pathogens (Table 3.1 and Figure 3.7): *Pseudomonas aeruginosa* (PA), *Staphylococcus aureus* (SA), *Methicillin-resistant Staphylococcus aureus* (MRSA), *Streptococcus pneumoniae* (SP), *Klebsiella pneumoniae* (KB), *Haemophilus influenzae* (HI), *Aspergillus fumigatus* (AF), *Morganella morganii* (MM), *Proteus mirabilis* (PM), *Proteus vulgaris* (PV), *Staphylococcus epidermidis* (SE), *Enterococcus faecalis* (EF), *Candida albicans* (CA), *Escherichia coli* (EC), *Helicobacter pylori* (HP), *Legionella pneumophila* (LP), *Clostridium difficile* (CD), *Campylobacter jejuni* (CJ), *Mycobacterium tuberculosis* (MT), *Giardia duodenalis* (GD), *Plasmodium falciparum* (PF), *Neisseria meningitidis* (NM) and *Moraxella catarrhalis* (MC).



**Figure 3.5-** Representation of the number of articles per year.



**Figure 3.6-** Representation of the number of experiments reported in each article.

**Table 3.1-** Pathogens referred in the selected papers, respective classification, analysis methods used to detect VOCs and number of experiments associated with each pathogen. Pathogens marked (\*) were studied in more than 3 experiments. *Pseudomonas aeruginosa* (PA), *Staphylococcus aureus* (SA), *Methicillin-resistant Staphylococcus aureus* (MRSA), *Streptococcus pneumoniae* (SP), *Klebsiella pneumoniae* (KB), *Haemophilus influenzae* (HI), *Aspergillus fumigatus* (AF), *Morganella morganii* (MM), *Proteus mirabilis* (PM), *Proteus vulgaris* (PV), *Staphylococcus epidermidis* (SE), *Enterococcus faecalis* (EF), *Candida albicans* (CA), *Escherichia coli* (EC), *Helicobacter pylori* (HP), *Legionella pneumophila* (LP), *Clostridium difficile* (CD), *Campylobacter jejuni* (CJ), *Mycobacterium tuberculosis* (MT), *Giardia duodenalis* (GD), *Plasmodium falciparum* (PF), *Neisseria meningitidis* (NM) and *Moraxella catarrhalis* (MC).

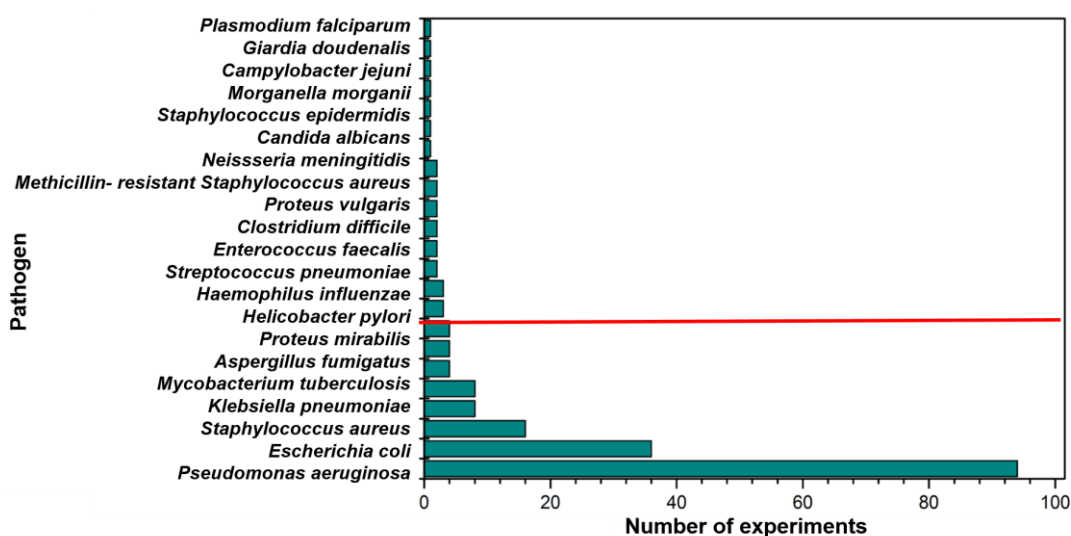
Pathogen	Classification	Analytical Method	Number of experiments	Type of sample	Refs.	Example
<i>Aspergillus fumigatus</i> *	Fungus	GC-MS; IMS; SIFT-MS	4	clinical isolate reporter-labeled strains	[24][37], [55], [99]	AF was detected in a breath sample headspace, associated with sinusitis [24]
<i>Campylobacter jejuni</i>	Gram <sup>-</sup> bacterium	GC-MS	1	clinical isolate	[100]	CJ was detected in faecal samples headspace, associated with gastroenteritis [100]
<i>Candida albicans</i>	Fungus	SIFT-MS	1	reporter-labeled strains	[101]	CA was inoculated into healthy males urine samples [101]
<i>Clostridium difficile</i>	Gram <sup>+</sup> bacterium	GC-MS	2	clinical isolate	[100] [102]	CD was detected in faecal samples associated with gastroenteritis [100]
<i>Enterococcus faecalis</i>	Gram <sup>+</sup> bacterium	SIFT-MS	2	clinical isolate reporter-labeled strains	[101] [103]	EF was inoculated into healthy males urine samples [101]
<i>Escherichia coli</i> *	Gram <sup>-</sup> bacterium	GC-MS; IMS; IMR-MS;PTR-MS; SESI-MS; SIFT-MS	36	clinical isolate reporter-labeled strains	[101] [61], [62], [70], [92], [103]–[110]	EC was detected in blood, urine and skin samples headspaces, associated with urinary tract infection [13]

Pathogen	Classification	Analytical Method	Number of experiments	Type of sample	Refs.	Example
<i>Giardia duodenalis</i>	Protozoa	GC-MS	1	clinical isolate	[111]	<i>GD</i> was detected in faecal samples headspace, associated with Giardiasis [111]
<i>Haemophilus influenzae</i>	Gram <sup>-</sup> bacterium	GC-MS	3	clinical isolate	[40], [112], [113]	<i>HI</i> was found in clinical isolates (origin not reported) from patients with sinusitis [113]
<i>Helicobacter pylori</i> <sup>*</sup>	Gram <sup>-</sup> bacterium	GC-MS; PTR-MS	4	clinical isolate reporter-labeled strains	[24][114]	<i>HP</i> was detected in breath sample headspace, associated with gastric infections [24][114]
<i>Klebsiella pneumoniae</i> <sup>*</sup>	Gram <sup>-</sup> bacterium	GC-MS; SIFT-MS	8	clinical isolate reporter-labeled strains	[101] [104] [62] [105] [110]	<i>KP</i> was found in clinical isolates (origin not reported) from patients with pneumonia [105] and from patients with urinary tract infections [110].
<i>Legionella pneumophila</i>	Gram <sup>-</sup> bacterium	GC-MS	1	unknown reporter-labeled strain	[113]	<i>LP</i> was inoculated in blood cultures, associated with Legionellosis [113]
<i>Methicillin-resistant Staphylococcus aureus</i>	Gram <sup>+</sup> bacterium	GC-MS	2	clinical isolate reporter-labeled strains	[104] [115]	<i>MRSA</i> was grown in Mueller Hinton broth and trypticase soy agar at 37°C [115]
<i>Moraxella catarrhalis</i>	Gram <sup>-</sup> bacterium	GC-MS	2	clinical isolate	[113] [40]	<i>MC</i> was found in clinical isolates (origin not reported) from patients with sinusitis [113]

Pathogen	Classification	Analytical Method	Number of experiments	Type of sample	Refs.	Example
<i>Morganella morganii</i>	Gram <sup>-</sup> bacterium	GC-MS	1	reporter-labeled strains	[106]	<i>MM</i> was inoculated into brain-heart-infusion broth, associated with urinary tract infections [106]
<i>Mycobacterium tuberculosis</i> *	Gram <sup>+</sup> bacterium	GC-MS; GC-SAW	8	clinical isolate reporter-labeled strains	[24][116]–[121]	<i>MT</i> was detected in breath [117], sputum [116] [119] and urine [24] [121] samples headspaces, associated with tuberculosis
<i>Neisseria meningitidis</i>	Gram <sup>-</sup> bacterium	GC-MS; SIFT-MS	2	clinical isolate reporter-labeled strains	[108] [122]	<i>NM</i> was detected in blood samples headspaces [108]
<i>Plasmodium falciparum</i>	Protozoa	GC-MS	1	clinical isolate	[12]	<i>PF</i> was detected in breath samples associated with malaria [12]
<i>Proteus mirabilis</i> *	Gram <sup>-</sup> bacterium	GC-MS; SIFT-MS	4	clinical isolate reporter-labeled strains	[103] [106] [110]	<i>PM</i> was detected in urine sample headspace, associated with urinary tract infection [103]
<i>Proteus vulgaris</i>	Gram <sup>-</sup> bacterium	IMR-MS; SIFT-MS	2	reporter-labeled strains	[101] [70]	<i>PV</i> was inoculated into healthy males urine samples [101]
<i>Pseudomonas aeruginosa</i> *	Gram <sup>-</sup> bacterium	GC-MS; IMS; IMR-MS; SESI-MS; SIFT-MS	94	clinical isolate reporter-labeled strains	[24][37][101] [61], [62], [92], [103], [104] [108] [70] [110] [40] [39], [123]–[130]	<i>PA</i> was detected in breath [24], blood, urine and skin [61] samples headspaces, associated with pneumonia

Pathogen	Classification	Analytical Method	Number of experiments	Type of sample	Refs.	Example
<i>Staphylococcus aureus</i> *	Gram <sup>+</sup> bacterium	GC-MS; IMS; SESI-MS; SIFT-MS	16	clinical isolate reporter-labeled strains	[101][62], [92], [103], [104] [105] [108] [40] [123][131]	SA was detected in breath samples headspaces, associated with pneumonia [123][131]
<i>Staphylococcus epidermidis</i>	Gram <sup>+</sup> bacterium	SIFT-MS	1	reporter-labeled strains	[101]	SE was inoculated into healthy males urine samples [101]
<i>Streptococcus pneumoniae</i>	Gram <sup>+</sup> bacterium	GC-MS; SIFT-MS	3	clinical isolate reporter-labeled strains	[108] [112] [40]	SP was detected in breath samples, from patients with pneumonia [112]

During this work, it was evident that there are pathogens more studied than others. The pathogen with more associated experiments is the bacterium *Pseudomonas aeruginosa* (94), corresponding to 47.24% of the total number of experiments. This pathogen is followed by *Escherichia coli* (36 associated experiments), *Staphylococcus aureus* (16 associated experiments), *Klebsiella pneumoniae* (8 associated experiments), *Mycobacterium tuberculosis* (8 associated experiments), *Aspergillus fumigatus* (4 associated experiments), *Proteus mirabilis* (4 associated experiments) and *Helicobacter pylori* (4 associated experiments) (Figure 3.7). The rest of the microorganisms has less than 4 associated experiments (Table 3.1).



**Figure 3.7-** Representation of the number of experiments associated with the 23 pathogens. The red line separates the pathogens with more than 3 associated experiments (below the line) from the pathogens with 3 or less associated experiments (above the line).

The experimental analysis of VOCs involves several steps that vary with the chosen analytical method [14][24][84]. The general steps include sampling (collection and storage, if needed, of the sample), headspace analysis [61][86][104], VOC quantification/identification [103][125], data processing [72], and, finally, as an output, putative VOC biomarkers should be obtained.

The experiments associated with each pathogen vary in the type of sample used (Figure 3.8). It was found that most of the experiments used commercial acquired microbial strains, known as reporter-labeled strains (50.2%), 16.6% of the experiments did not indicate the strain used in the study, referring only the microbe species group. The remaining 33.2% of the experiments used human clinical isolates collected from urine (11.1%), breath (8.5%), blood (6.1%), skin (5.5%) and faeces (2.0%). The high percent of experiments using urine and breath may be due to the growing interest in non-invasive diagnostic tools [58][59][60].



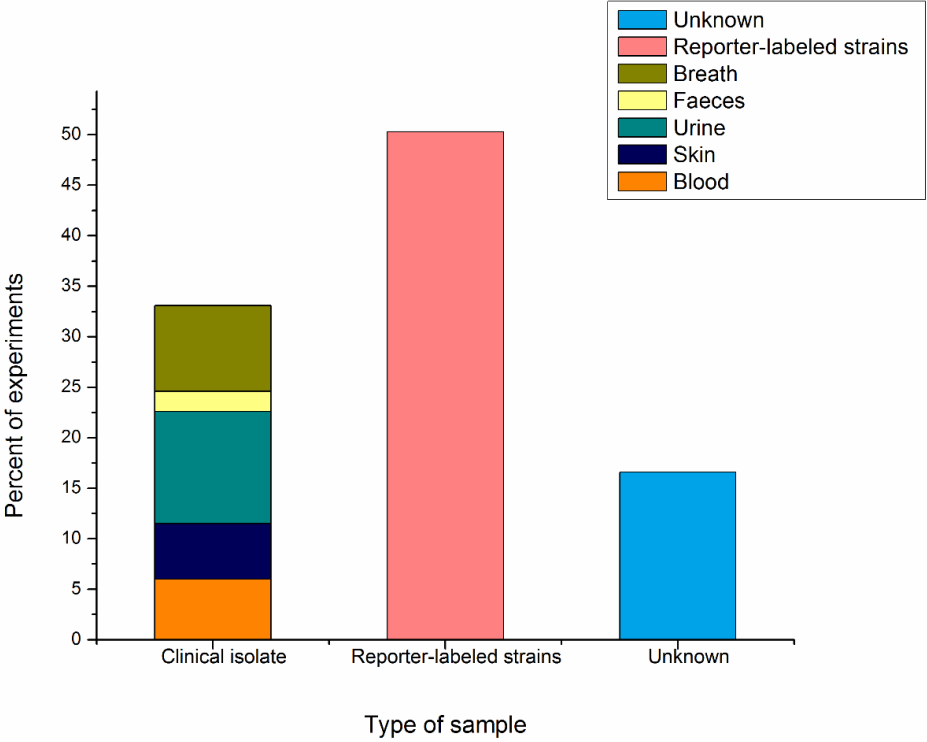
Different methods of analysis were used in the studied publications and sometimes more than one analysis method was employed in the same article. Clear differences were observed, regarding the most oftenly used methods (Figure 3.9). The 3 mostly used analytical methods to detect and quantify VOCs were IMS, SIFT-MS and GC-MS, mainly due to their high sensitivity (ppt<sub>v</sub>-ppb<sub>v</sub>) [16][58][61]–[65]. Interestingly, IMS and SIFT-MS allow real-time analysis while GC-MS only offers an off-line analysis, despite remaining the gold standard method [77][87]. Together these three analytical methods account for 96% of the total number of experiments, while the remaining 4 methods (GC-SAW, PTR-MS, IMR-MS and SESI-MS) only account for 4%.

The reported VOCs were grouped according to their chemical class to allow an easier overall comparison. Some compounds could be fitted in more than one class. In these situations, one of those chemical classes was randomly chosen. For instance, methyl thiocyanate contains both nitrogen and sulfur and it was classified as a sulfur containing compound (See Appendix 1).

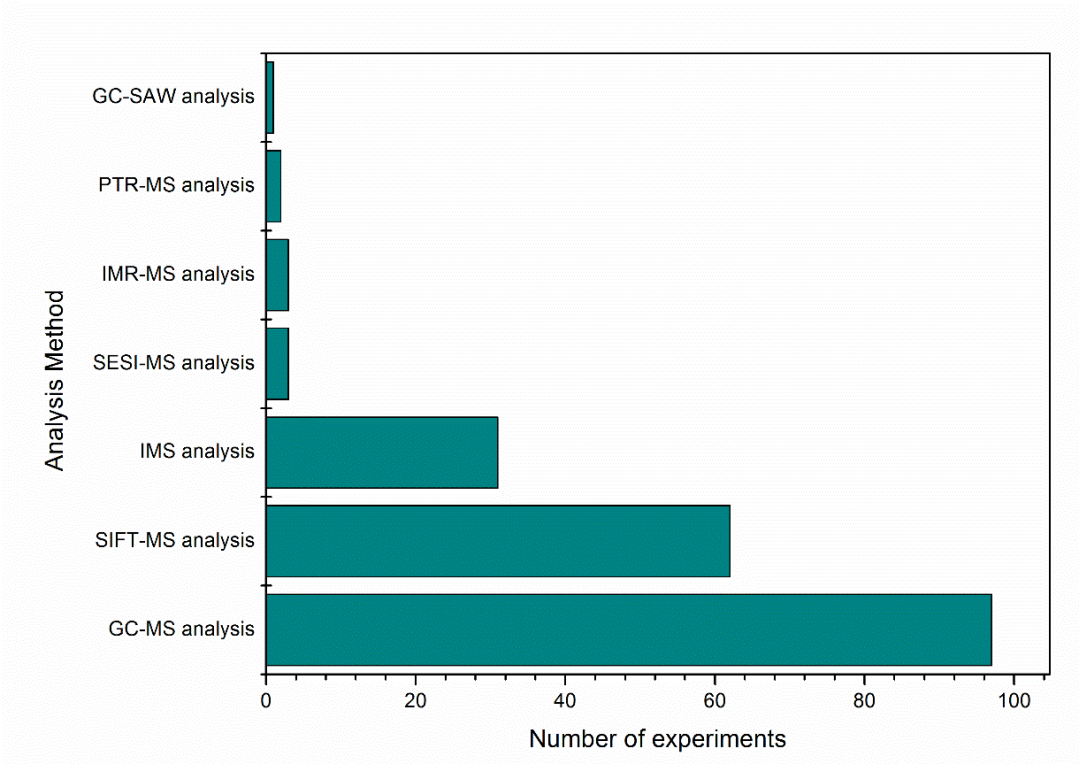
The relative abundance of each class was calculated by dividing the number of hits of each class by the total number of hits concerning all the classes (eq.1). The order of abundance of each compound class in all of the experiments was the following: alcohols > hydrocarbons > nitrogen-containing > ketones > sulfur-containing > aldehydes > esters > acids > furans and ethers > others > halogen-containing (Figure 3.10). The most abundant class is the class of the alcohols. However, the abundance values for the 5 most abundant classes are in the same order of magnitude, being almost equally distributed.

$$\text{Relative abundance (\%)} = \frac{\text{Number of hits}_{\text{class}}}{\text{Total number of hits}} \times 100$$

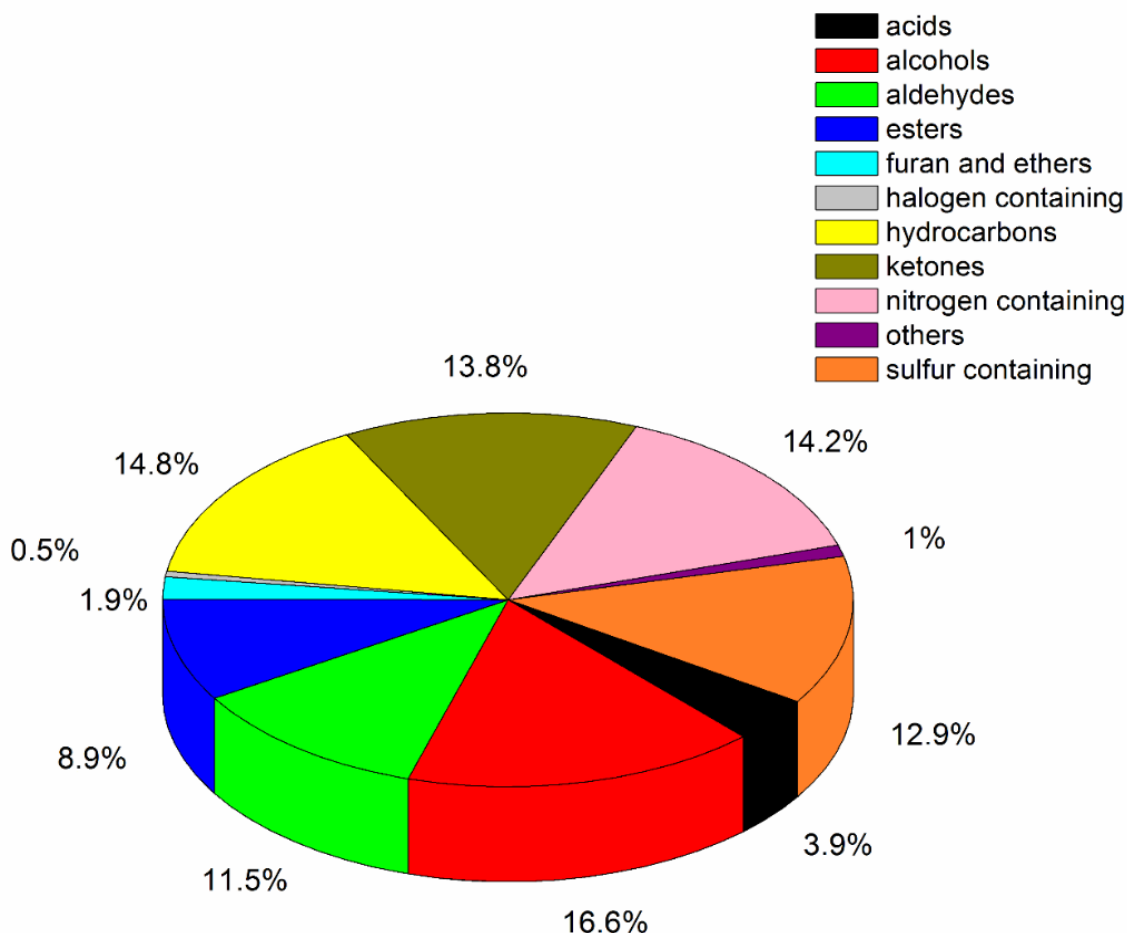
(eq.1)



**Figure 3.8-** Samples used to obtain VOC biomarkers.

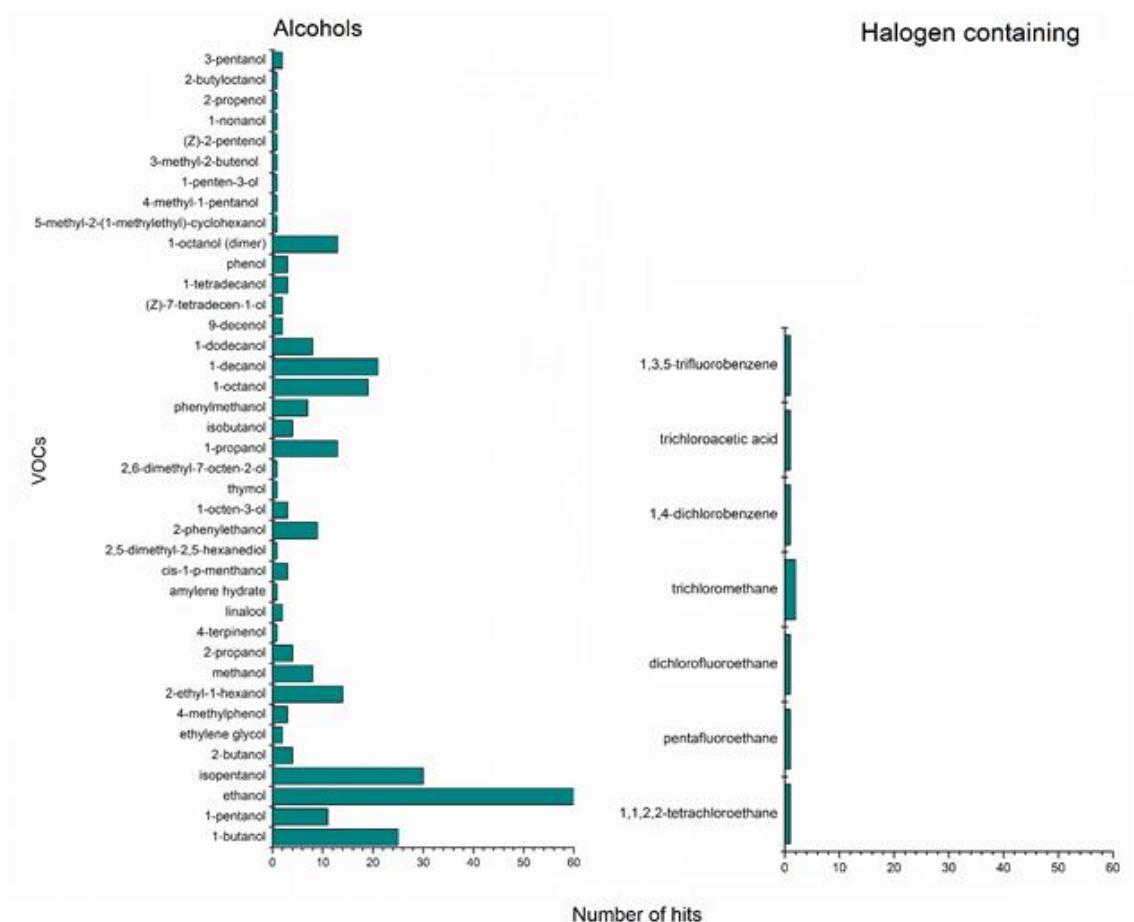


**Figure 3.9-** Analysis methods used and the corresponding number of experiments.



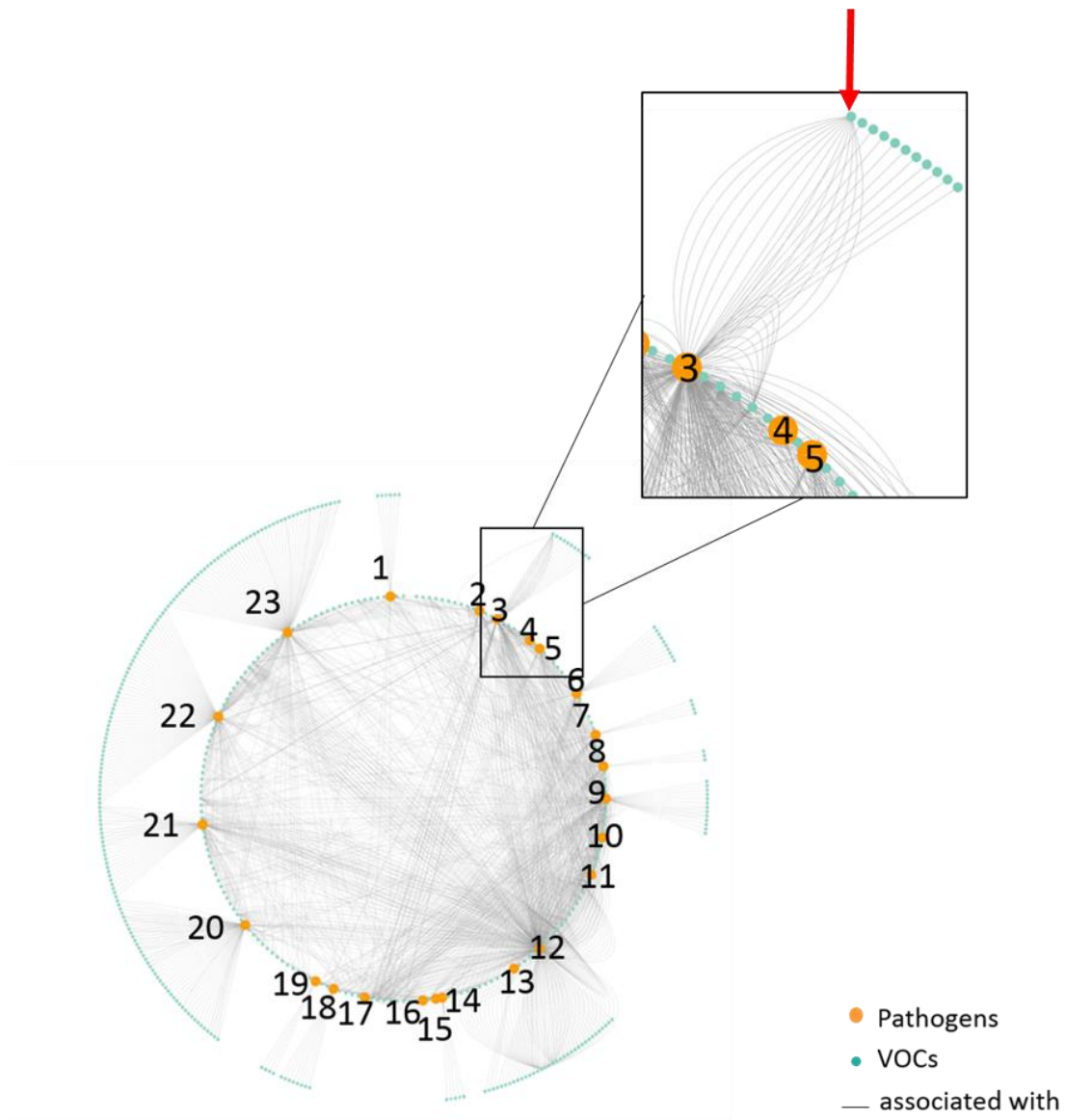
**Figure 3.10-** Piechart with the relative number of hits (% relative abundance) of compounds in each class that have been detected in all the experiments.

Ethanol is the most frequently detected VOC in pathogen culture headspaces. This volatile is produced by 14 of the 23 studied pathogens (AF, CA, CD, EF, EC, GD, HP, KP, MRSA, NM, PM, PA, SA, SP) (Figure 3.11). Regarding hydrocarbons, 1-undecene is the most reported, but is associated to only 2 microorganisms (PA, KB), (See Appendix 1) and the most referred nitrogen containing compound is hydrogen cyanide, associated with PA and HP (See Appendix 1). The class of hydrocarbons contains 106 distinct VOCs, being the most diverse class (See Appendix 1), while the less diversified are the halogen containing, and acids classes (See Appendix 1).



**Figure 3.11** - Representation of the number of hits found for each individual VOC, in all experiments, for the alcohols and the halogen containing classes.

The literature search resulted in 44 articles reporting 23 associated pathogens, each of them associated to a VOC pattern. In the circular graph representation shown in figure 3.12 the complexity of the pathogen -VOC relations is notable. While some VOCs are associated with more than one pathogen (the ones represented in the circumference), others were only reported to be associated with one pathogen (the ones outside the circle). For instance, for pathogen 3 (*Escherichia coli*) there is a group of VOCs that is exclusively associated with this bacterium, but these relations were referred only one time (one hit). Those are represented as nodes outside the circle with a straight line connecting them to the *Escherichia coli* node. It is also notable that one of the *Escherichia coli* exclusive VOCs (outside the circle) has been mentioned in several experiments (more than one hit), represented by the oval shaped lines connecting it to *Escherichia coli* and evidenced by a red arrow (inset of Figure 3.12).

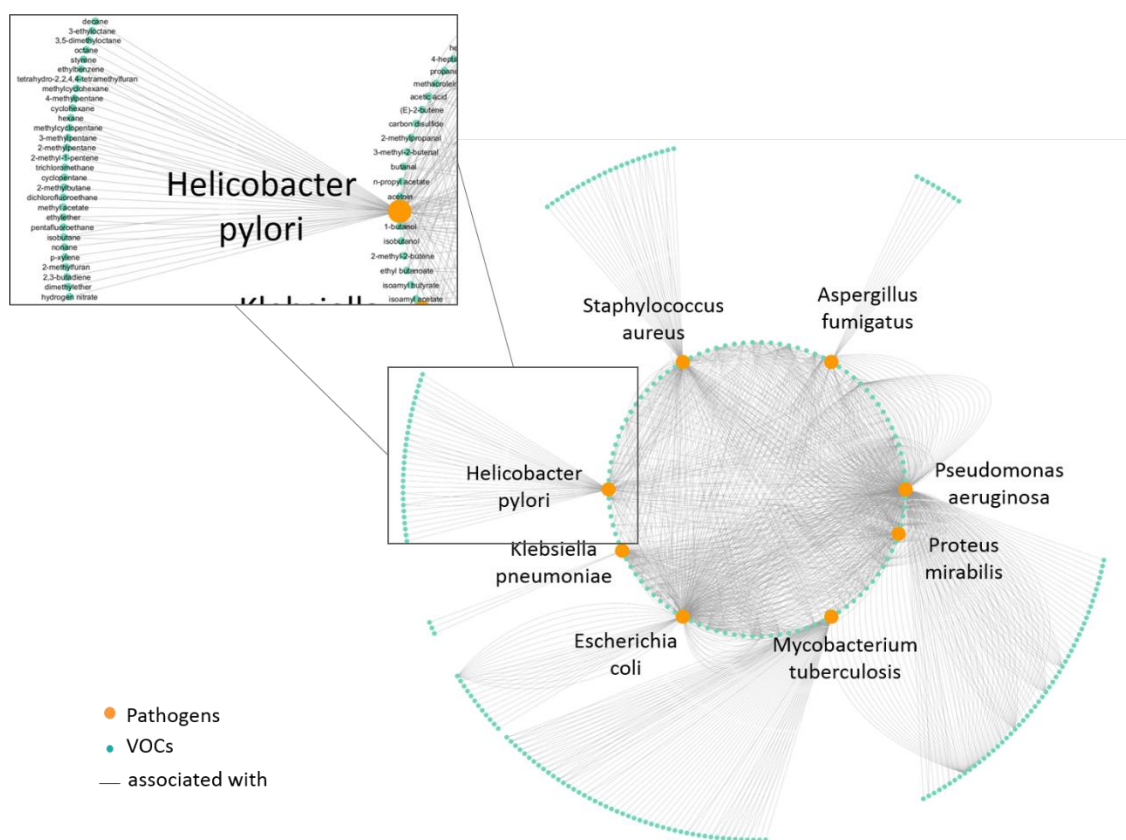


**Figure 3.12-** Representation of all the 23 pathogens and 418 associated VOCs. 1- *Aspergillus fumigatus*, 2- *Enterococcus faecalis*, 3- *Escherichia coli*, 4- *Morganella morganii*, 5- *Proteus mirabilis*, 6- *Haemophilus influenzae*, 7- *Streptococcus pneumoniae*, 8- *Klebsiella pneumoniae*, 9- *Staphylococcus aureus*, 10- *Proteus vulgaris*, 11- *Staphylococcus epidermidis*, 12- *Pseudomonas aeruginosa*, 13- *Plasmodium falciparum*, 14- *Neisseria meningitidis*, 15- *Legionella pneumophila*, 16- *Moraxella catarrhalis*, 17- *Candida albicans*, 18- *Giardia duodenalis*, 19- *Methicilin-resistant staphylococcus aureus*, 20- *Mycobacterium tuberculosis*, 21- *Helicobacter pylori*, 22- *Campylobacter jejuni*, 23- *Clostridium difficile*. The arrow in the inset highlights a VOC that is associated with one pathogen and that was reported in more than one experiment.

### 3.3.2 VOCs distinguishing pathogens

It was evident from the circular layout representing the full dataset (Figure 3.12), that a distinction between the pathogens based on the reported associated VOCs could be possible. The next step was to identify the set of VOCs that play the major roles in this distinction. A supervised machine-learning algorithm was used to devise an automated VOC-based pathogen classifier. The VOC-pathogen interaction subset data, including pathogens with more than 3

experiments (8 pathogens, 269 VOCs) was used as input for the algorithm to ensure a better equilibrium between the information in the model. The circular graph for the 8 pathogens (Figure 3.13) shows the significant reduction of complexity, compared with the representation of the full dataset (Figure 3.12). For this set of microorganisms there are still shared VOCs (those in the circumference), however, it is visually remarkable the existence of a group of VOCs exclusive to almost each pathogen (outside the circle), as it is the case of the *Helicobacter pylori* (inset of Figure 3.13).



**Figure 3.13-** Circular layout of the 8 pathogens with more than 3 associated experiments, and the 269 associated VOCs.

From the 269 VOCs in the dataset, a profile with 26 VOCs was identified by the algorithm as the set of VOCs that allows to better separate the 8 pathogens. This selection was based on the VOCs score contributions for the separation model (Table 3.2).

**Table 3.2-** Set of 26 VOCs that better separates the 8 analyzed pathogens, selected interactively, by an identification approach. The VOCs are listed from the highest to the lowest score.

Set of VOCs that better separates the 8 analyzed pathogens	
1.	indole
2.	3-methylbutanoic acid
3.	1,3,5-trimethylbenzene
4.	cymol
5.	isobutane
6.	2-phenylanisol
7.	1,1,2,2-tetrachloroethane
8.	1-decanol
9.	1-undecene
10.	isopentanol
11.	acetoin
12.	1-heptene
13.	1-hydroxy-2-butanone
14.	2-methylbutanal
15.	2-pentanone
16.	2-phenylethanol
17.	dimethyl disulfide
18.	ammonia
19.	1-pentanol
20.	3-octanone
21.	acetic acid
22.	hydrogen nitrate
23.	1-dodecanol
24.	1-propanol
25.	isoprene
26.	1-hydroxy-2-propanone

The classification model was validated using the “leave-one-out” cross-validation method. The limitation of evaluating a model without cross-validation is that we do not know how well the classifier will do when it is asked to make new predictions for data that it has never seen before. In our case, the obtained accuracy rate was of 81%, slightly less than when no cross-validation was used (94%) (Table 3.3).

Despite having high accuracy rates, in both cases, is important to refer that the classes (pathogens) are not equally represented in the model, since the number of associated experiments varies. So, the reason to the high accuracy may be due to the existence of imbalanced data, because the model looks at the data and decides that the best thing to do is to always predict *Pseudomonas aeruginosa*, since it is the class with more instances.

**Table 3.3-** Accuracy rates obtained by classification with and without cross-validation.

Accuracy rate	
With cross-validation	0.810
Without cross-validation	0.936

The confusion matrix for the classification with highest accuracy rate (without cross-validation) value shows the correct predictions (diagonal) and the types of incorrect predictions made by the model (which pathogens are being confused by another) (Table 3.4). It can be seen that the classifier never confuses *H.pylori* or *M.tuberculosis* with other pathogens. There are 3 microorganisms that are only mislabeled once, namely: *A. fumigatus* as *P.aeruginosa*, both associated with sinusitis (Table 3.5), *P.mirabilis* as *E.coli*, both associated with urinary tract infections, and *P.aeruginosa* as *E.coli*, both found in urinary tract infections. In addition, *E.coli* was mislabeled twice as *P. aeruginosa*. Finally, *S.aureus* and *K.pneumoniae* were mislabeled 3 times each. The first was confused 2 times with *E.coli* (both found in urinary tract infections) and 1 with *P.aeruginosa* (both found in urinary tract infections, pneumonia and cystic fibrosis). The second was mislabeled 1 time with *E.coli* (both associated with urinary tract infections) and 2 with *P.aeruginosa* (both found in urinary tract infections, pneumonia, blood and bone infections), respectively). So, in total there were 163 true positives (predicted pathogen was the same as the actual pathogen) and 11 false positives, corresponding to a classification accuracy rate of 93.6%. This confusion can be due to the fact that the mislabeled pathogens emit similar compounds. For instance, *P.aeruginosa* and *S. aureus* both share 36 VOCs with *E.coli*, being the pathogens with higher number of shared compounds. Also, regarding these 3 pathogens, within the shared compounds, the most relevant classes are the aldehydes, ketones, alcohols, nitrogen containing and sulfur containing. *P.aeruginosa* and *E.coli*, share 8 aldehydes and 7 ketones (which contain both a carbonyl group), 5 alcohols, 5 sulfur containing and 6 nitrogen containing compounds. Similarly, *E.coli* and *S. aureus*, have 7 aldehydes, 6 ketones, 6 sulfur containing, 5 nitrogen containing compounds and 5 alcohols in common.

Given the class unbalance, in cases where there are reduced samples of the pathogen, the trivial classification as not the pathogen (uninformed accuracy) has already a high accuracy level. To better report the improvements of the classifier learning, we present the base accuracy and compare with the final classification accuracy after selecting the feature subset and train the classifier (Table 3.6). As a result, we obtained a putative set of VOCs that allows a separation between the pathogens with high final classification accuracy values.



**Table 3.4-** Confusion matrix for the classification without cross-validation, where the column labels represent the predicted pathogen classification and the line labels represent the actual pathogen identity. Red represents the incorrect predictions made by the model and green the correct predictions (diagonal).

			Predicted							
Actual			<i>A.fumigatus</i>	<i>E.coli</i>	<i>H.pylori</i>	<i>K.pneumoniae</i>	<i>M.tuberculosis</i>	<i>P.mirabilis</i>	<i>P.aeruginosa</i>	<i>S.aureus</i>
	Fungus	<i>A.fumigatus</i>	3	0	0	0	0	0	1	0
	Gram <sup>-</sup> bacterium	<i>E.coli</i>	0	34	0	0	0	0	2	0
	Gram <sup>-</sup> bacterium	<i>H.pylori</i>	0	0	4	0	0	0	0	0
	Gram <sup>-</sup> bacterium	<i>K.pneumoniae</i>	0	1	0	5	0	0	1	1
	Gram <sup>+</sup> bacterium	<i>M.tuberculosis</i>	0	0	0	0	8	0	0	0
	Gram <sup>-</sup> bacterium	<i>P.mirabilis</i>	0	1	0	0	0	3	0	0
	Gram <sup>-</sup> bacterium	<i>P.aeruginosa</i>	0	1	0	0	0	0	93	0
	Gram <sup>+</sup> bacterium	<i>S.aureus</i>	0	2	0	0	0	0	1	13

These pathogens are mainly associated with respiratory infections (tuberculosis (MT), pneumonia (SA, KP, PM, PA), cystic fibrosis (AF, SA, PA) and sinusitis (AF, SA, PA)) urinary tract infections (SA, EC, KP, PM, PA), gastric (SA, HP) and gastrointestinal infections (EC, PA) (Table 3.5). Tuberculosis is one of the clinically most relevant disease, since, although it is curable and preventable, affects every part of the world and an earlier diagnosis is essential to a positive outcome. Also, pneumonia is the largest infectious cause of death in children and elderly, worldwide and an early detection of the disease and identification of the involved microorganism have vital importance. Therefore, the determination of pathogen specific biomarkers is crucial.

Biomarkers have been emerging as a dynamic and powerful approach to screen and detect a disease. In this study we have successfully determinate a set of VOCs that allow pathogen separation, based on VOCs released by microbial pathogens by consolidating results obtained by several authors, respecting pathogens causing infections.

A recent review descriminate 1840 VOC compounds identified from healthy humans, and the respective bodily fluids [133]. Although the typical VOC concentrations ranges in the normal and disease states are not described, it is important to see whether the sets of VOCs that better separates the 8 pathogens, obtained by identification and verification approaches are present in the healthy body or not, and if so in which bodily fluid they can be found. By comparing the healthy body data with the set of VOCs that better separates the studied pathogens, obtained by an identification approach, 3 VOCs were found that are not reported, so far, in the healthy human body (Table 3.7). These compounds are 2-phenylanisol, 1-hydroxy-2-butanone and hydrogen nitrate. Interestingly, each of these 3 VOCs was reported as being associated with only one pathogen and is part of the set of VOCs that better separates determined in this work for that pathogen: 2-phenylanisol was associated twice with *M.tuberculosis* (reporter-labeled strain) [119][120], 1-hydroxy-2-butanone was associated once with *K.pneumoniae* (unknown type of sample) [105] and hydrogen nitrate was associated once with *H.pylori* (reported-labeled strain) [24]. By comparing the list of VOCs that better separates the 8 pathogens, obtained by a verification approach, with the VOCs found in the healthy body, 4 more VOCs were never found in the headspace of any healthy bodily fluid: 1-methylethenyl-pyrazine associated with *A.fumigatus* [37], 3-methylcyclohexene, associated with *K.pneumoniae* [104], 2,3,4,5-tetrahydropyridazine and 2-methylbutanoate, associated with *S.aureus* [105] (Table 3.8). Therefore, the detection of any of these 7 VOCs in some human sample's headspace may facilitate the distinction of pathogens present in an infection.

Knowing the human bodily fluids where a VOC is normally found may facilitate the choice of the most suitable fluid to be analyzed. We verified that when testing for the presence of *E.coli* in

urinary tract infections, the type of collected sample varies between blood, skin and urine [61]. Curiously, indole is reported to be present in healthy skin and urine samples but not in breath, milk and blood samples. Therefore, when investigating the presence of *E.coli*, the presence of indole in blood may be an indicator of an infection caused by that pathogen.

**Table 3.5** – Associated diseases/infections to each of the 8 studied pathogens. AR- allergic reactions. BLI- blood infections. BI- bone infections. CF- cystic fibrosis. GI- gastric infections. GTI- gastrointestinal infections. HI- heart infections. KI- kidney infections. PN- pneumonia. SI- sinusitis. SKI- skin infections. TB- tuberculosis. UTI- urinary tract infections.

Pathogen		Associated diseases/infections												
		AR	BLI	BI	CF	GI	GTI	HI	KI	PN	SI	SKI	TB	UTI
<i>A. fumigatus</i>	Fungus	x			x						x			
<i>S. aureus</i>	Gram <sup>+</sup> bacterium		x	x	x	x		x		x	x	x		x
<i>E. coli</i>	Gram <sup>-</sup> bacterium						x							x
<i>H. pylori</i>	Gram <sup>-</sup> bacterium					x								
<i>K. pneumoniae</i>	Gram <sup>-</sup> bacterium		x	x						x				x
<i>M. tuberculosis</i>	Gram <sup>+</sup> bacterium												x	
<i>P. mirabilis</i>	Gram <sup>-</sup> bacterium								x	x				x
<i>P. aeruginosa</i>	Gram <sup>-</sup> bacterium		x	x	x		x	x		x	x			x

**Table 3.6-** Representation of the results obtained by the verification approach.

Pathogen	Number of experiments	Set of VOCs that better separates	Uninformed accuracy	Final classification accuracy
<i>A. fumigatus</i>	4	1-methylethenyl-pyrazine 2-pentylfuran cyclohexanone pentanal	0.98	1.0
<i>E. coli</i>	36	indole	0.79	0.92
<i>H. pylori</i>	4	isobutane hydrogen nitrate	0.98	1.0
<i>K. pneumoniae</i>	8	1-hydroxy-2-butanone 3-methylcyclohexene	0.95	0.97
<i>M. tuberculosis</i>	8	1,3,5-trimethylbenzene cymol 2-phenylanisol	0.95	1.0
<i>P. mirabilis</i>	4	(E)-2-butene	0.98	0.98

Pathogen	Number of experiments	Set of VOCs that better separates	Uninformed accuracy	Final classification accuracy
<i>P. aeruginosa</i>	94	ethanol 1-decanol 3-methylbutanal 2-phenylethanol 1,3,5-trimethylbenzene cymol hydrogen cyanide 2-pentene ethane	0.54	0.86
<i>S. aureus</i>	16	3-methylbutanoic acid 1,1,2,2-tetrachloroethane 2,3,4,5-tetrahydropyridazine 2-methylbutanoate pyrimidine	0.91	0.95

**Table 3.7-** Comparison between the set of VOCs that better separates the group of 8 pathogens, obtained by identification approach, and the VOCs reported in the literature as present (X) or absent ( ) in healthy bodily fluids. NR-not reported.

VOCs	PubChem ID	Healthy body						
		Faeces	Urine	Breath	Skin	Milk	Blood	Saliva
indole	798	x	x		x			x
3-methylbutanoic acid	10430	x		x	x			
1,3,5-trimethylbenzene	7947			x				
cymol	7463		x	x	x			x
isobutane	6360			x				
2-phenylanisol	6835	NR	NR	NR	NR	NR	NR	NR
1,1,2,2-tetrachloroethane	6591		x				x	
1-decanol	8174	x			x			
1-undecene	13190			x				
isopentanol	31260	x	x				x	x
acetoin	179	x		x				x
1-heptene	11610			x				
1-hydroxy-2-butanone	521300	NR	NR	NR	NR	NR	NR	NR
2-methylbutanal	7284	x	x	x	x			x
2-pentanone	7895	x	x	x	x	x		x
2-phenylethanol	6054	x			x	x		x
dimethyl disulfide	12232	x	x			x		x
ammonia	222		x	x			x	
1-pentanol	6276	x	x	x	x	x	x	x
3-octanone	246728	x	x	x		x		x
acetic acid	176	x	x	x	x	x		x
hydrogen nitrate	944	NR	NR	NR	NR	NR	NR	NR
1-dodecanol	8193	x			x			x
1-propanol	1031	x	x	x		x	x	x
isoprene	6557			x	x	x	x	
1-hydroxy-2-propanone	8299	x		x				

**Table 3.8-** Comparison between the VOCs that better separates the 8 pathogens, obtained by a verification approach, and the VOCs reported in the literature as (X) or absent ( ) in healthy bodily fluids. NR-not reported. (+) -positive biomarker. (-) – negative biomarker.

Pathogen	VOCs	PubChem ID	Healthy body						
			Faeces	Urine	Breath	Skin	Milk	Blood	Saliva
AF	1-methylethenyl-pyrazine (+)	62897	NR	NR	NR	NR	NR	NR	NR
	2-pentylfuran (+)	19602	x	x	x		x		x
	cyclohexanone (+)	7967		x	x			x	
	pentanal (+)	8063	x	x	x		x		x
EC	indole (+)	798	x	x		x			x
HP	isobutane (+)	6360			x				
	hydrogen nitrate (+)	944	NR	NR	NR	NR	NR	NR	NR
KP	1-hydroxy-2-butanone (+)	521300	NR	NR	NR	NR	NR	NR	NR
	3-methylcyclohexene (+)	11573	NR	NR	NR	NR	NR	NR	NR
MT	1,3,5-trimethylbenzene (+)	7947			x				
	cymol (+)	7463		x	x	x			x
	2-phenylanisol (+)	6835	NR	NR	NR	NR	NR	NR	NR
PM	(E)-2-butene (-)	62695			x				
PA	ethanol (+)	702	x	x	x			x	x
	1-decanol (-)	8174	x			x			
	3-methylbutanal (+)	11552	x	x	x				x
	2-phenylethanol (+)	6054	x			x	x		x
	1,3,5-trimethylbenzene (-)	7947			x				
	cymol (-)	7463		x	x	x			x
	hydrogen cyanide (+)	768			x				
	2-pentene (+)	12585			x				
	ethane (+)	6324			x				
SA	3-methylbutanoic acid (+)	10430	x		x	x			
	1,1,2,2- tetrachloroethane (+)	6591		x				x	
	2,3,4,5- tetrahydropyridazine (+)	not available	NR	NR	NR	NR	NR	NR	NR
	2-methylbutanoate (+)	22253297	NR	NR	NR	NR	NR	NR	NR
	pyrimidine (+)	9260				x			

Other authors also focused on the determination of VOC biomarkers [24][134][135]. The published databases acquired data through extensive literature searches. Our results are compared with the VOC fingerprints suggested in those studies in Table 3.9. However, different methods were employed to determine the VOC biomarkers. In our study, as described above, we used a verification method to obtain a set of VOCs that allow a separation between a specific pathogen and the other 7 studied pathogens, based on the information collected from published research. This method selected both negative and positive biomarkers (identified as – or + in Table 3.9). In the mVOC database, available in <http://bioinformatics.charite.de/mvoc/> [135], a similarity determination between the compound of interest and the compounds of the mVOC database, is used based on the Tanimoto coefficient.

In the Bos et al. review [134] the compounds referred in the selected articles were organized in tables per functional group. Then, all the compounds produced by at least one of the bacteria were included in an interaction graph that connected the compounds with all bacteria known to produce them. This step allowed a visual representation of VOCs that were produced by only one pathogen. Those volatiles were suggested as possible biomarkers for the pathogen. In the Sethi review [24], the VOC biomarkers/profiles referred in each article were summarized. In both studies, there was no further processing of the information contained in those articles.

Although the methods to obtain the VOC fingerprints were distinct between the compared databases, some of the compounds identified as putative biomarkers for a certain pathogen were identified in more than one database. For instance, hydrogen cyanide was identified in 3 out of 4 databases as *P.aeruginosa* biomarker, while 2-pentylfuran, indole, isobutane, cymol, methyl thiocyanate and 3-methylbutanoic acid were identified in 2 of the compared databases as *A. Fumigatus*, *E.coli*, *H.pylori*, *M.tuberculosis*, *P.aeruginosa* and *S.aureus* biomarkers, respectively (Table 3.9). The consistent finding of the same putative biomarkers in distinct databases using different data pre-processing and processing methods empowers the possibility of these compounds actually being biomarkers of the pathogens in question.

**Table 3.9-** Comparison between the set of VOCs obtained in our study, the mVOC database and two existing reviews, where “+” means positive biomarker and “-“ negative biomarker.

Pathogen		VOC fingerprint			
		this work	mVOC database [135]	Bos et al. review [134]	Sethi et al. review [24]
<i>Fungus</i>	<i>A. fumigatus</i>	1-methylethenyl-pyrazine +	-	-	-
		2-pentylfuran +	-	-	2-pentylfuran
		cyclohexanone +	-	-	-
		pentanal +	-	-	-
		-	2,4-pentadione	-	-
		-	3-methyl-1,3-pentadione	-	-
<i>Gram<sup>-</sup> bacterium</i>	<i>E.coli</i>	indole +	-	indole	-
		-	-	methanol	-
		-	-	1-pentanol	-
		-	-	ethyl acetate	-
		-	pentyl cyclopropane	-	-
	<i>H.pylori</i>	isobutane +	-	-	isobutane
		hydrogen nitrate +	-	-	-
		-	-	-	2-butanone
		-	-	-	ethyl acetate
	<i>P. aeruginosa</i>	ethanol +	-	-	-
		1-decanol -	-	-	-
		3-methylbutanal +	-	-	-
		2-phenylethanol +	-	-	-
		1,3,5-trimethylbenzene -	-	-	-
		cymol -	-	-	-
		hydrogen cyanide +	-	hydrogen cyanide	hydrogen cyanide
		2-pentene +	-	-	-
		ethane +	-	-	-
		-	-	1-undecene	-
		-	-	2-butanone	-
		-	-	2,4-dimethylheptane	-
		-	-	methyl thiocyanate	methyl thiocyanate
		-	-	4-methyl-quinazoline	-
		-	-	-	2-aminoacetophenone



### Chapter 3: Looking for infectious disease VOC biomarkers

Pathogen		VOC fingerprint				
		this work	mVOC database [135]	Bos et al. review [134]	Sethi et al. review [24]	
	<i>K.pneumoniae</i>	1-hydroxy-2-butanone + 3-methylcyclohexene +	- -	- -	- -	
	<i>P.mirabilis</i>	(E)-2-butene -	-	-	-	
<i>Gram<sup>+</sup> bacterium</i>	<i>M. tuberculosis</i>	1,3,5-trimethylbenzene + cymol + 2-phenylanisol + - - - - - - -	- - - - - - - - - -	- - - - - - - - - -	1-methyl-naphthalene 1,4-dimethylcyclohexane - - - - - -	- cymol - - - o-xylene isopropyl acetate 3-pentanol - - dimethylstyrene
		3-methylbutanoic acid + 1,1,2,2-tetrachloroethane + 2,3,4,5-tetrahydropyridazine + 2-methylbutanoate + pyrimidine +	- - - - -	3-methylbutanoic acid - - - -	- - - - -	
	<i>S. aureus</i>					

### 3.4 Conclusions

Advances in analytical technologies for detecting and measuring VOCs in clinical matrices have generated an increasing interest in the exploitation of VOCs as biomarkers of different diseases. In this work we were able to conclude that the volatolome of *Pseudomonas aeruginosa* is the most studied, which is mostly associated with urinary tract infections and pulmonary infections, such as pneumonia and cystic fibrosis. The collected articles reported mostly reporter-labeled strains. However, when they used clinical isolates, the most used human sample was urine, probably due to the existing standardized procedures for handling and analysis. The GC-MS analysis was the gold standard method to identify the compounds, followed by the SIFT-MS. After grouping the reported VOCs according to their chemical class, it was found that the alcohols class was the most abundant one, while halogen containing compounds were the less abundant. Our systematic literature search resulted in 44 articles, published between 1977 and 2016, reporting 23 distinct pathogens and 418 associated VOCs. We have employed a machine learning method to classify pathogens with more than 3 associated experiments based on the emitted VOCs. VOC-pathogen interaction data was used to build an input data matrix with 8 clinically relevant pathogens (*Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Aspergillus fumigatus*, *Escherichia coli*, *Helicobacter pylori*, *Proteus mirabilis*, *Klebsiella pneumoniae* and *Mycobacterium tuberculosis*) and 269 VOCs. The classifier was able to distinguish the 8 pathogens based on 26 of the 269 VOCs. That set of VOCs was compared with the reported VOCs emitted from a healthy human body [133] and it was found that 3 VOCs (2-phenylanisol, 1-hydroxy-2-butanone and hydrogen cyanide) are not reported, so far, in the healthy human body. We have identified a minimal set of VOCs that allowed the separation of a specific pathogen from the others and compared those VOC lists with the ones found in other studies, for the same pathogens. It was found that indole for *E.coli*; 2-pentylfuran for *A.fumigatus*; isobutane for *H.pylori*; cymol for *M.tuberculosis*; hydrogen cyanide and methyl thiocyanate for *P.aeruginosa*; and 3-methylbutanoic acid for *S.aureus*, were referred in ours and other of the compared databases that used distinct data processing methods. Therefore, these compounds have strong probability of being biomarkers. Nonetheless, more work is required to define the range of normality/disease state in VOCs from humans in terms of concentration ranges in all bodily fluids. This data could then be used to interpretate the constitution of each collected sample obtained from patients, and to monitor their health state or infer about possible pathogen invasions.

## 4. Tailoring biogels selectivity

### 4.1 Introduction

Over the last years, GC-MS and associated methodologies became the most widely used analytical gas detectors for clinical diagnosis. However, these instruments are large, expensive and require trained operators, which represent significant limitations to their massive application for diagnosis purposes [26]. E-noses have become target of research, because of their potential ability of being non-invasive, simple and fast tools for detecting VOCs from the human body [84]. An e-nose is a device that comprises of an array of chemical sensors with different selectivity, a signal-preprocessing unit and a pattern recognition system [79]. The interaction of VOCs with an array of sensors generates a characteristic fingerprint which can then be recognized by comparing it with previously recorded patterns in the recognition system [78][79][84]. Several diseases have already been detected using electronic noses, such as urinary tract infections [136][137], tuberculosis [117] and ashtma [49]. However, despite the advances in e-nose research areas, sensors selectivity to detect VOCs remains a major challenge.

Liquid crystals (LCs) are intermediate phases between solid and liquid states (mesophase) [138], in which the matter has fluid properties like liquids and anisotropic properties like crystals [139]. Nowadays, LCs are well known mainly due to their application in electronic display devices. Due to their optical properties, materials that present LC phases are also attractive for other applications, such as chemical sensors [89][139][140]. One of their most advantageous characteristics is that LC molecules are able to rotate the polarization of light that passes through them [141]. Nowadays, the detection principle of LC-based chemical sensors relies on the disruption of the orientations of thermotropic LC molecules at LC/solid or LC/aqueous interfaces upon interaction with analytes [142]. This orientational change of the LC molecules can be observed in a microscope using crossed polarizers due to the LCs birefringence [142]. LCs are promising materials for VOCs detection due to high sensitivity to changes in their molecular ordering under external influences. Several LC based sensors have been sucessfully tested. For example, Ding *et al.* [142] tested a LC based optical sensor to detect butylamine in the air, Winterbottom and colleagues [140] have also used cholesteric LCs to detect ethanol, water vapour, and vaporous analytes such as amines in air and Sen *et al.* [143] have reported a LC based sensor to selectively detect nitrogen dioxide.

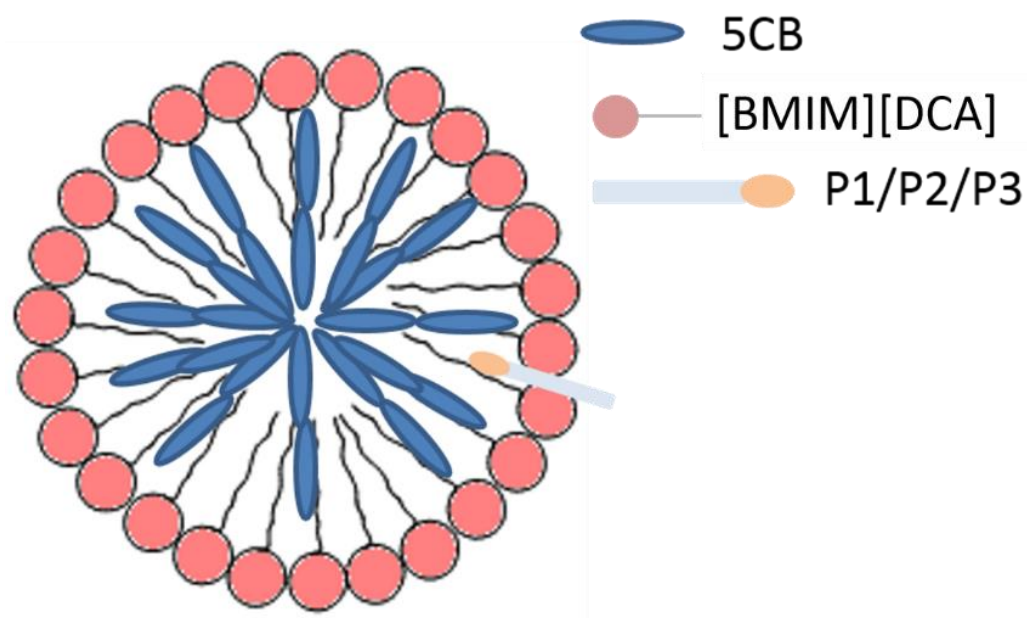
Ionic liquids (ILs) are organic salts with a stable liquid phase over a wide temperature range around room temperature, whose cations and anions can be varied at will to change their

chemical and physical properties. ILs have drawn attention to polymer chemistry, as well as extraction processes of volatile organic solvents worldwide, since they offer a potentially clean method to carry out chemical processes [144] [145]. ILs can act as surfactants [146] [147]. By taking advantage of this property, a proprietary composite material was developed at the Biomolecular Engineering Lab. It is possible to obtain micelles with LC inside and IL at the surface (Figure 4.1). By using a biopolymer and water the micelles are immobilized in the biopolimeric matrix, forming a thin film [148] [149]. When exposed to VOCs, the LC molecules change their organization [150] [151] that can be detected by Polarizing Optical Microscopy (POM) [152].

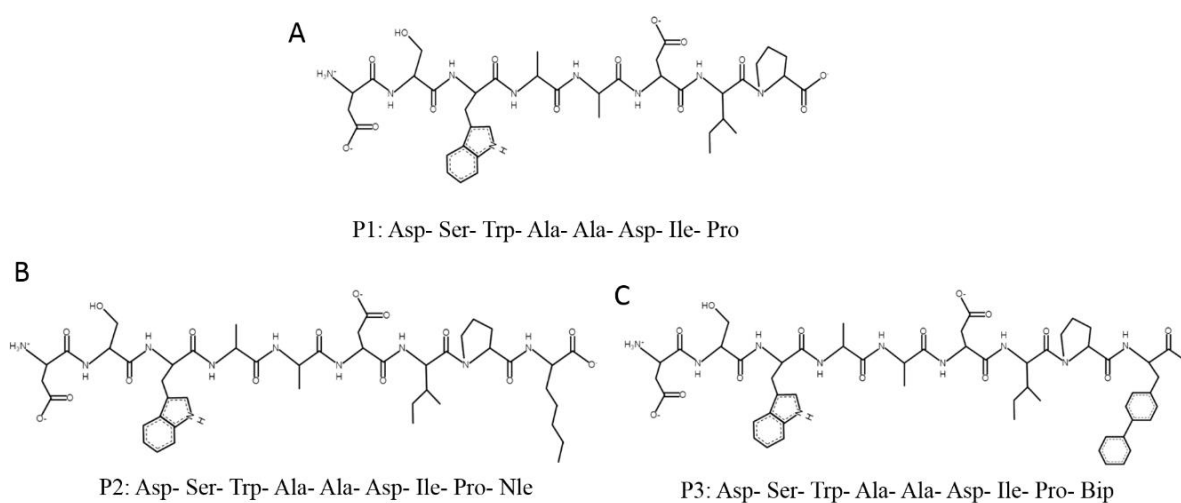
In this work we used the proprietary composite gel-like material composed by the LC 5CB, the IL [BMIM][DCA] and the biopolymer gelatin. This material is able to form thin films responsive to VOCs. The films, here referred as biogels, produce an optical response in presence of VOC molecules due to the change of conformation of 5CB molecules within 5CB-[BMIM][DCA] micelle structures. The optical response is observable by POM and quantified using an e-nose developed in-house. The e-nose consists of an array of sensing elements (each composed by a LED, a biogel thin film placed between two crossed polarizers and a LDR), and a signal processing module that quantifies optical response of the biogels.

Some VOCs are recognized as disease biomarkers. Therefore, tailoring the selectivity of the e-nose response towards certain VOC biomarkers would benefit its usability in disease detection. Ju *et al.* [153] demonstrated that tailor-made small peptides can be promising specific receptors for VOCs detection. In their study they identified a specific peptide, with the aminoacid sequence DSWAADIP (Figure 4.2 A), that showed selectivity towards benzene over toluene, xylene, hexane, acetone and ethanol. The behavior of this peptide could provide a very useful foundation for qualitative and quantitative sensing of VOCs for future applications, such as non-invasive testing of health conditions or environmental risk monitoring.

In this work, we have accessed the feasibility of adding VOC-selectivity to biogels thin films by incorporating in the standard biogel the benzene-sensitive peptide identified by Ju *et al.* [153] and two modified versions of it, that contained norleucine (P2) or biphenylalanine (P3) added to their C-terminal (Figure 4.2 B and C), to facilitate its entry into the micelles. The biogels doped with these modified peptides were observed by POM, the location of the added peptides was verified by FITC labeling and the respective optical response to several VOCs was tested on the in-house developed e-nose.



**Figure 4.1-** Micelles with LC molecules inside, IL at the surface and simulation of the peptides proposed interaction with the micelles.

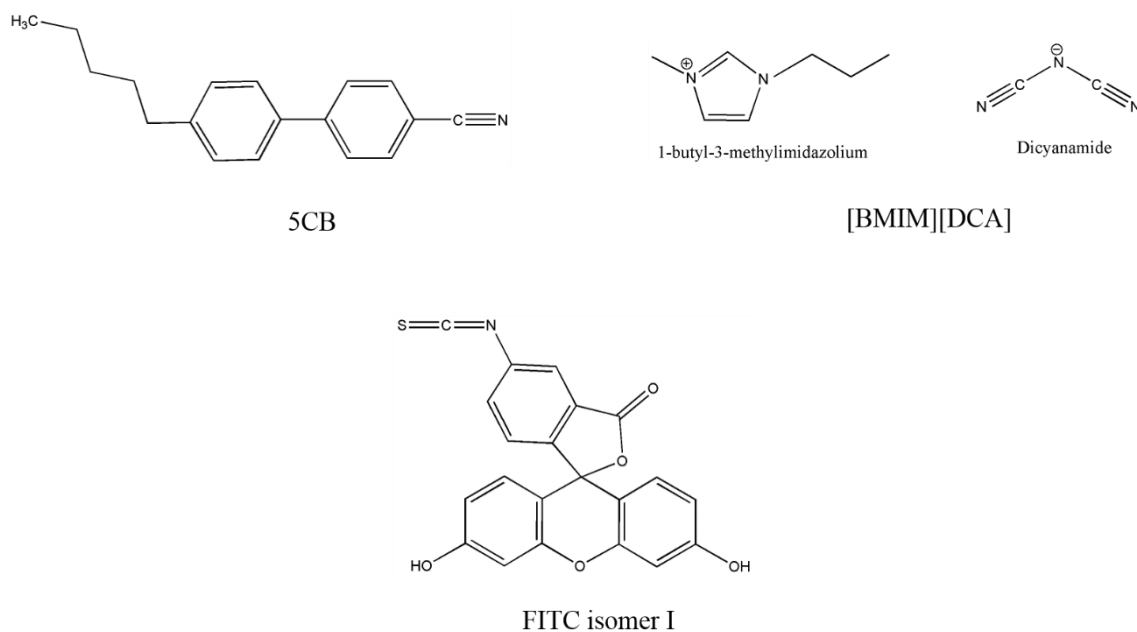


**Figure 4.2-** Structure of peptides P1 (A), P2 (B) and P3 (C).

## 4.2 Materials and methods

### 4.2.1 Materials

**Materials:** 4-cyano-4-pentylbiphenyl (5CB, >98% purity, purchased from TCI) (Figure 4.3), 1-butyl-3-methylimidazolium dicyanamide ([BMIM][DCA], >98% purity, purchased from io-littec) (Figure 4.3), Ammoniumperoxodisulfate (APS,  $\geq 98\%$  purity, purchased from Roth), Tetramethylethylenediamine (TEMED,  $\geq 98\%$ , purchased from nzytech), acrylamide/bis-acrylamide solution, Fluorescein isothiocyanate (FITC) Isomer I (99% purity, purchased from Sigma) (Figure 4.3), agarose (ultrapure grade, from nzytech), gelatin from bovine skin (purchased from Sigma), acetone (>99.5% purity, purchased from Roth), toluene (>99.5% purity, purchased from Panreac), benzene (>98% purity, purchased from Quimilabo), xylene (>98% purity, purchased from Riedel de-Haen), hexane (95% purity, purchased from PA Fisher), ethanol (96% purity, purchased from Panreac), peptide P1 (Asp- Ser- Trp- Ala- Ala- Asp- Ile- Pro, 98.9% purity, purchased from GeneCust), peptide P2 (Asp- Ser- Trp- Ala- Ala- Asp- Ile- Pro- Nle, 98.2% purity, purchased from GeneCust), peptide P3 (Asp- Ser- Trp- Ala- Ala- Asp- Ile- Pro- Bip, 99.3% purity, purchased from GeneCust).



**Figure 4.3** - Structures of 4-cyano-4-pentylbiphenyl (5CB), 1-butyl-3-methylimidazolium dicyanamide ([BMIM][DCA]) and Fluorescein isothiocyanate (FITC) Isomer I.

## 4.2.2 Methods

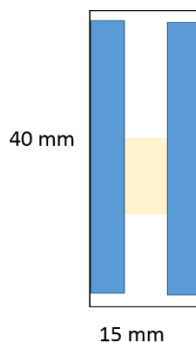
### 4.2.2.1 Production of gelatin biogels

For the production of biogels with liquid crystal-ionic liquid micelles immobilized in gelatin a procedure developed at FCT/UNL by the research group of Prof. Dr. Ana Cecília Roque was followed. A glass vial containing a small magnetic stirrer was pre-heated to 37°C in a stirring hotplate (Echotherm <sup>TM</sup> HS40, Torrey Pines Scientific®) 75 µl of [BMIM][DCA] were introduced in the vial and stirred at 340 rpm for 5 min, after which, 5 µl of 5CB were added and kept stirring for 10 min. Then, 25 mg of gelatin were slowly added and kept stirring at 600 rpm for 10 min to obtain a homogeneous mixture. Finally, 25 µl of milliQ water at 37°C were added and the mixture was kept stirring for 10 min. Negative controls (G<sub>0</sub>, G<sub>1</sub>, G<sub>2</sub>) were also produced (Table 4.1).

A drop (45 µl) of the mixtures was pipetted into warm (37°C) glass slides with adhesive tape on the sides (Figure 4.4), spread with a glass rod and the film was left to dry.

**Table 4.1-** Reagents and corresponding quantities used to produce biogels (G<sub>3</sub>) and the respective negative controls (G<sub>0</sub>, G<sub>1</sub>, G<sub>2</sub>). G<sub>0</sub>- gelatin + milliQ water; G<sub>1</sub>- [BMIM][DCA] + gelatin + milliQ water; G<sub>2</sub>- 5CB + gelatin + milliQ water; G<sub>3</sub>- [BMIM][DCA]+ 5CB + gelatin + milliQ water.

	[BMIM][DCA]	5CB	gelatin	milliQ water
<b>G<sub>0</sub></b>	0 µl	0 µl	25 mg	105 µl
<b>G<sub>1</sub></b>	75 µl	0 µl	25 mg	30 µl
<b>G<sub>2</sub></b>	0 µl	5 µl	25 mg	75 µl + 25 µl
<b>G<sub>3</sub></b>	75 µl	5 µl	25 mg	25 µl



**Figure 4.4-** Representation of the glass plate, after spreading the gel.

#### 4.2.2.2 Production of agarose biogels

To test agarose as a support matrix to the 5CB/[BMIM][DCA] micelles, a similar protocol was followed. Namely, a hotplate was heated to 45°C and a small magnetic stirrer was placed in a glass vial. Then, 75 µl of [BMIM][DCA] was introduced in the vial and stirred at 340 rpm for 5 min, 5 µl of 5CB was added to the mixture and was kept stirring for 10 min. After that, 84.5 µl of 5% agarose were added and kept stirring at 600 rpm for 10 min. Negative controls were also produced (A<sub>0</sub>, A<sub>1</sub>, A<sub>3</sub>) (Table 4.2). A drop (45 µl) of the mixtures was pipetted into warm (45°C) glass slides with adhesive tape on the sides (Figure 4.4), spread with a glass rod and the film was left to dry.

**Table 4.2-** Reagents and corresponding quantities used to produce agarose biogels (A<sub>3</sub>), and the respective negative controls (A<sub>0</sub>, A<sub>1</sub>, A<sub>2</sub>). A<sub>0</sub>- agarose + milliQ water; A<sub>1</sub>- [BMIM][DCA] + agarose + milliQ water; A<sub>2</sub>- 5CB + agarose + milliQ water; A<sub>3</sub>- [BMIM][DCA]+ 5CB + agarose + milliQ water.

	[BMIM][DCA]	5CB	agarose	milliQ water
A <sub>0</sub>	0 µl	0 µl	84.5 µl	80 µl
A <sub>1</sub>	75 µl	0 µl	84.5 µl	5 µl
A <sub>2</sub>	0 µl	5 µl	84.5 µl	75 µl
A <sub>3</sub>	75 µl	5 µl	84.5 µl	0 µl

#### 4.2.2.3 Production of polyacrylamide biogels

An acrylamide/bisacrylamide solution was prepared by adding 1.67 ml of acrylamide/bisacrylamide solution to 19 µl of PSA and 1.25 µl TEMED. A glass vial containing a small magnetic stirrer was pre-heated to 45°C in a stirring hotplate (Echotherm™ HS40, Torrey Pines Scientific®). Then, 75 µl of [BMIM][DCA] was introduced in the vial and stirred at 340 rpm for 5 min, after which, 5 µl of 5CB were added and kept stirring for 10 min. Then, 84.5 µl of the previously prepared polyacrylamide were added and kept stirring at 600 rpm for 10 min. Negative controls were also produced (P<sub>0</sub>, P<sub>1</sub>, P<sub>2</sub>) (Table 4.3). A drop (45 µl) of the mixtures was pipetted into warm (45°C) glass slides with adhesive tape on the sides (Figure 4.4), spread with a glass rod and the resulting film was left to dry.



**Table 4.3-** Reagents and corresponding quantities used to produce biogels with polyacrylamide (P<sub>3</sub>), and the respective negative controls (P<sub>0</sub>, P<sub>1</sub>, P<sub>2</sub>). P<sub>0</sub>- polyacrylamide + milliQ water; P<sub>1</sub>- [BMIM][DCA] + polyacrylamide + milliQ water; P<sub>2</sub>- 5CB + polyacrylamide + milliQ water; P<sub>3</sub>- [BMIM][DCA]+ 5CB + polyacrylamide + milliQ water.

	[BMIM][DCA]	5CB	polyacrylamide	milliQ water
<b>P<sub>0</sub></b>	0 µl	0 µl	84.5 µl	80 µl
<b>P<sub>1</sub></b>	75 µl	0 µl	84.5 µl	5 µl
<b>P<sub>2</sub></b>	0 µl	5 µl	84.5 µl	75 µl
<b>P<sub>3</sub></b>	75 µl	5 µl	84.5 µl	0 µl

#### 4.2.2.4 Coomassie staining of the biogels

The biogels produced using agarose and polyacrylamide as support matrices were stained with coomassie blue R-250 solution. For that, the glass slides were immersed in coomassie blue R-250 solution for 30 min, under gentle agitation. Then, the staining solution was discarded and the biogels were washed several times with ddH<sub>2</sub>O. Finally, the glass slides were immersed in destaining solution for 30 min, slowly agitating. After this step the destaining solution was removed and the gels were left to dry in the hotte.

#### 4.2.2.5 Labeling of peptide P1 with FITC

For FITC labeling, 30 mg/ml stock solution of peptide P1 in carbonate-bicarbonate buffer (0.1 M pH = 9.0) was prepared and 0.44 mg of FITC were diluted in 1 ml of DMSO. Then, 300 µl of the FITC solution were added, in aliquots of 10 µl, slowly while agitating, to 100 µl of P1 solution. The final mixture was left to incubate in the dark overnight. P1-FITC conjugate was recovered by HPLC. Since the resulting sample was very diluted (0.80 mg/ml P1 to 0.30 mg/ml FITC) it was then concentrated (3.46 mg/ml P1 to 1.40 mg/ml FITC) by using a rotary evaporator.

#### 4.2.2.6 Determination of P1 and FITC concentrations in the P1-FITC sample

To determine P1 and FITC concentrations in the P1-FITC sample, two aliquots were taken before and after using the rotary evaporator, respectively, and the absorbance values (280 nm) and fluorescence intensity (485-535 nm) were measured in a microplate reader (infinite 200, Tecan i-control). P1 in carbonate-bicarbonate buffer solutions were prepared, in 5 different concentrations (6 mg/ml, 3 mg/ml, 1.5 mg/ml, 0.75 mg/ml and 0.375 mg/ml) and the respective absorbance values were measured at 280 nm (infinite 200, Tecan i-control). FITC in DMSO

solutions were prepared in 6 distinct concentrations (13 mg/ml, 2.6 mg/ml, 1.3 mg/ml, 0.65 mg/ml, 0.325 mg/ml and 0.1625 mg/ml) and the fluorescence intensity was measured at 485-535 nm in a microplate reader (infinite 200, Tecan i-control). Two calibration lines were constructed and the concentrations of P1 and FITC, for the two P1-FITC aliquots, were calculated.

#### 4.2.2.7 Incorporation of P1-FITC in a gelatin biogel

The gelatin biogel was produced following the protocol detailed in 4.2.2.1 with slight modifications: P1-FITC solution (as obtained after the evaporation of excess solvent) was added to the mixture before the addition of the gelatin, and no water was added (Table 4.4).

**Table 4.4-** Reagents and corresponding quantities used to incorporate the peptides P1 labeled with FITC in the liquid crystal-ionic liquid micelles, and the respective negative control. F<sub>0</sub>- [BMIM][DCA] + 5CB + gelatina + milliQ water; F<sub>1</sub>- [BMIM][DCA] + 5CB + P1-FITC + gelatin.

	[BMIM][DCA]	5CB	gelatin	P1-FITC	milliQ water
<b>F<sub>0</sub></b>	75 µl	5 µl	25 mg	0 µl	25 µl
<b>F<sub>1</sub></b>	75 µl	5 µl	25 mg	25 µl	0 µl

#### 4.2.2.8 Incorporation of peptides P1, P2 and P3 in gelatin biogels

Different concentrations of P1 in water were tested to optimize the protocol. The most successful one is detailed hereafter. First, 75 µl of [BMIM][DCA] were added to a glass vial containing a small magnetic stirrer and pre-heated to 37°C for 5 min with stirring at 340 rpm. Then, 5 µl of 5CB were added and kept stirring for 10 min. After that, 10 µl of a 10 mg/ml P1 solution or 11.3 µl of a 10 mg/ml P2 solution or 12.5 µl of a 10 mg/ml P3 solution were added to the mixture and kept stirring for 10 min. After that, 25 mg of gelatin were slowly added and kept stirring at 600 rpm for 10 min to obtain an homogeneous mixture. Finally, 25 µl of warm milliQ water were added and the mixture was kept stirring for 10 min. Negative controls (G<sub>0</sub><sup>1</sup>, G<sub>1</sub><sup>1</sup>, G<sub>2</sub><sup>1</sup>, G<sub>0</sub><sup>2</sup>, G<sub>1</sub><sup>2</sup>, G<sub>2</sub><sup>2</sup>, G<sub>0</sub><sup>3</sup>, G<sub>1</sub><sup>3</sup>, G<sub>2</sub><sup>3</sup>) were also produced, for each peptide (Table 4.5).

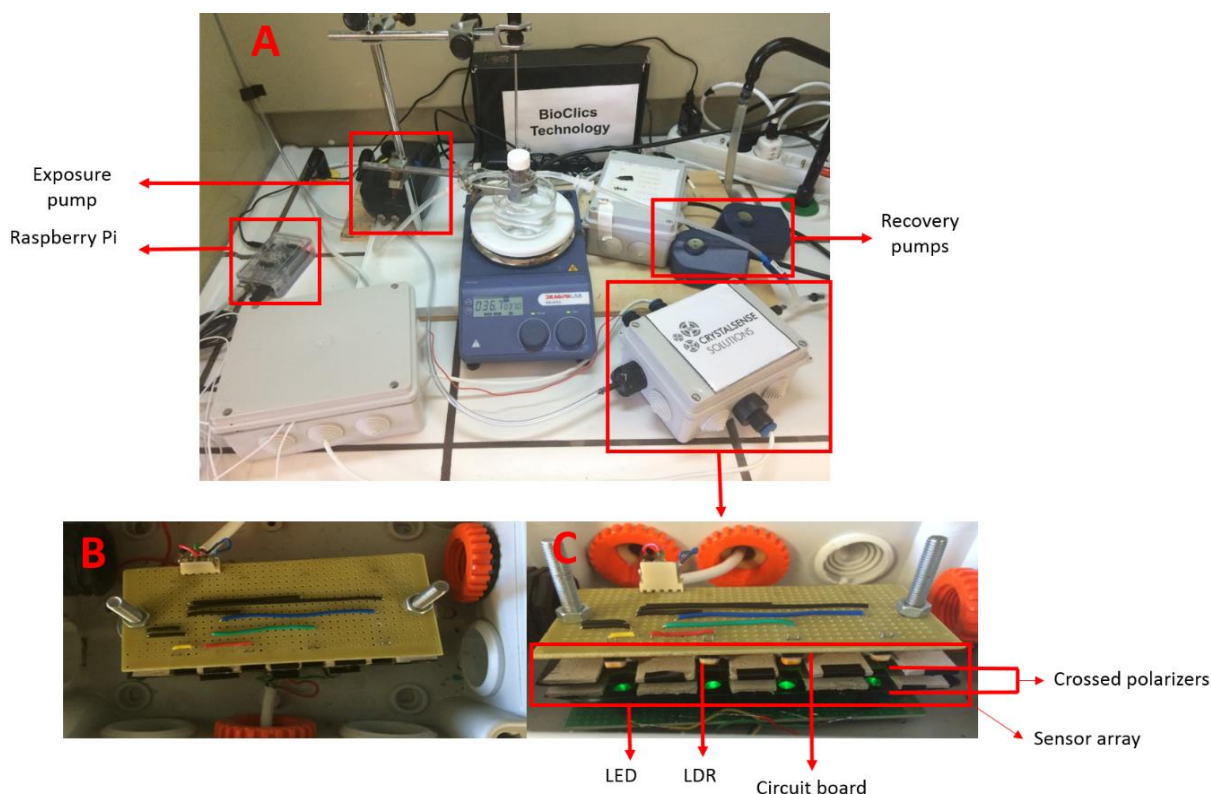
**Table 4.5-** Reagents and corresponding quantities used to incorporate the peptides P1 ( $G_3^1$ ), P2 ( $G_3^2$ ) and P3 ( $G_3^3$ ) in the biogels, and to produce the respective negative controls ( $G_0^1$ ,  $G_1^1$ ,  $G_2^1$ ), ( $G_0^2$ ,  $G_1^2$ ,  $G_2^2$ ) and ( $G_0^3$ ,  $G_1^3$ ,  $G_2^3$ ).  $G_0^1$ - gelatin + P1 + milliQ water;  $G_1^1$ - [BMIM][DCA] + P1 + gelatin + milliQ water;  $G_2^1$ - 5CB + P1 + gelatin + milliQ water;  $G_3^1$ - [BMIM][DCA]+ 5CB + P1 + gelatin + milliQ water.  $G_0^2$ - gelatin + P2 + milliQ water;  $G_1^2$ - [BMIM][DCA] + P2 + gelatin + milliQ water;  $G_2^2$ - 5CB + P2 + gelatin + milliQ water;  $G_3^2$ - [BMIM][DCA]+ 5CB + P2 + gelatin + milliQ water.  $G_0^3$ - gelatin + P3 + milliQ water;  $G_1^3$ - [BMIM][DCA] + P3 + gelatin + milliQ water;  $G_2^3$ - 5CB + P3 + gelatin + milliQ water;  $G_3^3$ - [BMIM][DCA]+ 5CB + P3 + gelatin + milliQ water.

		[BMIM][DCA]	5CB	gelatin	Peptide	milliQ water
<b>Biogels with P1</b>	<b><math>G_0^1</math></b>	0 $\mu$ l	0 $\mu$ l	25 mg	10 $\mu$ l	75+15 $\mu$ l
	<b><math>G_1^1</math></b>	75 $\mu$ l	0 $\mu$ l	25 mg	10 $\mu$ l	15 $\mu$ l
	<b><math>G_2^1</math></b>	0 $\mu$ l	5 $\mu$ l	25 mg	10 $\mu$ l	75+15 $\mu$ l
	<b><math>G_3^1</math></b>	75 $\mu$ l	5 $\mu$ l	25 mg	10 $\mu$ l	15 $\mu$ l
<b>Biogels with P2</b>	<b><math>G_0^2</math></b>	0 $\mu$ l	0 $\mu$ l	25 mg	11.3 $\mu$ l	75+13.7 $\mu$ l
	<b><math>G_1^2</math></b>	75 $\mu$ l	0 $\mu$ l	25 mg	11.3 $\mu$ l	13.7 $\mu$ l
	<b><math>G_2^2</math></b>	0 $\mu$ l	5 $\mu$ l	25 mg	11.3 $\mu$ l	75+13.7 $\mu$ l
	<b><math>G_3^2</math></b>	75 $\mu$ l	5 $\mu$ l	25 mg	11.3 $\mu$ l	13.7 $\mu$ l
<b>Biogels with P3</b>	<b><math>G_0^3</math></b>	0 $\mu$ l	0 $\mu$ l	25 mg	12.5 $\mu$ l	75+12.5 $\mu$ l
	<b><math>G_1^3</math></b>	75 $\mu$ l	0 $\mu$ l	25 mg	12.5 $\mu$ l	12.5 $\mu$ l
	<b><math>G_2^3</math></b>	0 $\mu$ l	5 $\mu$ l	25 mg	12.5 $\mu$ l	75+12.5 $\mu$ l
	<b><math>G_3^3</math></b>	75 $\mu$ l	5 $\mu$ l	25 mg	12.5 $\mu$ l	12.5 $\mu$ l

#### 4.2.2.9 Evaluation of the optical response of gelatin biogels in the presence of different VOCs

The different gelatin biogel films ([BMIM][DCA] + 5CB + milliQ water + gelatin, [BMIM][DCA] + 5CB + P1+ milliQ water + gelatin, [BMIM][DCA] + 5CB + P2 + milliQ water + gelatin, [BMIM][DCA] + 5CB + P3 + milliQ water + gelatin, and the correspondent controls) were positioned in the sensor array of the in-house developed e-nose (Figure 4.5) and tested regarding their responses to acetone, hexane, benzene, xylene, toluene and ethanol vapours. For all the solvents the biogels were subjected to 5 cycles of exposure/recovery (60 s of exposure followed by 100 s of recovery, totalizing 15 min per test). The solvents were kept in a bath thermostated at 37°C and the respective vapours were pumped to the sensors array chamber (during the exposure time) alterned with clean air (during the recovery time).

The optical signal intensities were calculated by determining the signal amplitude corresponding to each biogel after being exposed to the solvents. The response fold-increase was calculated by the ratio of the biogels with P1, P2 or P3 and the standard biogel signal intensities.



**Figure 4.5-** In-house developed e-nose assembly, when exposing a biogel to a solvent (A). E-nose sensor array, seen from above (B) and from the side (C). LED- Light Emitting Diode. LDR- Light Dependent Resistor.

#### 4.2.2.10 Optical characterization of the biogels by Microscopy

To characterize the biogels regarding the formation of LC/IL micelles and the respective morphology, the glass slides containing the biogels were observed under polarized light using a ZEISS, Observer.Z1 optical microscope equipped with a ZEISS, AxioCam 503 color camera. To observe the biogels containing P1-FITC, the same microscope was employed, using a source of UV light and a green fluorescent filter. The imaging software for microscopy (ZEN 2.3 (blue edition), ZEISS) was used to process and analyze the images.

### 4.3 Results and discussion

#### 4.3.1 Doping gelatin biogels with P1, P2 and P3 peptides

As shown by Ju *et al.* the peptide Asp- Ser- Trp- Ala- Ala- Asp- Ile- Pro (P1) has the ability to recognize benzene over other volatile molecules [153]. As a case-study for tailoring the selectivity of the biogels, new peptide-containing biogels were produced by incorporating this peptide or its analogs P2 and P3 in the composition of the biogels. The resulting biogels were then evaluated regarding their optical response towards different volatile solvents to access the effect of adding a selective component in the material.

The proprietary composite gel-like material composed by the LC 5CB, the IL [BMIM][DCA] and te biopolymer gelatin allow the formation of optically active micelles with LC inside and IL at the surface, observable by POM (Figure 4.6).

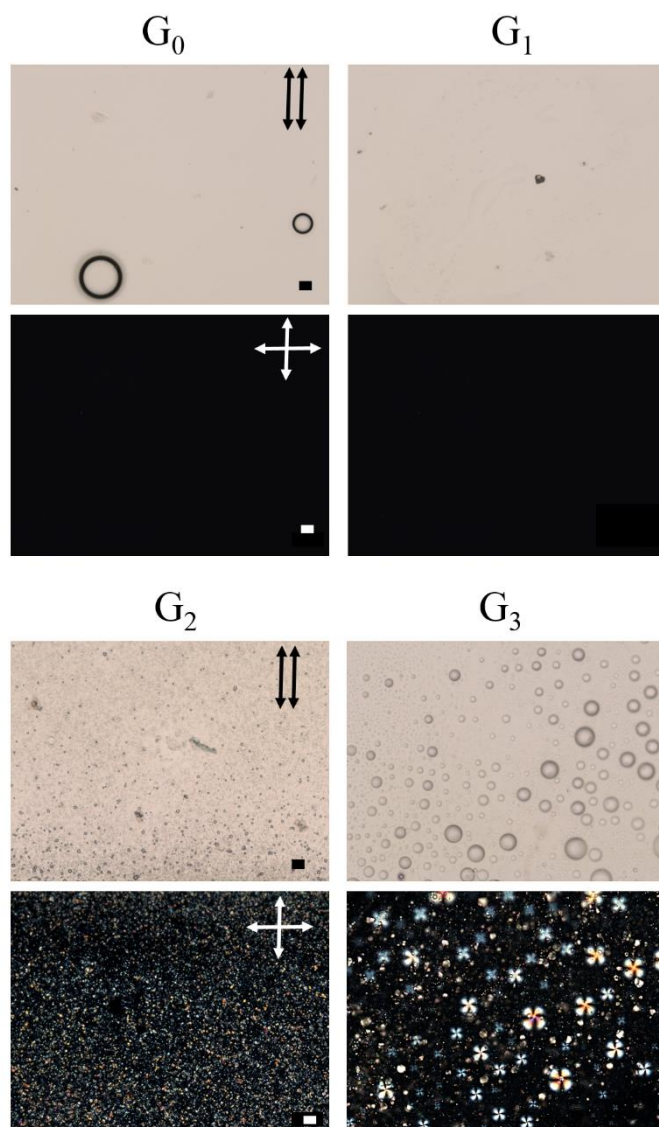
Controls without the addition of the peptides were produced (Figure 4.7 and Appendix 1) and we verified that micelles were only formed in a mixture containing [BMIM][DCA], 5CB and water (Figure 4.8). The IL acts as a surfactant and, since LC molecules are hydrophobic, occurs the self-assembly of 5CB/[BMIM][DCA] in micellar structures with radial configuration [154]. This configuration has one point of defect at the micelle center, causing the micelles to appear as crosses, when observed by POM with crossed-polarizers [154] [155].

We found that by adding P1 to liquid crystal-ionic liquid mixture, micelles were also formed. However, the negative controls with gelatin, P1, milliQ water and gelatin, [BMIM][DCA], P1, milliQ water presented birefringent structures, which could indicate that the peptide may have LC properties [156] [157] and self-assemble in some structure with birefringent properties (Figure 4.9).

After finding that the incorporation of the peptide P1 in the biogels was possible and did not affect the formation of micelles (See Appendix 1), we proceeded to the incorporation of two similar peptides, containing an artificial aminoacid in the C-terminal (norleucine in P2 and bifenilalanine in P3) that could act as hydrophobic tails in the peptides termination, to favour their entry into the micelles.

The addition of either P2 or P3 to the liquid crystal-ionic liquid mixture also resulted in the formation of micelles (See Appendix 1). Regarding P2 controls, we verified that the 5CB/P2 gels presented droplets and micelles, and that the [BMIM][DCA]/P2 gels contained rod-shaped birefringent structures (Figure 4.10), also possibly due to the molecular rigidity [156], while the P3 controls with only [BMIM][DCA], P3 and water, presented micelles (Figure 4.11). P3 contains biphenylalanine which has a similar structure to 5CB (Figure 4.3), which could lead to these peptide behaving similar to 5CB, under these conditions.

Despite the fact that micelles formed in the presence of the three peptides, the biogel that contained the higher number of micelles per biogel area was the one containing [BMIM][DCA], 5CB and P3 with 23497 micelles and a mean micelle area of approximately  $312 \mu\text{m}^2$  (19.9  $\mu\text{m}$  diameter). However, despite the standard gelatin biogel (without peptide) being the one with the smaller number of micelles formed (992) it was the gel that contained micelles with the highest mean area value ( $1098 \mu\text{m}^2$  and 37.9  $\mu\text{m}$  diameter) (Table 4.6). This finding suggests that the addition of small peptides to the gelatin biogel composition tends to reduce the size and increase the number of micelles.

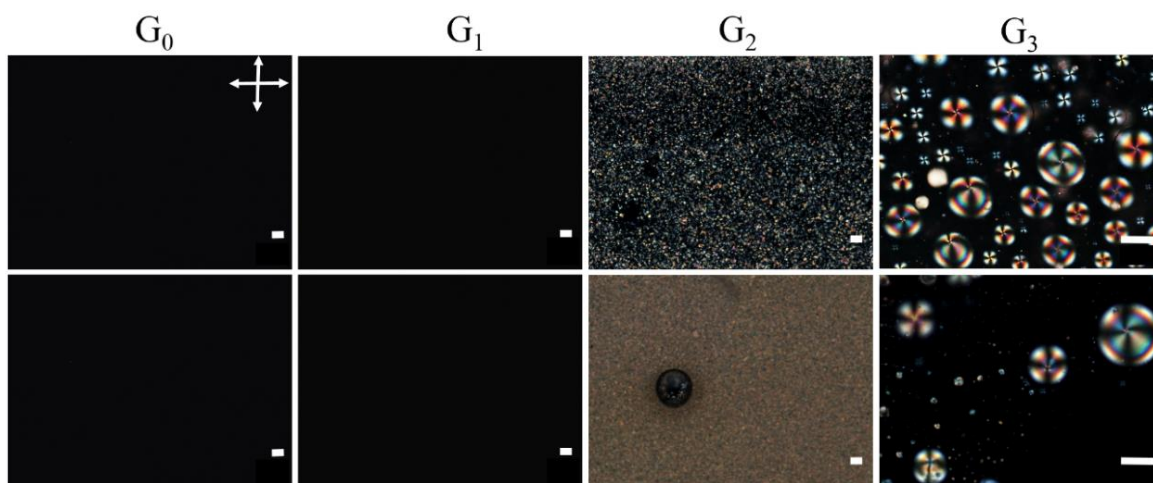


**Figure 4.6-** POM images of standard biogel (G<sub>3</sub>- Gelatin + [BMIM][DCA] + 5CB + milliQ water) and respective controls. G<sub>0</sub>- Gelatin + milliQ water. G<sub>1</sub>- Gelatin + [BMIM][DCA] + milliQ water. G<sub>2</sub>- Gelatin + 5CB + milliQ water. All the images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .

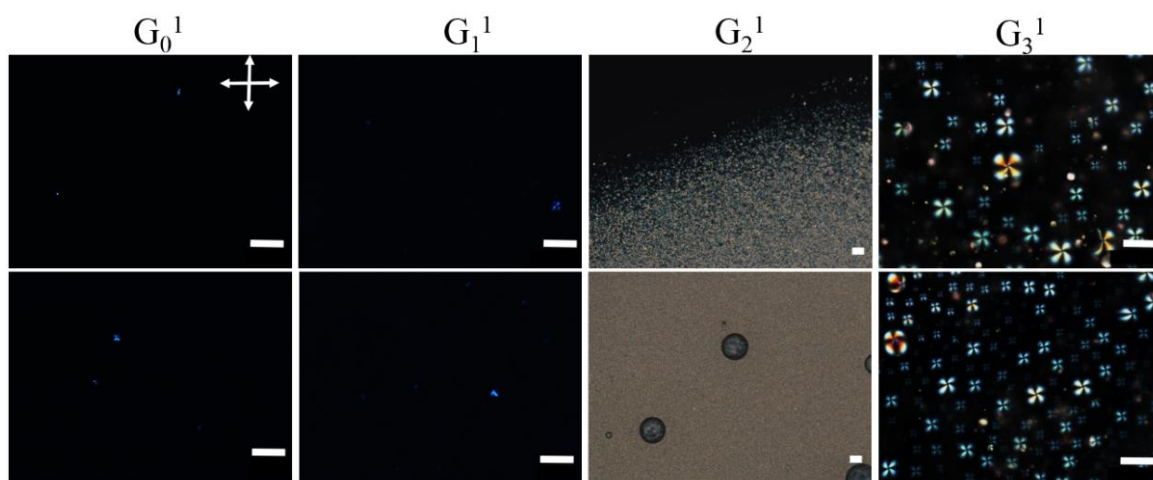




**Figure 4.7-** Visualization of a [BMIM][DCA]/5CB/gelatin biogel, with crossed polarizers. A: visualization of the whole biogel. B- detail of a biogel region.

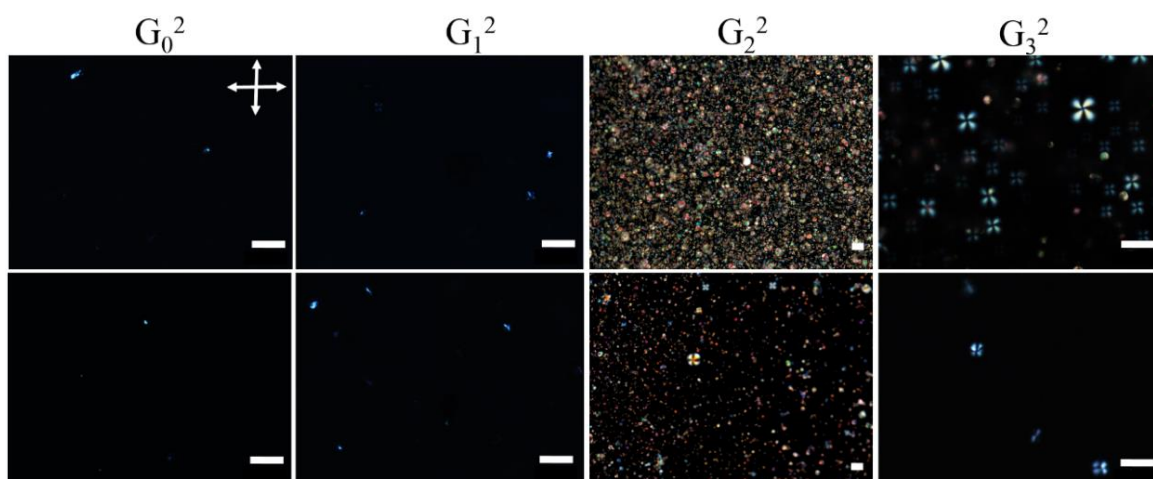


**Figure 4.8-** Standard gelatin biogel and respective controls, observed by POM, with crossed polarizers, in 2 different fields of view.  $G_0$ : Gelatin + milliQ water.  $G_1$ : Gelatin + [BMIM][DCA] + milliQ water.  $G_2$ : Gelatin + 5CB + milliQ water.  $G_3$ : Gelatin + [BMIM][DCA] + 5CB + milliQ water. The scale bar corresponds to 50  $\mu\text{m}$ .

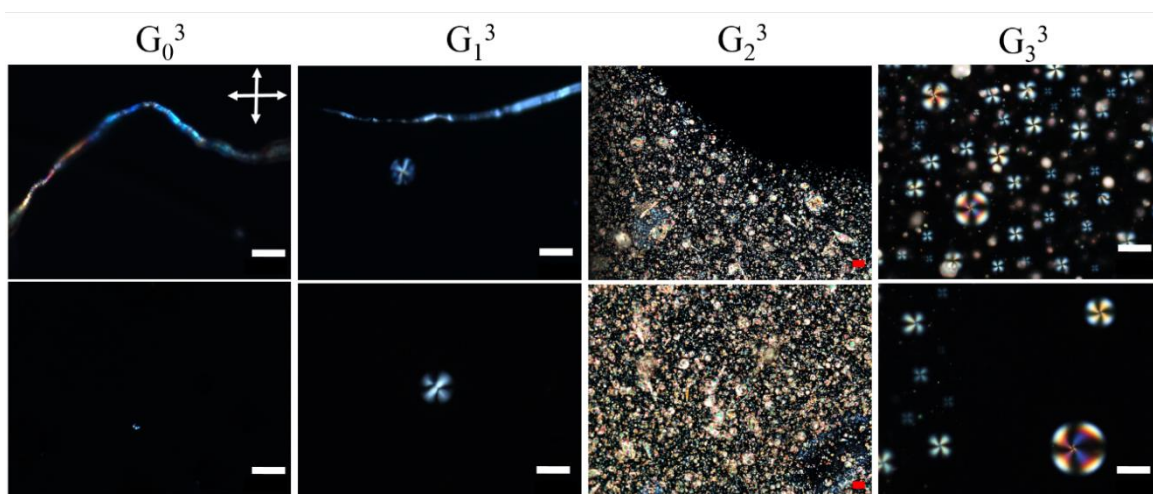


**Figure 4.9-** Gelatin biogels with P1 added, visualized by POM, with crossed polarizers in 2 different fields of view.  $G_0^1$ : Gelatin + P1 + milliQ water.  $G_1^1$ : Gelatin + [BMIM][DCA] + P1 + milliQ water.  $G_2^1$ : Gelatin + 5CB + P1 + milliQ water.  $G_3^1$ : Gelatin + [BMIM][DCA] + 5CB + P1 + milliQ water. The scale bar corresponds to 50  $\mu\text{m}$ .





**Figure 4.10-** Gelatin biogels with P2 added, visualized by POM, with crossed polarizers in 2 different fields of view.  $G_0^2$ : Gelatin + P2 + milliQ water.  $G_1^2$ : Gelatin + [BMIM][DCA] + P2+ milliQ water.  $G_2^2$ : Gelatin + 5CB + P2 + milliQ water.  $G_3^2$ : Gelatin + [BMIM][DCA] + 5CB + P2 + milliQ water. The scale bar corresponds to 50  $\mu\text{m}$ .



**Figure 4.11 -** Gelatin biogels with P3 added, visualized by POM, with crossed polarizers in 2 different fields of view.  $G_0^3$ : Gelatin + P3 + milliQ water.  $G_1^3$ : Gelatin + [BMIM][DCA] + P3+ milliQ water.  $G_2^3$ : Gelatin + 5CB + P3 + milliQ water.  $G_3^3$ : Gelatin + [BMIM][DCA] + 5CB + P3 + milliQ water. The scale bar corresponds to 50  $\mu\text{m}$ .

**Table 4.6** - Biogels with distinct composition and corresponding number of micelles formed and corresponding mean area ( $\mu\text{m}^2$ ).

Biogel composition	Number of micelles	Mean micelle area ( $\mu\text{m}^2$ )	Area with micelles (%)	Biogel area ( $\mu\text{m}^2$ )	Number of micelles per biogel area (micelles/ $\mu\text{m}^2$ )
[BMIM][DCA] + 5CB + milliQ water + gelatin	992	1098.14	2.6	1089350.06	0.0009
[BMIM][DCA] + 5CB + P1 + milliQ water + gelatin	3905	849.50	4.9	3317295.01	0.0012
[BMIM][DCA] + 5CB + P2 + milliQ water + gelatin	1744	833.22	2.3	1453143.60	0.0012
[BMIM][DCA] + 5CB + P3 + milliQ water + gelatin	23497	312.67	8.1	7346790.87	0.0032

### 4.3.2 Tracking peptides location within biogels

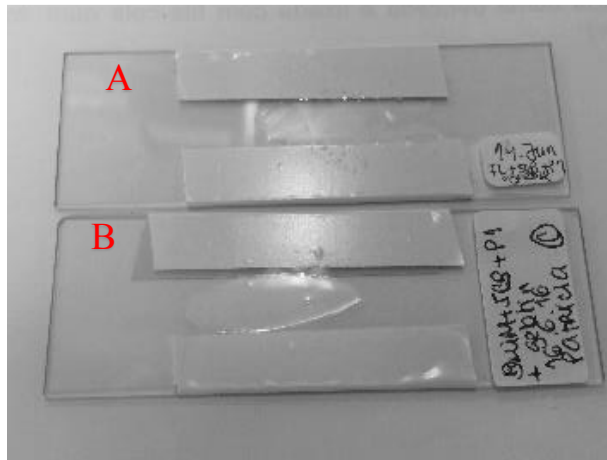
#### 4.3.2.1 Staining with Coomassie blue

To study the morphology of the new peptide-containing biogels we needed a strategy to verify if those peptides were located in the micelles or dispersed in the biogel matrix. Since the aim was to locate the peptides, by specifically staining them, the gelatin component of the biogels was removed, because it is a proteic component and the coomassie blue would also stain it. The gelatin component was substituted by agarose or polyacrylamide, for further coomassie staining of those biogels.

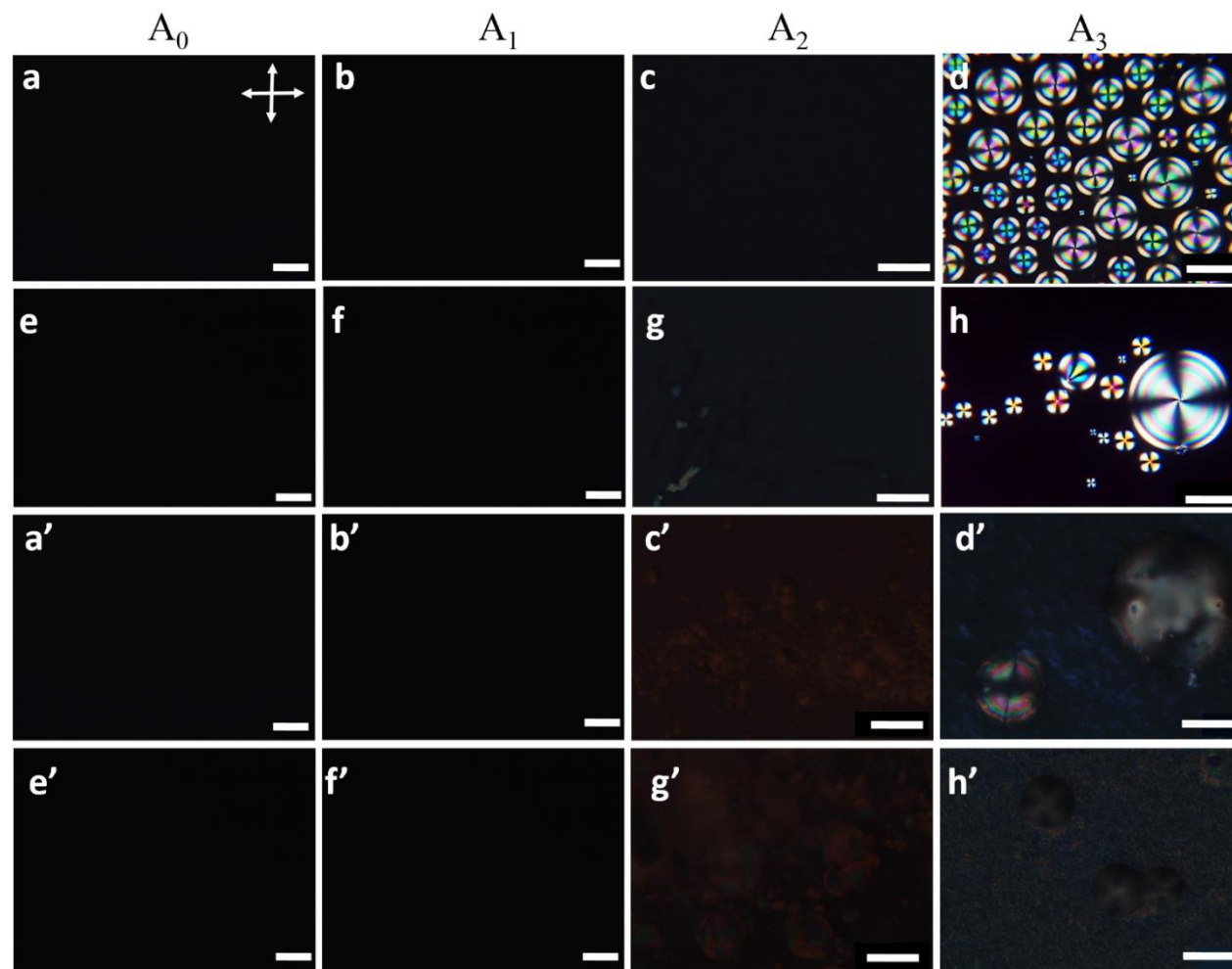
We verified that the biogels obtained with agarose and polyacrylamide in their composition did not have the same consistency of the ones produced with gelatin (Figure 4.12). The gelatin biogel formed more resistant, peelable films while the agarose and polyacrylamide ones resulted in sticky films. By POM we verified that, when using agarose, the liquid crystal-ionic liquid micelles were successfully produced (Figure 4.13 a to h). On the other hand, the polyacrylamide gels did not allow the formation of micellar structures, producing other birefringent structures instead (Figure 4.14 a to h). Curiously, the negative control composed by 5CB, polyacrylamide

and milliQ water by itself produced liquid crystal droplets and some micelles (Figure 4.14 c and g).

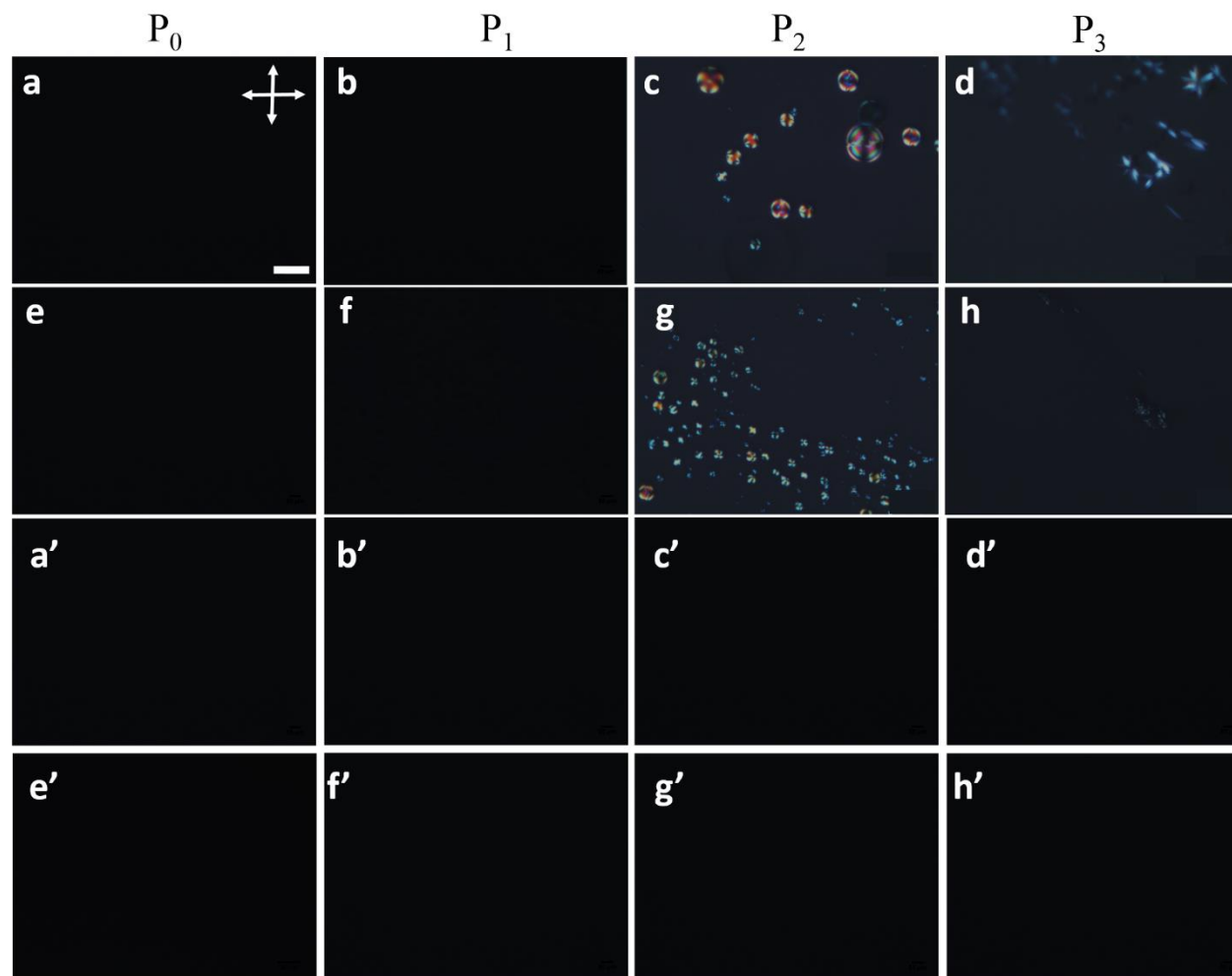
The agarose and the polyacrylamide gels were both stained with Coomassie Blue R-250 and observed by POM. We verified that the staining procedure damaged both biogels and destroyed the micelles in the agarose support (Figures 4.13 and 4.14). In the polyacrylamide gels we did not see any birefringent structures after the staining (Figure 4.14). Micelles formation was only observed in the agarose biogels, however, those gels were also damaged by the staining procedure. Therefore, peptides incorporation was not carried out in agarose and polyacrylamide biogels.



**Figure 4.12** - Appearance of biogels produced using agarose (A) and gelatin (B), as the micelles support.



**Figure 4.13** - Agarose gels visualization by POM, with crossed polarizers, in 2 different fields of view, before (a to h) and after (a' to h') coomassie staining. A<sub>0</sub>- Agarose + milliQ water. A<sub>1</sub>- Agarose + [BMIM][DCA] + milliQ water. A<sub>2</sub>- Agarose + 5CB + milliQ water. A<sub>3</sub>- Agarose + [BMIM][DCA] + 5CB + milliQ water. The scale bar corresponds to 50  $\mu\text{m}$ .

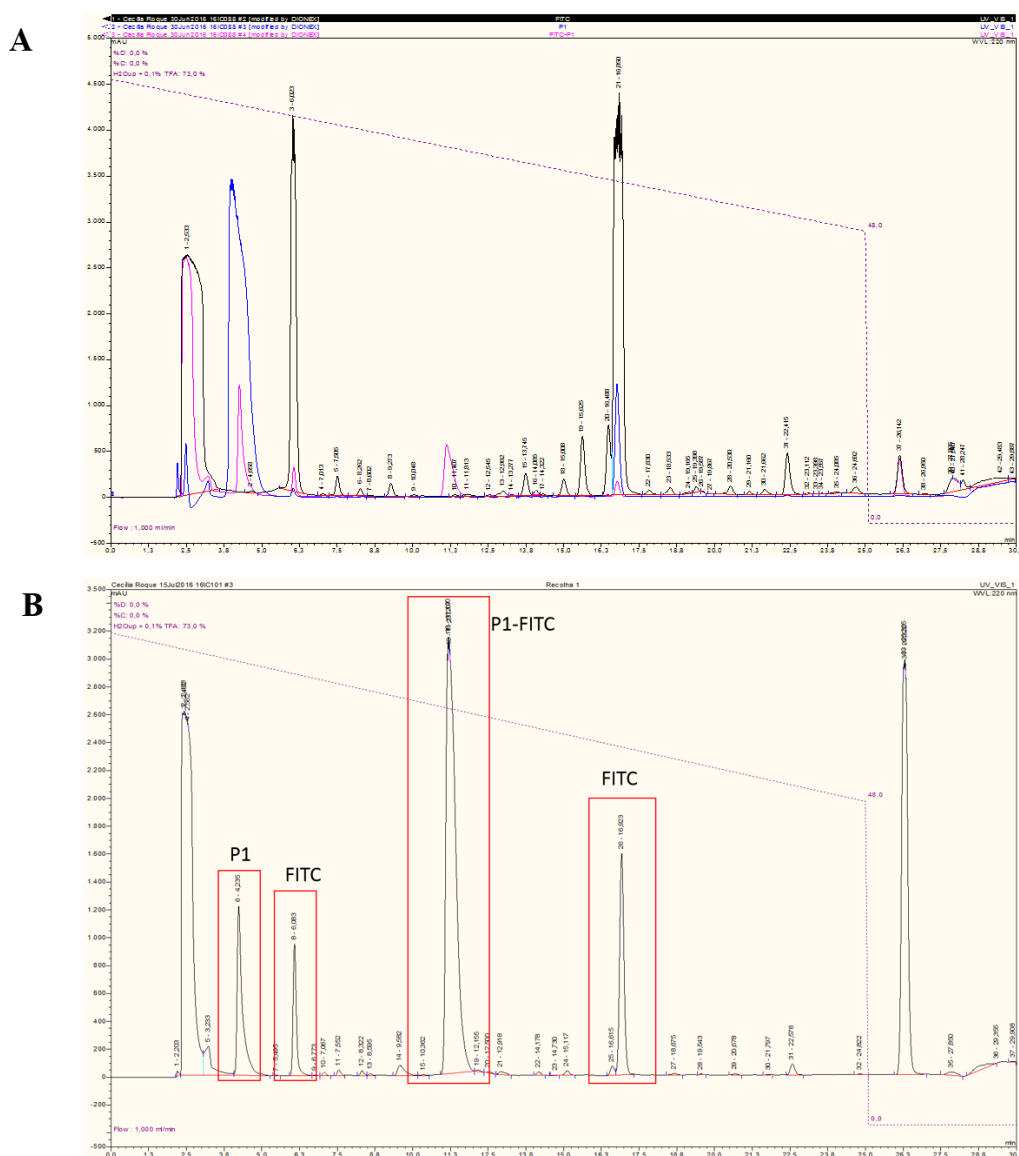


**Figure 4.14** - Polyacrylamide gels visualization by POM, with crossed polarizers, in 2 different fields of view, before coomassie staining (a to h) and after (a' to h'). P<sub>0</sub>- Polyacrylamide + milliQ water. P<sub>1</sub>- Polyacrylamide + [BMIM][DCA] + milliQ water. P<sub>2</sub>- Polyacrylamide + 5CB + milliQ water. P<sub>3</sub>- Polyacrylamide + [BMIM][DCA] + 5CB + milliQ water. All the images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .

### 4.3.2.2 Tracking peptides by fluorescence

Since the staining with coomassie blue R-250 was destructive for the biogels, a different strategy has to be devised for tracking the peptides in the biogels. We applied the method of labeling the peptide P1 with FITC so that it could be detected by observing the biogel by fluorescence microscopy.

After the P1-FITC conjugation reaction, HPLC was performed in order to identify each component and separate P1-FITC conjugate from unlabeled peptide P1 and eventual remains of FITC present in the reaction mixture (Figure 4.15 A and Appendix 1). After identifying each component, the fraction eluted at 11.3-12.5 min, corresponding to the P1-FITC conjugate was collected (Figure 4.15 B).



**Figure 4.15** - Chromatograms obtained for peptide P1, FITC and P1-FITC conjugate samples, overlaid (A) and identification of each fraction (B)

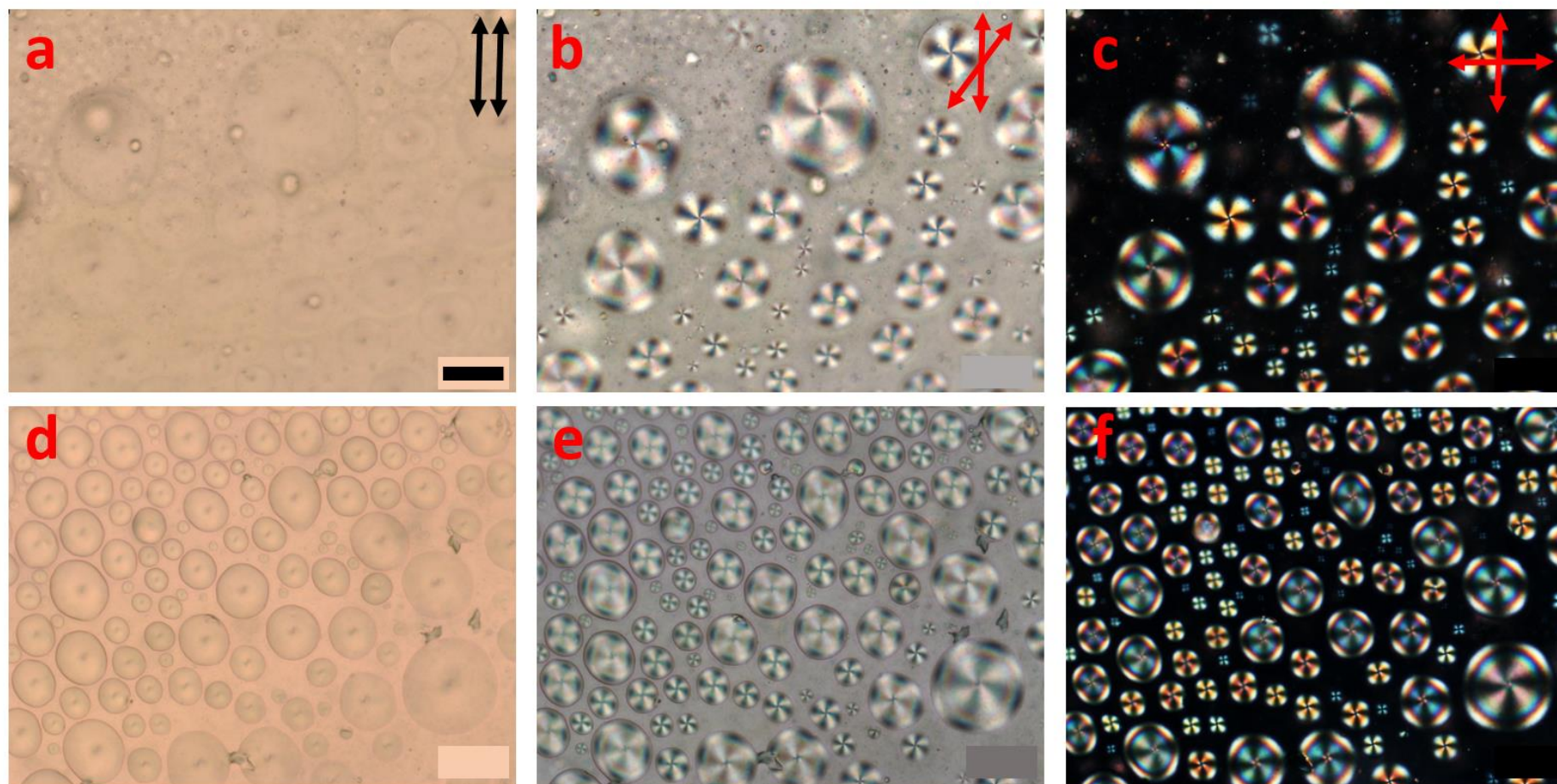
The collected P1-FITC fraction was very diluted, therefore, the rotary evaporator was used to concentrate the sample. After that, P1 and FITC concentrations in the collected fraction were determined. We verified that with the rotary evaporator we were able to concentrate the sample, approximately 4 times. Also, although the FITC and the P1 were added in a 1:1 ratio ( $1.14 \times 10^{-7}$  mol), the final conjugate contained those components in a 1:2.5 ratio, respectively (Table 4.7), which may indicate that some amount of the added FITC did not bind to P1.

**Table 4.7** - Values of absorbance (280 nm) and fluorescence (485-535 nm) for P1 and FITC and the respective concentrations of each compound, in the P1-FITC sample before and after using the rotary evaporator.

	Abs 280 nm	Fluorescence 485-535 nm	Concentration (mg/ml)
<b>P1-FITC sample before rotary evaporator</b>	0.129	3562	P1 = 0.80 FITC = 0.30
<b>P1-FITC sample after rotary evaporator</b>	0.346	6976	P1 = 3.46 FITC = 1.40

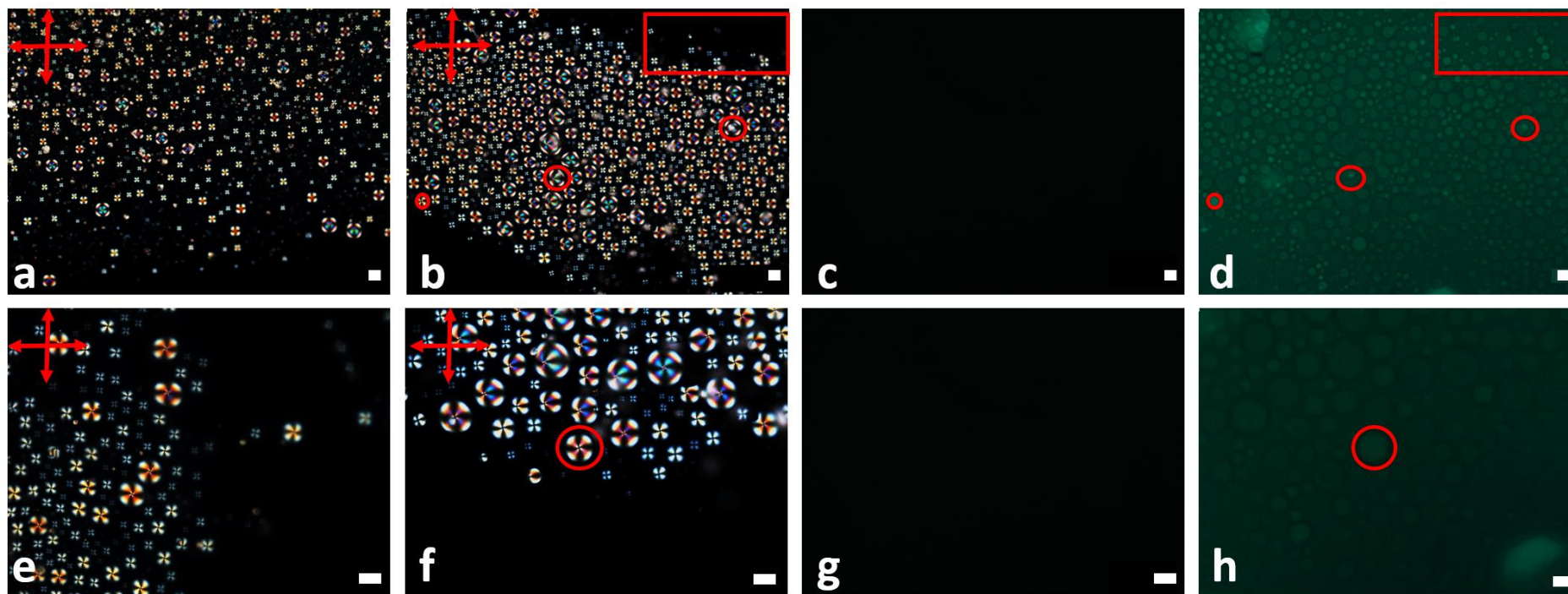
The concentrated P1-FITC sample (3.46 to 1.40 mg/ml) was incorporated in the biogels with gelatin during their production and the resulting films were observed by POM and fluorescence microscopy. POM revealed that in the films containing [BMIM][DCA], 5CB, and gelatin the formation of micelles was not affected by the addition of P1-FITC (Figure 4.16). By observing the biogel with FITC labeled P1 with fluorescence microscopy we concluded that the matrix and the micelles were both green (Figure 4.17, d and h), compared to the control biogel, which did not present any fluorescence (Figure 4.17, c and g), some bubbles also seem to have encapsulated P1-FITC, since they present fluorescence (red rectangle in Figure 4.17). However the inside of the micelles presented a brighter colour than the surrounding matrix, indicating that P1 was successfully incorporated within the micelles and probably a much smaller amount was also dispersed in the gel (red highlights in Figure 4.17). It is assumed that, due to the similarity in the aminoacid sequence, peptides P2 and P3 behave like P1 and will also be partially incorporated into the micelles.





**Figure 4.16** - POM images of gelatin biogels without any peptide added (a to c) and P1-FITC (d to f) with uncrossed (a and d), semi-crossed (b and e) and crossed polarizers (c and f). All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .





**Figure 4.17** - Gelatin biogels with (b and f) and without P1-FITC (a and e) visualized by POM, with crossed polarizers, and by fluorescence microscopy (d, h, and c, g) in 2 different fields of view. a, c, e and g: control biogel composed by gelatin, [BMIM][DCA], 5CB, P1 and milliQ water. b, d, f and h: biogel composed by gelatin, [BMIM][DCA], 5CB, P1-FITC and milliQ water. The scale bar corresponds to 50  $\mu\text{m}$ . The red marks highlight fluorescent bubbles (rectangle) and Bright spots inside the micelles (circles).

### 4.3.3 Effect of P1, P2 and P3 in the response of gelatin biogels to different VOCs

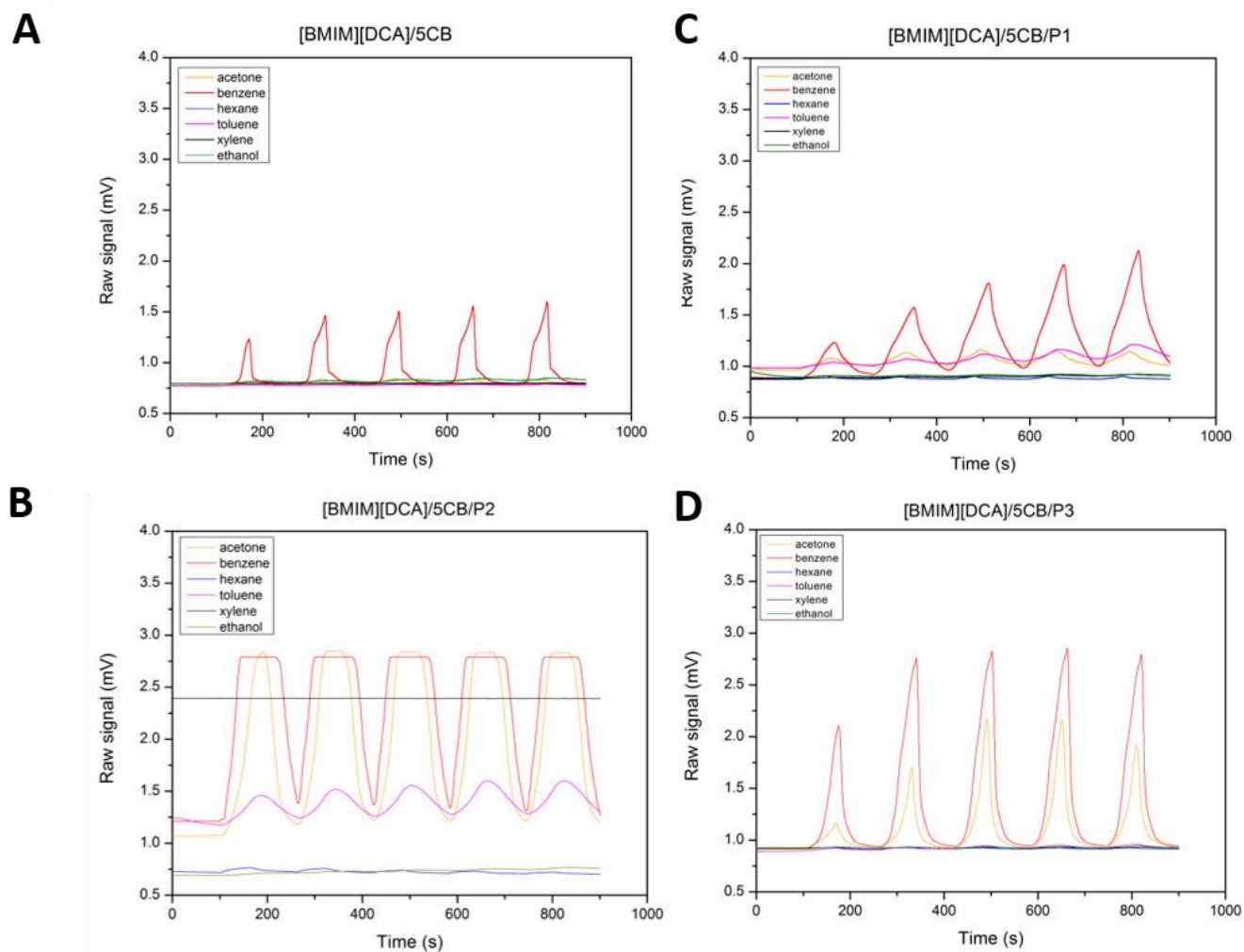
After incorporating the peptides into the biogels, micelles were still produced, therefore, we exposed those materials to a group of solvent vapours in order to evaluate the effect of the peptides in the VOC response of the new materials compared to the standard one. The responses to VOCs were evaluated using a proprietary custom built e-nose. Since we verified that, in some cases, micelles or rod-shaped birefringent structures were formed without the addition of liquid crystal, we also exposed those control films to the same solvents (See Appendix 1).

The solvents to which the biogels were exposed were the same tested in [153] (acetone, benzene, xylene, toluene, hexane and ethanol). These compounds belong to three of the most abundant chemical classes (ketones, hydrocarbons and alcohols) found in Chapter 3 and ethanol was found to be the most frequently detected VOC in pathogen culture headspaces in that same chapter.

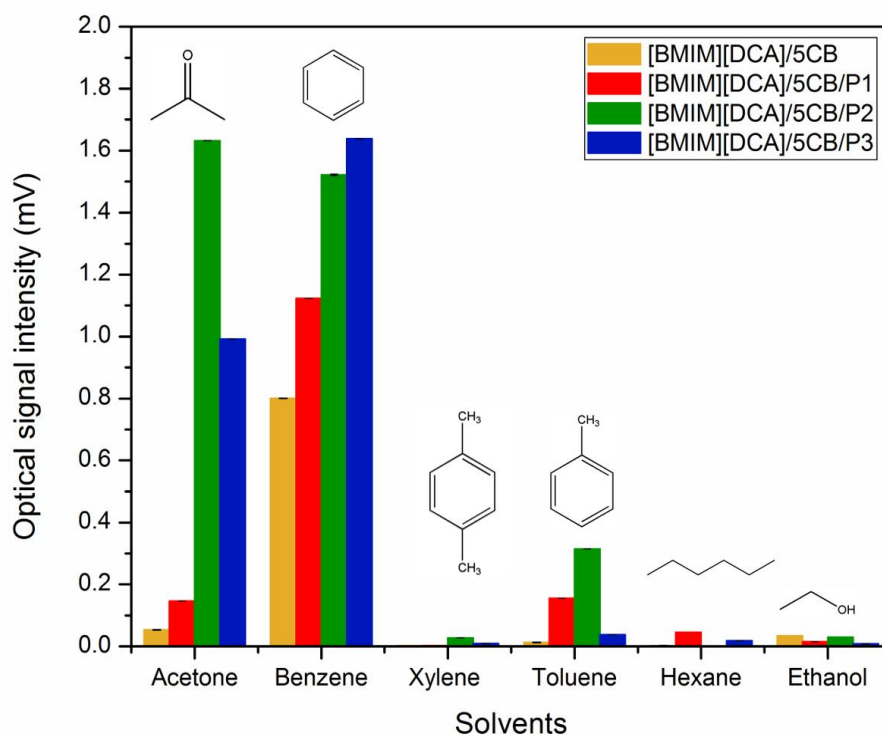
According to Ju *et al.* [153], the peptide P1 presented great selectivity towards benzene, but also presented a minimal response to toluene.

We observed that, although the standard biogel already responded significantly to benzene, compared to other VOCs (Figures 4.18 A and 4.19), the biogel containing P1 presented a much more intense response (Figures 4.18 C and 4.19).

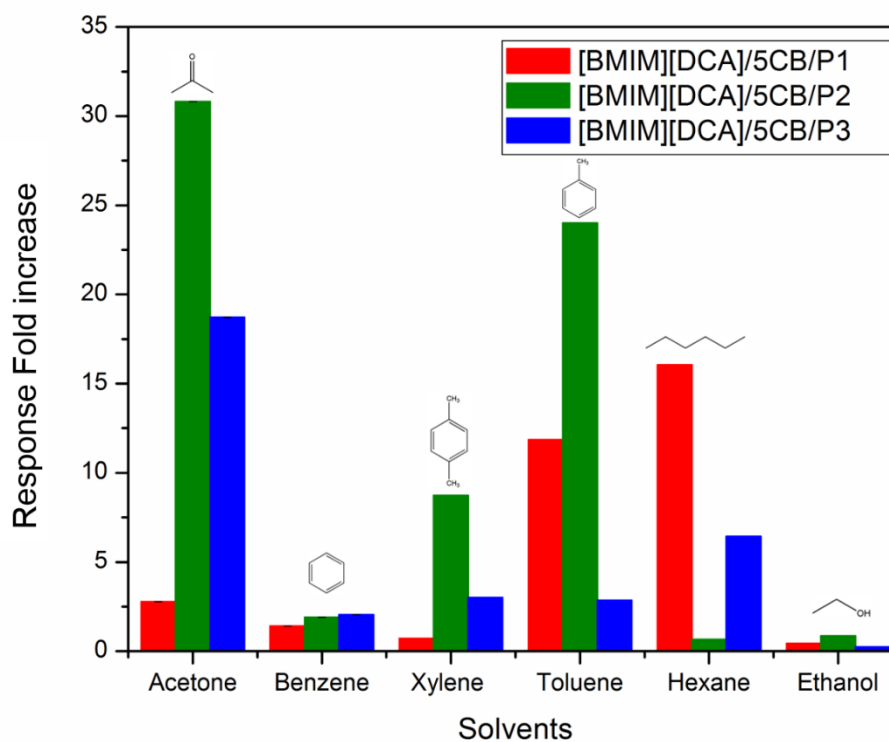
The incorporation of P1 also increased the biogel response to acetone, hexane and toluene, when compared to the material without P1 (Figures 4.18 A and C and Figure 4.19). However, the highest increase occurs for toluene and hexane (Figure 4.20). Biogels containing the modified versions of P1 (P2 and P3) were also tested to see if the addition of norleucine and biphenylalanine, respectively, changed the signal response observed for P1. The incorporation of P2 in the biogels modified the sensors response (Figure 4.19). The sensor responded with a higher signal intensity to benzene, acetone and toluene when P2 was present in the biogel (Figures 4.18 B and 4.19). The greatest response increase occurred in presence of acetone and toluene (Figure 4.20). Regarding the incorporation of P3 we noted that both acetone and benzene produced a pronounced response of the biogel (Figures 4.18 D) and the higher response increase occurs for acetone and hexane (Figure 4.20). In fact, the biogel containing P3 was the one that responded the most when exposed to benzene and the biogel containing P2 responded the most to acetone, xylene, toluene and (Figure 4.19).



**Figure 4.18** -Overlaid signals of the biogels containing [BMIM][DCA]/5CB (A), [BMIM][DCA]/5CB/P1 (C), [BMIM][DCA]/5CB/P2 (B) and [BMIM][DCA]/5CB/P3 (D) when exposed to acetone, benzene, hexane, toluene, xylene and ethanol.



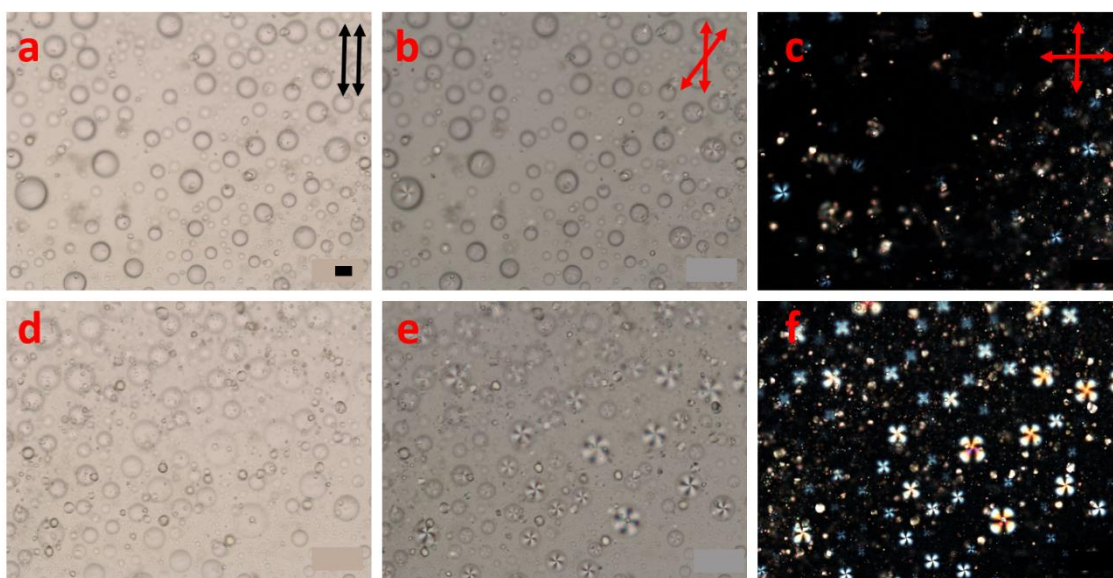
**Figure 4.19** - Optical signal intensity of each biogel to the different solvents.



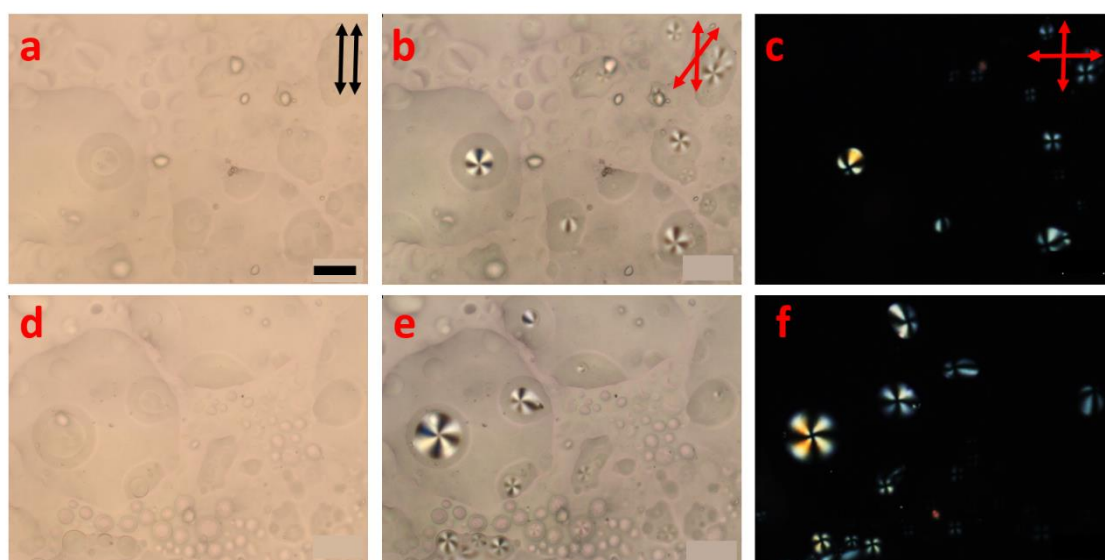
**Figure 4.20** - Response fold increase of each biogel, after P1, P2 and P3 incorporation in the standard biogel.

#### 4.3.4 Biogels stability after VOCs exposure

After VOCs exposure the biogels were observed again by POM. Although some micelles appear to remain stable, all the gels showed that bubbles and some micelles were destroyed (Figures 4.21-4.24). The material containing P3 is the one that remains more stable (Figure 4.24). However, one of the VOCs was an alcohol and the gels stability could be affected after being exposed to it, since it may interfere with the ionic liquid at the micelles surface [157] [156], altering their structure and destabilize the gelatin component [158] [159].

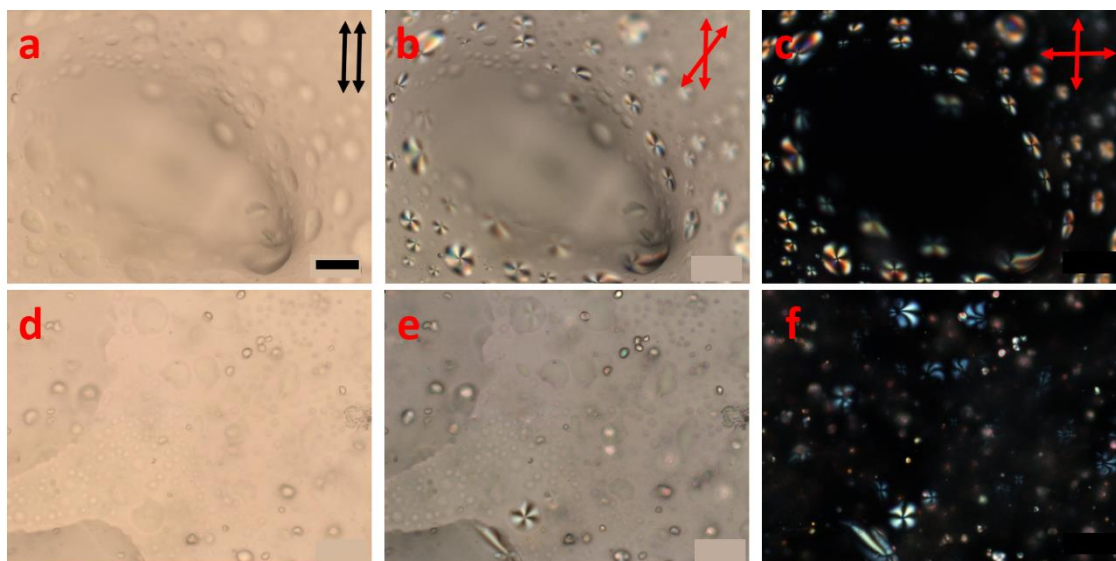


**Figure 4.21** - POM images of standard gelatin biogels (gelatin + [BMIM][DCA] + 5CB), after VOCs exposure, in 2 different fields of view, with uncrossed (a and d), semi-crossed (b and e) and crossed (c and f) polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .

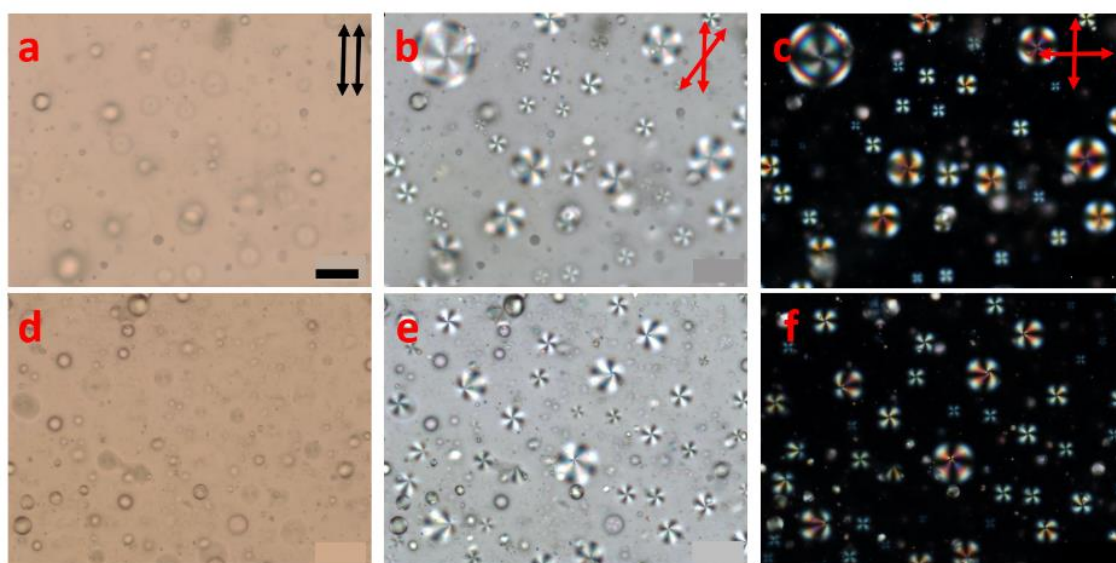


**Figure 4.22** - POM images of P1 biogels (gelatin + [BMIM][DCA] + 5CB + P1), after VOCs exposure, in 2 different fields of view, with uncrossed (a and d), semi-crossed (b and e) and crossed (c and f) polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .





**Figure 4.23** - POM images of P2 biogels (gelatin + [BMIM][DCA] + 5CB + P2), after VOCs exposure, in 2 different fields of view, with uncrossed (a and d), semi-crossed (b and e) and crossed (c and f) polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .



**Figure 4.24** - POM images of P3 biogels (gelatin + [BMIM][DCA] + 5CB + P3), after VOCs exposure, in 2 different fields of view, with uncrossed (a and d), semi-crossed (b and e) and crossed (c and f) polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .

#### 4.4 Conclusions

Tailoring selectivity of VOC-responding materials was conducted by adding three different peptides (P1, P2 and P3) to a standard biogel. Selectivity was assessed using a proprietary electronic nose based on the optical response of the biogels upon their interaction with VOCs.

In this work we used a biogel, a proprietary composite gel-like sensor material composed by the LC 5CB and the IL [BMIM][DCA] self-assembled in micelle structures immobilized within the biopolymer gelatin. This material is able to form thin films responsive to VOCs. The biogel produce an optical response in presence of VOC molecules. The optical response is observable by POM and quantified using an e-nose developed in-house.

In this work, we have accessed the feasibility of adding VOC-selectivity to biogels thin films by incorporating in the standard biogel the benzene-sensitive peptide identified by Ju *et al.* (P1) and two modified versions of it, that contained norleucine (P2) or biphenylalanine (P3) added to their C-terminal to facilitate self-assembly and LC encapsulation within the micelles.

By labeling P1 with FITC we were able to verify that it was successfully incorporated in the micelles, although some P1 was also dispersed in the gel. P1, P2 and P3 were incorporated in the biogels and micelles were always produced.

We were also able to verify by POM that the peptides self-assemble in some structures with birefringent properties. P3, for instance, forms micelles with only the addition of [BMIM][DCA] and water. By testing a set of VOCs in a proprietary custom built e-nose we verified that the biogels without any peptide added responded more sharply to benzene and acetone. The addition of P1 seemed to amplify the response to benzene and toluene. The addition of P2 and P3 amplified the response signal to both acetone and benzene. Addition of P2 also increased the response intensity to toluene when compared to the addition of P1 to the standard biogel. Since the standard biogel already responded significantly towards benzene it would be of interest to incorporate P1 in biogels that do not respond when exposed to that solvent and test if a selectivity improvement towards benzene occurs.





## 5. Concluding remarks and future perspectives

In this work the distinction between 8 clinically relevant pathogens based on the emitted VOCs reported in literature was investigated. Data of interest was collected and machine learning methods were employed to classify the pathogens, based on the emitted VOCs. Data regarding VOC-pathogen interaction found in research articles between 1977 and 2016 was used to build an input data matrix with *Pseudomonas aeruginosa*, *Staphylococcus aureus*, *Aspergillus fumigatus*, *Escherichia coli*, *Helicobacter pylori*, *Proteus mirabilis*, *Klebsiella pneumoniae* and *Mycobacterium tuberculosis* and 269 VOCs. That set of VOCs was compared with the reported VOCs emitted from a healthy human body [133] to assess whether those compounds were also present in the healthy body. It was found that 3 VOCs (2-phenylanisol, 1-hydroxy-2-butanone and hydrogen cyanide) are not reported, so far, in the healthy human body. We have identified a minimal set of VOCs that allowed the separation of a specific pathogen from the others and compared those VOC lists with the ones found in other studies, for the same pathogens. It was found that indole for *E.coli*; 2-pentylfuran for *A.fumigatus*; isobutane for *H.pylori*; cymol for *M.tuberculosis*; hydrogen cyanide and methyl thiocyanate for *P.aeruginosa*; and 3-methylbutanoic acid for *S.aureus*, were referred in ours and other databases. Therefore, these compounds have strong probability of being biomarkers. Nonetheless, more work is required to define the range of normality/disease state in VOCs from humans in terms of concentration ranges in all bodily fluids. This data could then be used to interpret the constitution of each collected sample obtained from patients, to monitor their health state or infer about possible pathogen invasions. Also, since there are pathogens much more studied than others, from a statistical point of view, it would be important to balance the classifier input data, choosing, for example, 4 experiments for each pathogen, randomly, and repeat this classification procedure several times to confirm the reliability of the classifier results.

In the second part of the work, the selectivity of VOC-responding materials (biogels) was also tailored by adding three different peptides (P1, P2 and P3) to a standard biogel. Selectivity was assessed using a proprietary electronic nose based on the optical response of the biogels upon their interaction with VOCs.

E-noses allow the development of non-invasive, simple and fast tools for detecting VOCs from the human body [84]. However, despite the advances in e-nose research areas, sensors selectivity to detect VOCs remains a major challenge. In this work we used a biogel, a proprietary composite gel-like sensor material composed by the LC 5CB and the IL [BMIM][DCA] self-assembled in micelle structures immobilized within the biopolymer gelatin. This material is able to form thin films responsive to VOCs. The biogel produce an optical response in presence of VOC molecules. The optical response is observable by POM and

quantified using an e-nose developed in-house. Since some VOCs are recognized as disease biomarkers, tailoring the selectivity of biogel response towards certain VOC biomarkers would benefit its usability in future applications, namely for non-invasive testing of health conditions or environmental risk monitoring. In this work, we have accessed the feasibility of adding VOC-selectivity to biogels thin films by incorporating in the standard biogel the benzene-sensitive peptide identified by Ju *et al.* (P1) and two modified versions of it, that contained norleucine (P2) or biphenylalanine (P3) added to their C-terminal to facilitate self-assembly and LC encapsulation within the micelles. By labeling P1 with FITC we were able to verify that it was successfully incorporated in the micelles, although some P1 was also dispersed in the gel. P1, P2 and P3 were incorporated in the biogels and micelles were always produced. We were also able to verify by POM that the peptides self-assemble in some structures with birefringent properties. P3, for instance, forms micelles with only the addition of [BMIM][DCA] and water. By testing a set of VOCs in a proprietary custom built e-nose we verified that the biogels without any peptide added responded more sharply to benzene and acetone. The addition of P1 seemed to amplify the response to benzene and toluene. The addition of P2 and P3 amplified the response signal to both acetone and benzene. Since the standard biogel already responded significantly towards benzene it would be of interest to incorporate P1 in biogels that do not respond when exposed to that solvent and test if a selectivity improvement towards benzene occurs.

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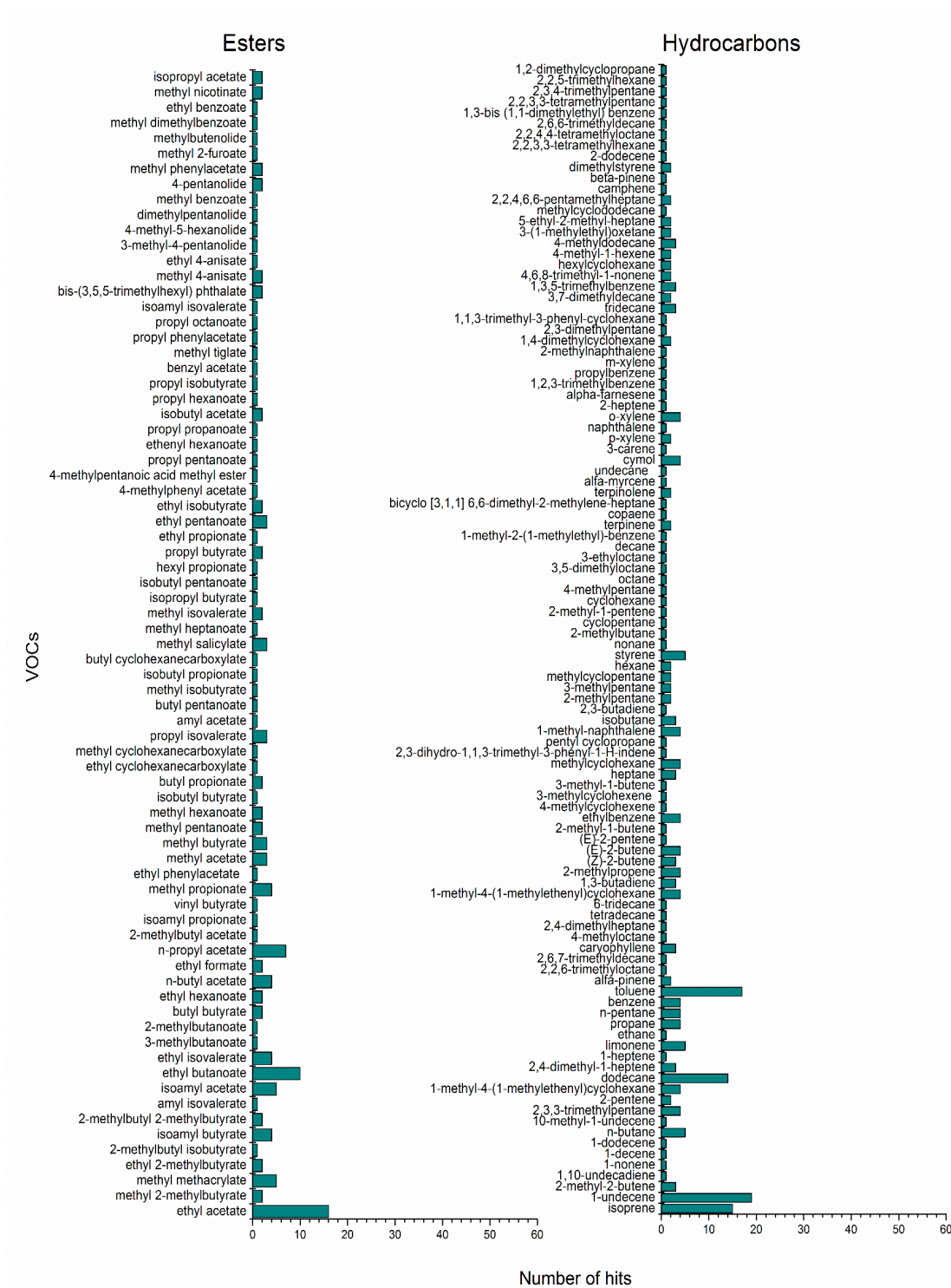
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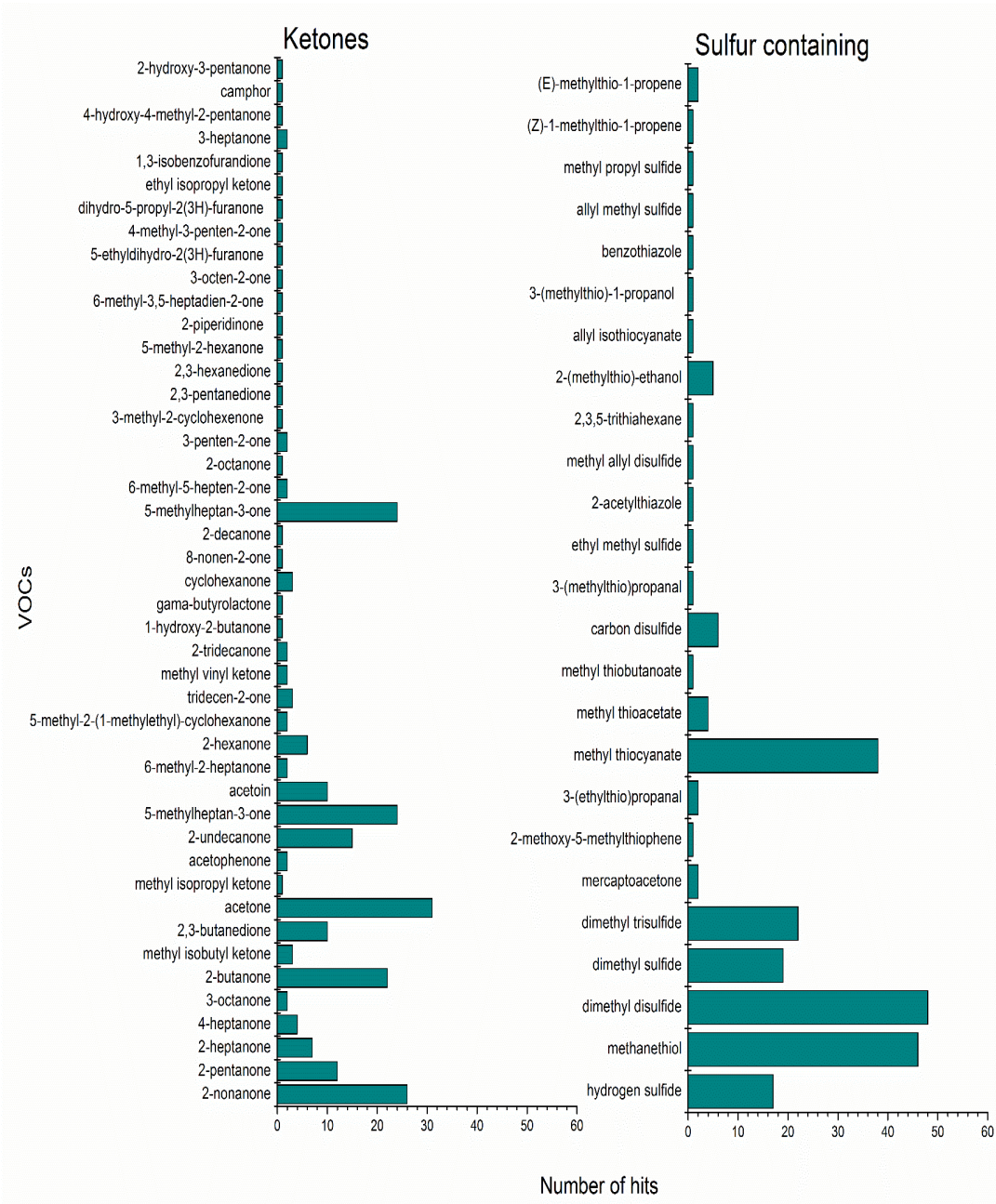
# Appendix

## Appendix 1

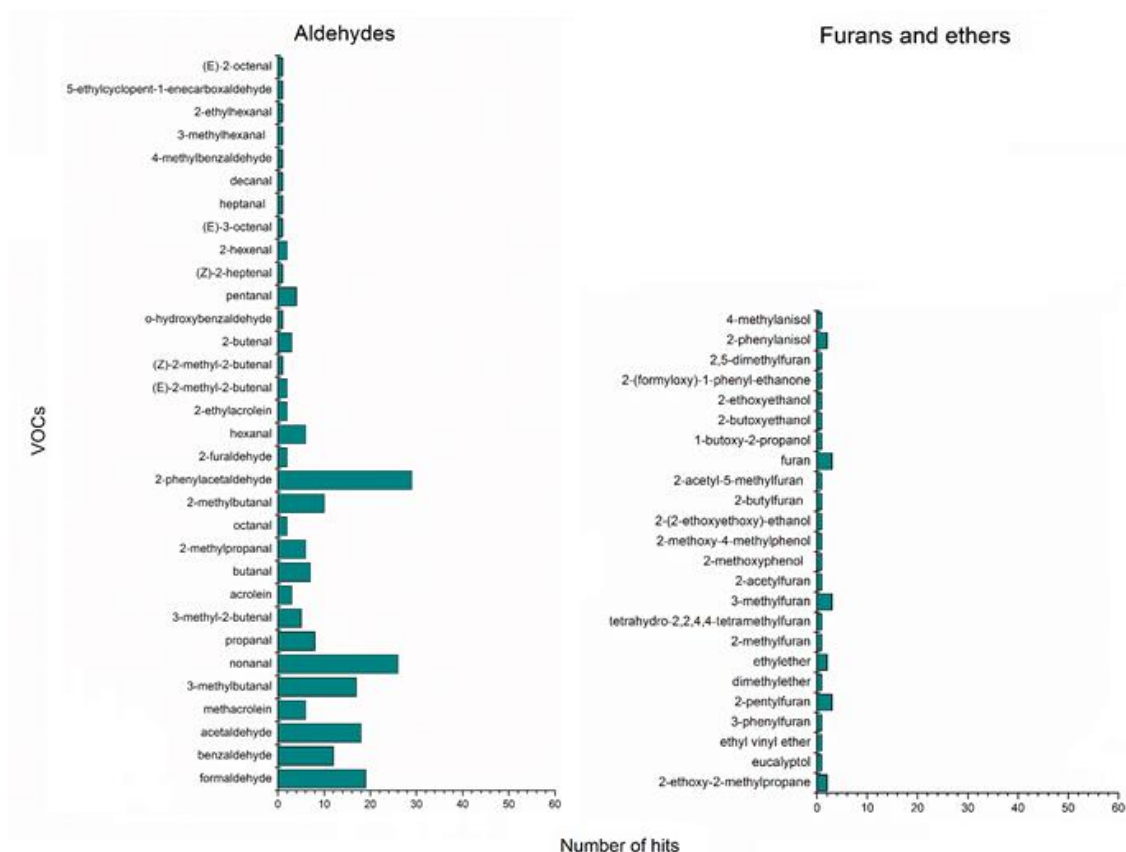


**Figure A1-** Representation of the number of hits for each individual VOC, in all the experiments, for the esters and hydrocarbons.

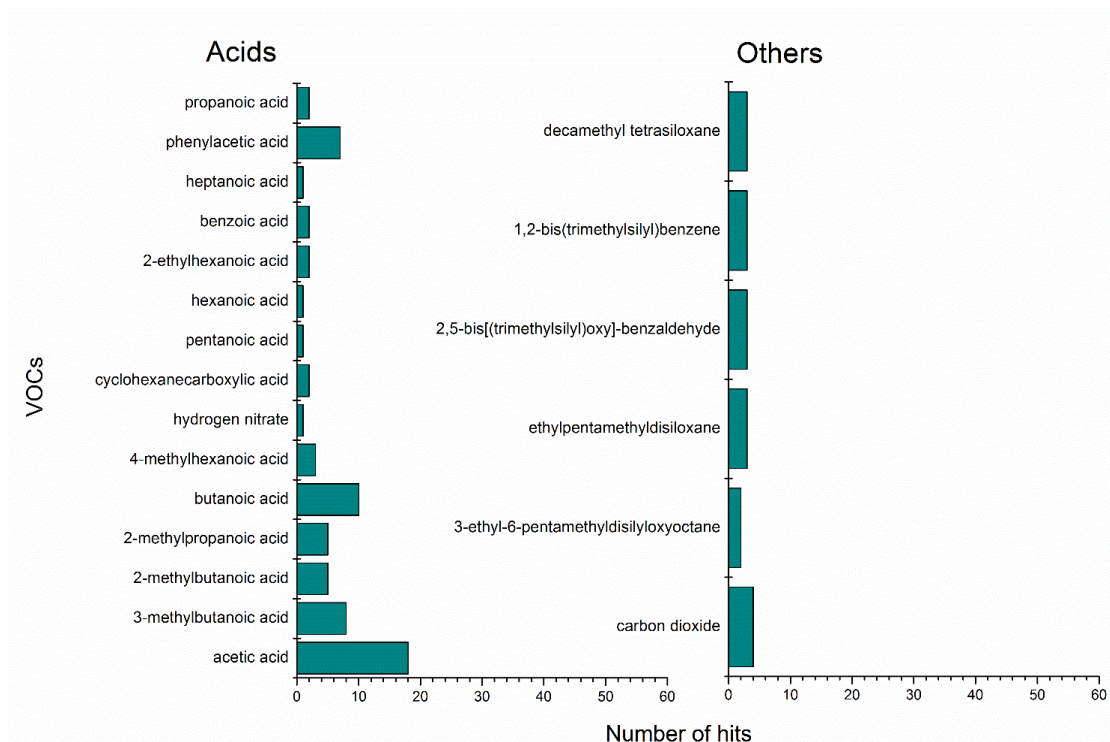




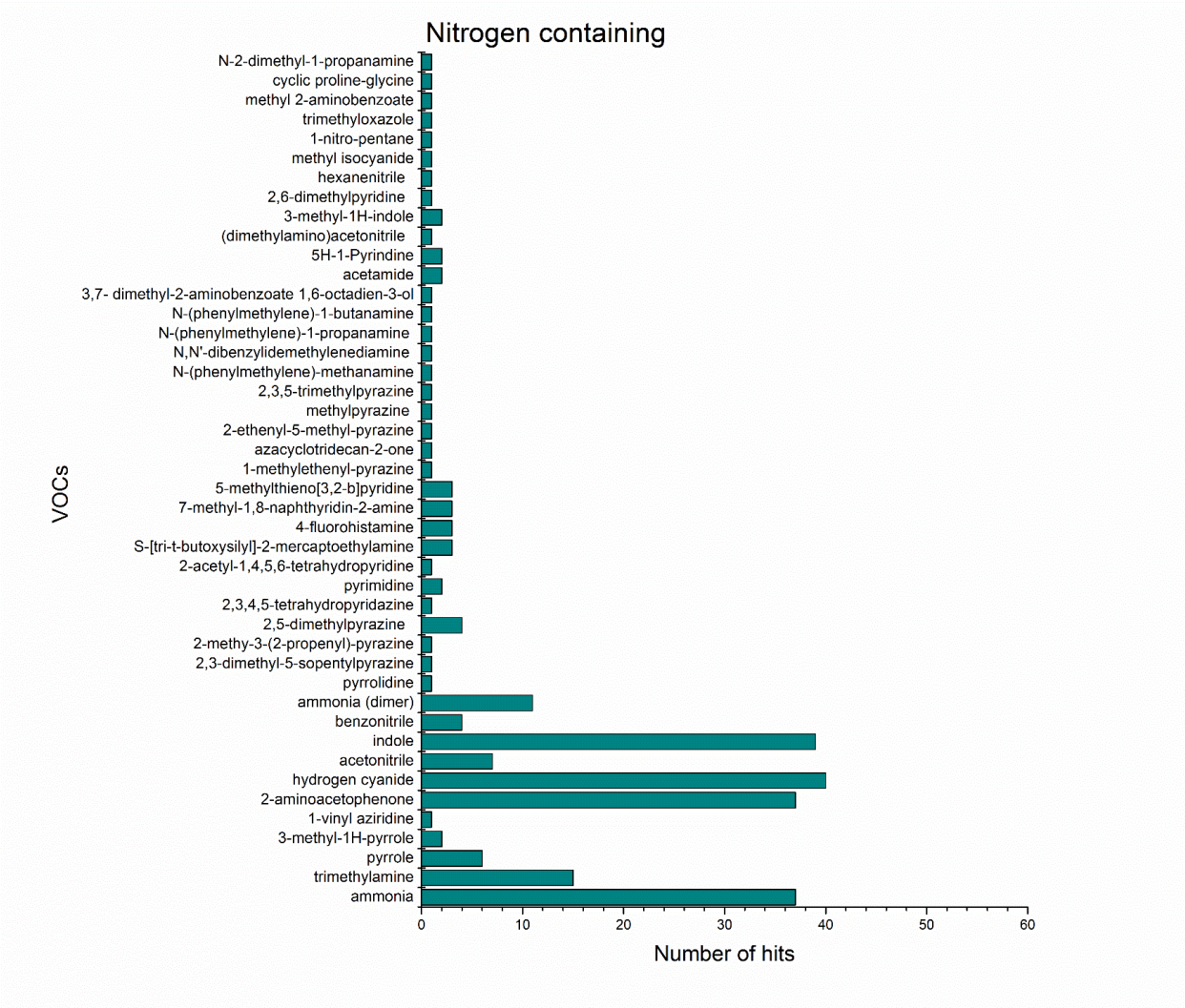
**Figure A2-** Representation of the number of hits for each individual VOC, in all the experiments, for the ketones and sulfur containing.



**Figure A3** - Representation of the number of hits for each individual VOC, in all the experiments, for aldehydes and furans and ethers.

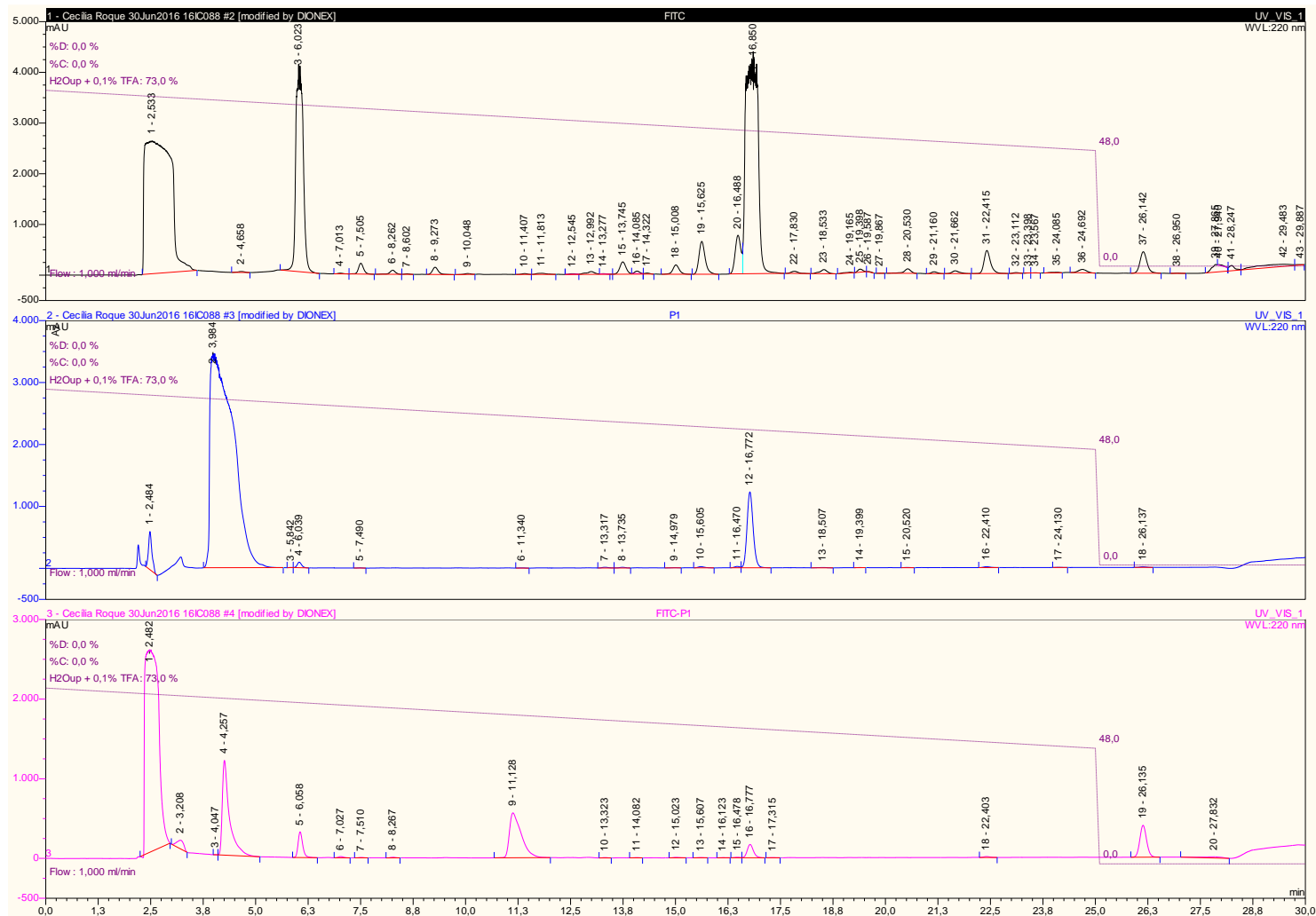


**Figure A4**- Representation of the number of hits for each individual VOC, in all the experiments, for acids and others.



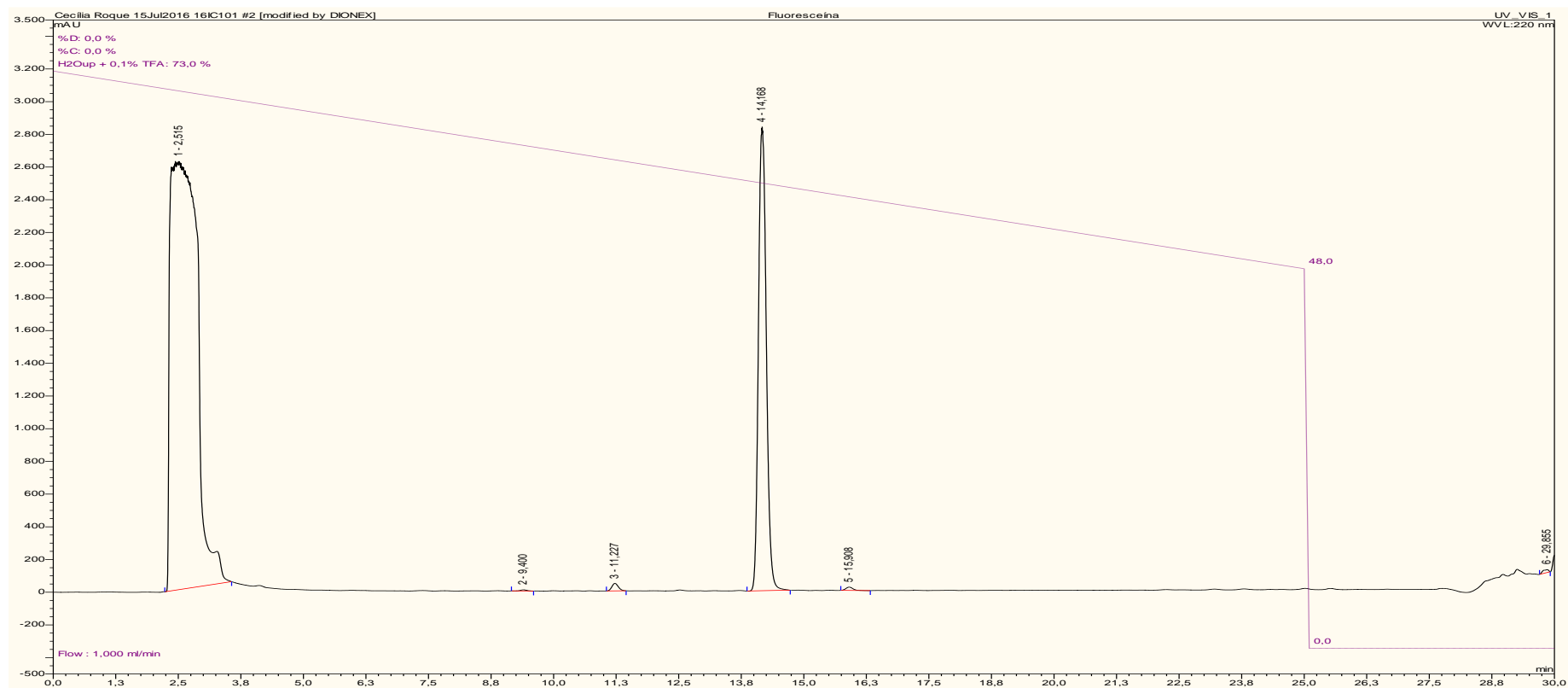
**Figure A5-** Representation of the number of hits for each individual VOC, in all the experiments, for the Nitrogen containing.



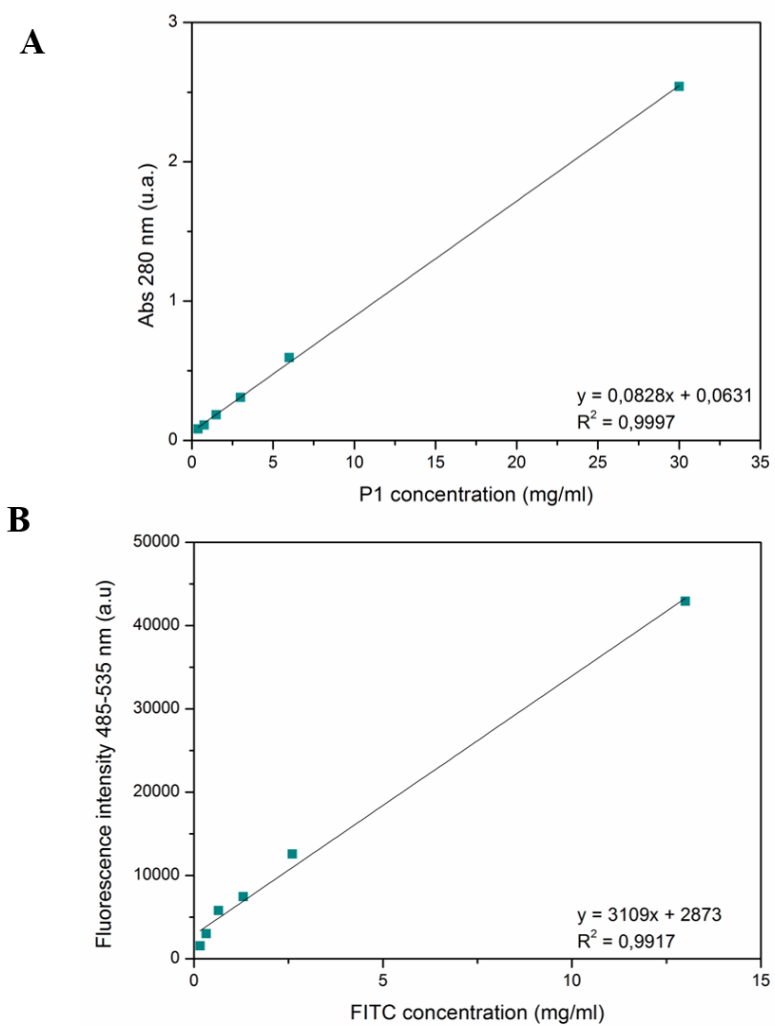


**Figure A6-** Chromatograms obtained for peptide P (blue), FITC (black) and P1-FITC conjugate (pink) samples.

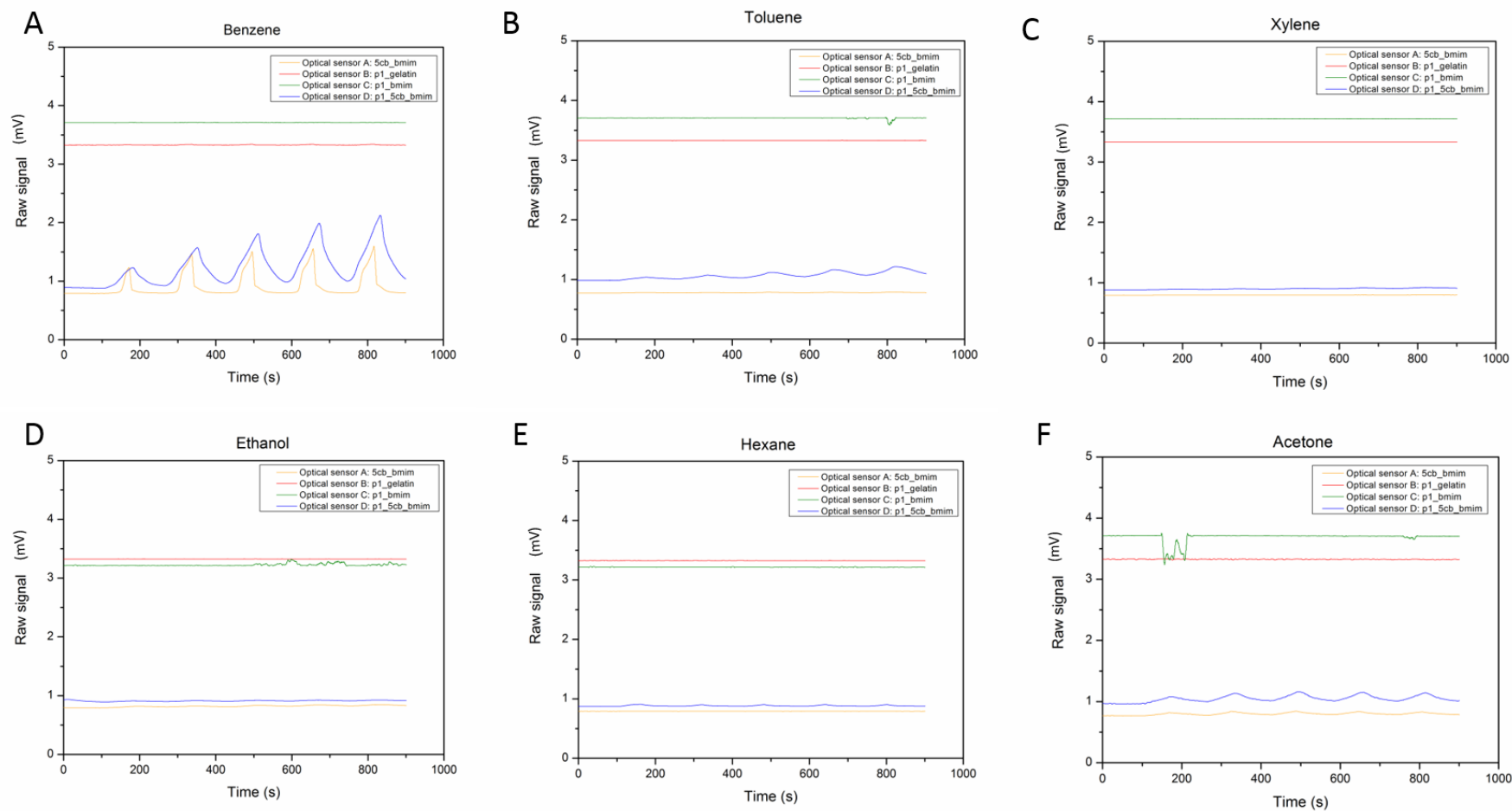
## Appendix



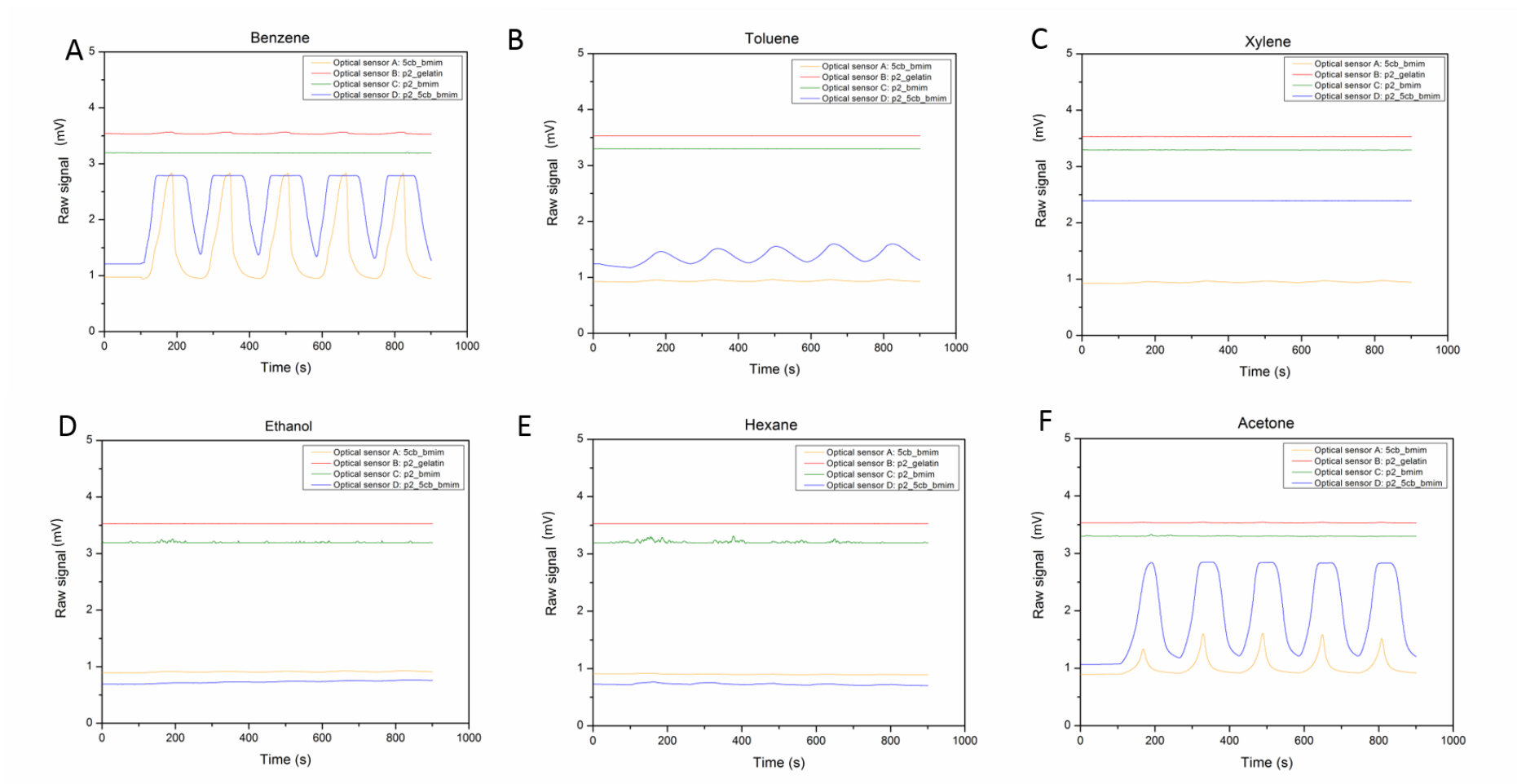
**Figure A7-** Chromatograms obtained for a fluorescein sample.



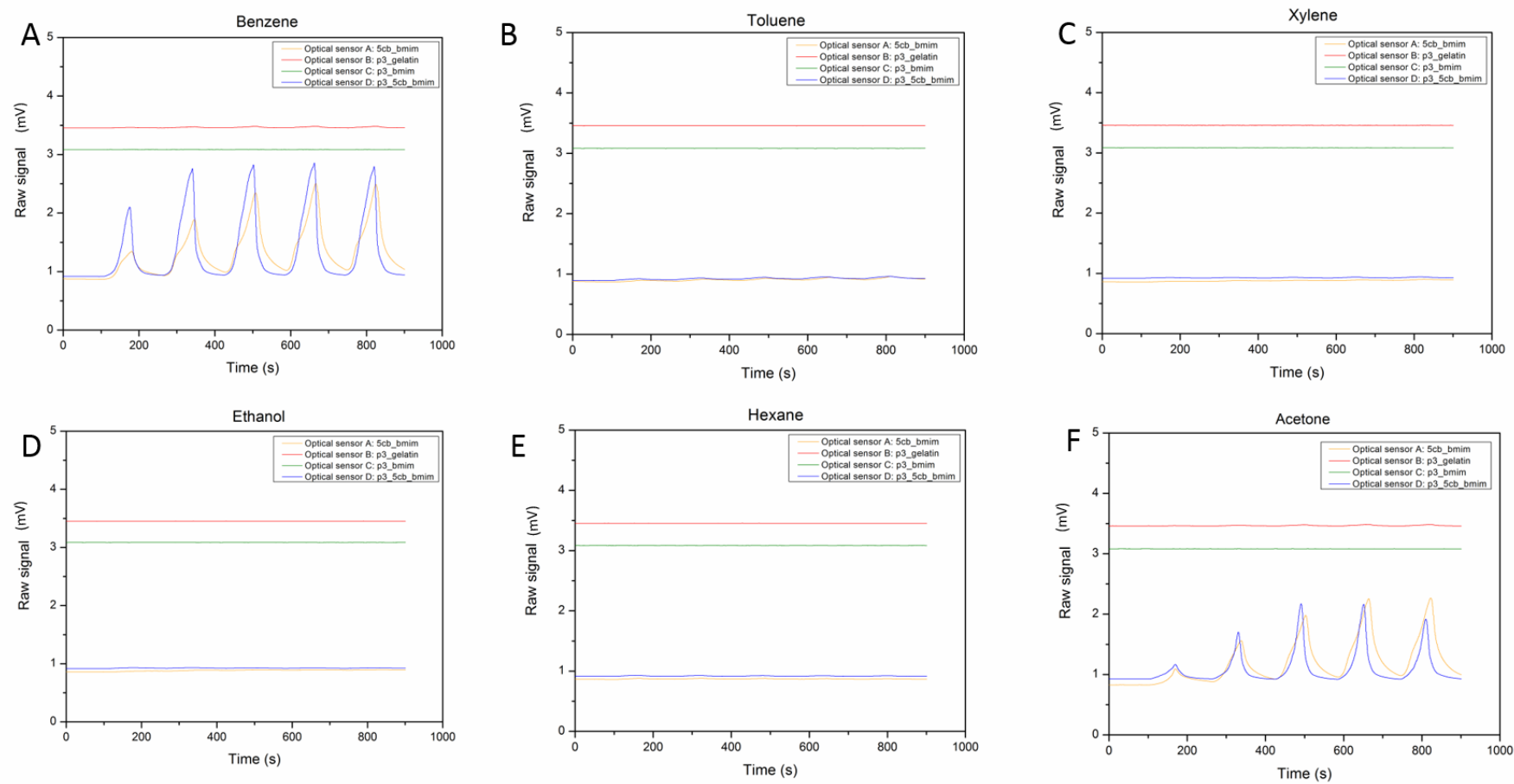
**Figure A8-** Calibration lines to quantify the amount of P1 (A) and FITC (B) present in the collected fraction.



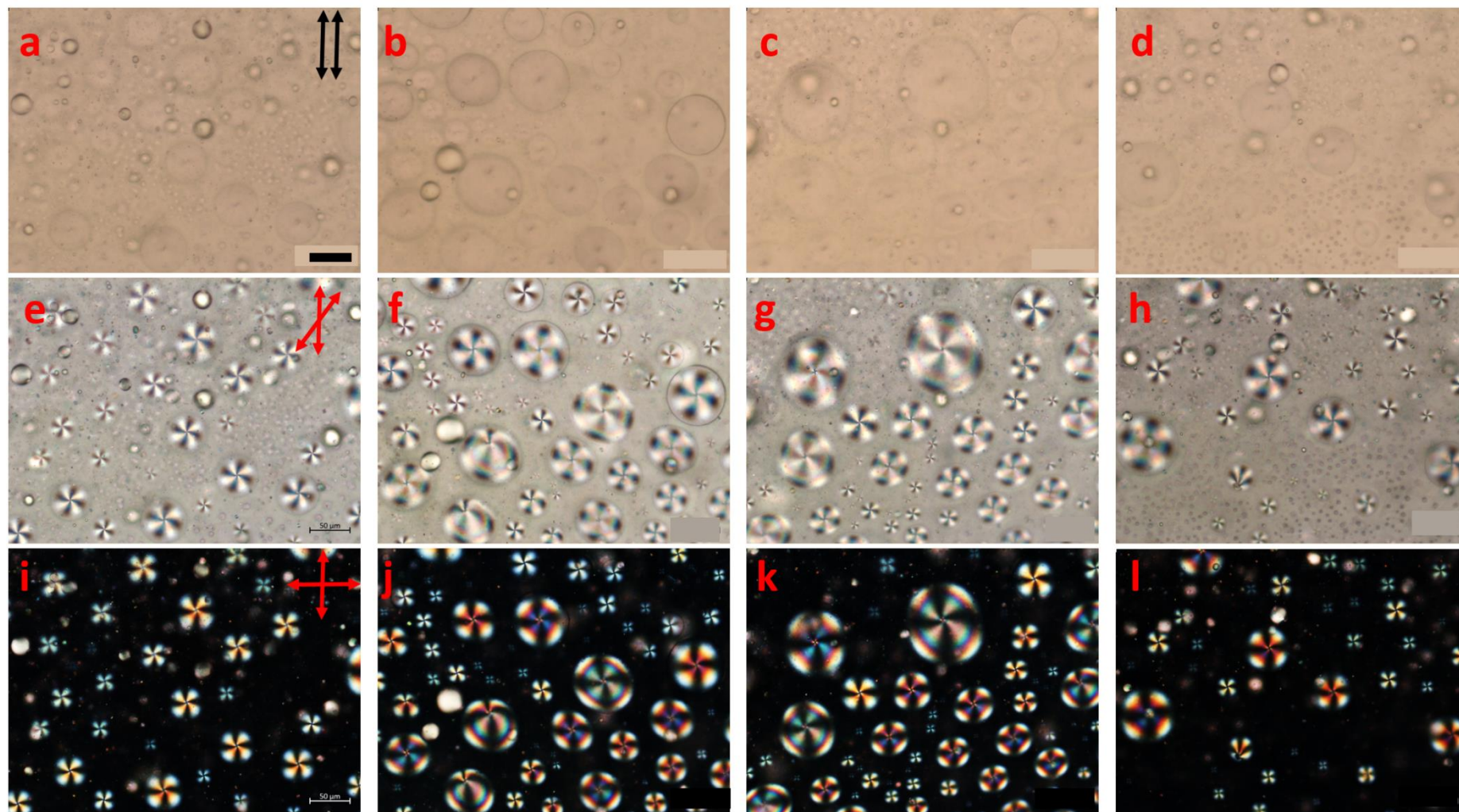
**Figure A9-** Responses of the biogels containing [BMIM][DCA]/5CB (yellow), gelatin/P1 (red), [BMIM][DCA]/P1 (green) and [BMIM][DCA]/5CB/P1 (blue) when exposed to benzene (A), toluene (B), xylene (C), ethanol (D), hexane (E) and acetone (F).



**Figure A10-** Responses of the biogels containing [BMIM][DCA]/5CB (yellow), gelatin/P2 (red), [BMIM][DCA]/P2 (green) and [BMIM][DCA]/5CB/P2 (blue) when exposed to benzene (A), toluene (B), xylene (C), ethanol (D), hexane (E) and acetone (F).

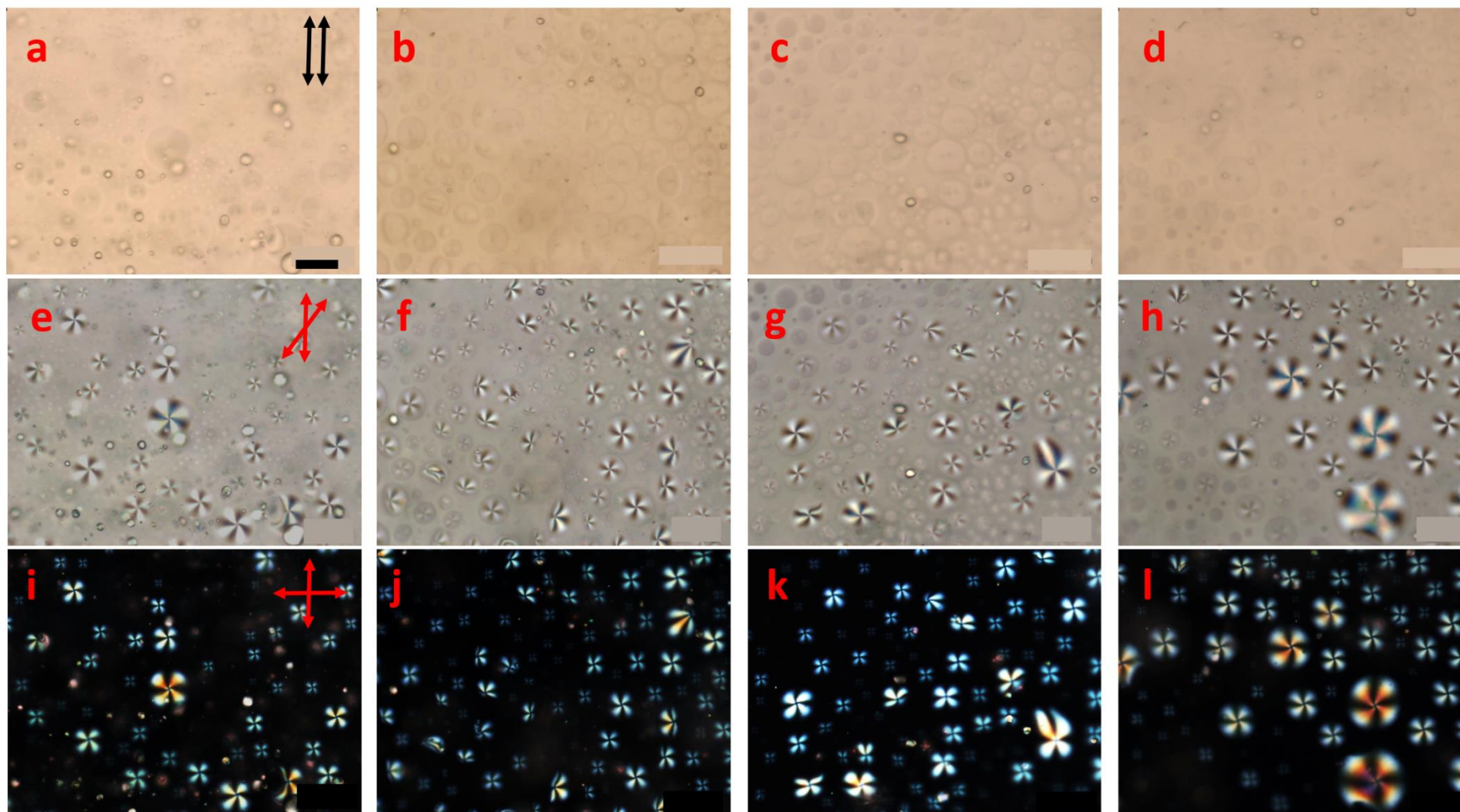


**Figure A11-** Responses of the biogels containing [BMIM][DCA]/5CB (yellow), gelatin/P3 (red), [BMIM][DCA]/P3 (green) and [BMIM][DCA]/5CB/P3 (blue) when exposed to benzene (A), toluene (B), xylene (C), ethanol (D), hexane (E) and acetone (F).



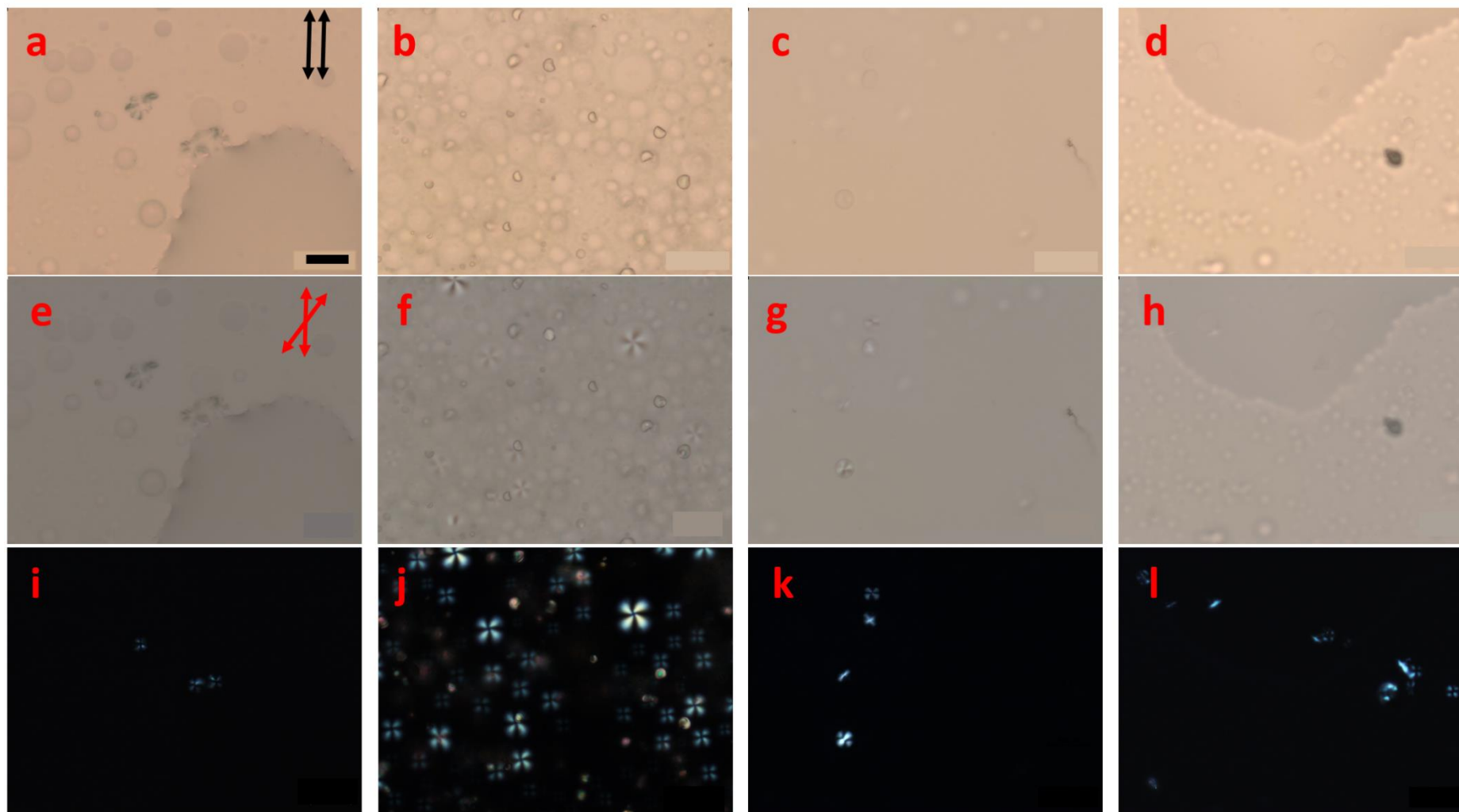
**Figure A12-** POM images of standard gelatin biogels ([BMIM][DCA]/5CB), in 5 different fields of view. a to d: uncrossed polarizers. e to h: semi-crossed polarizers. i to l: crossed polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .



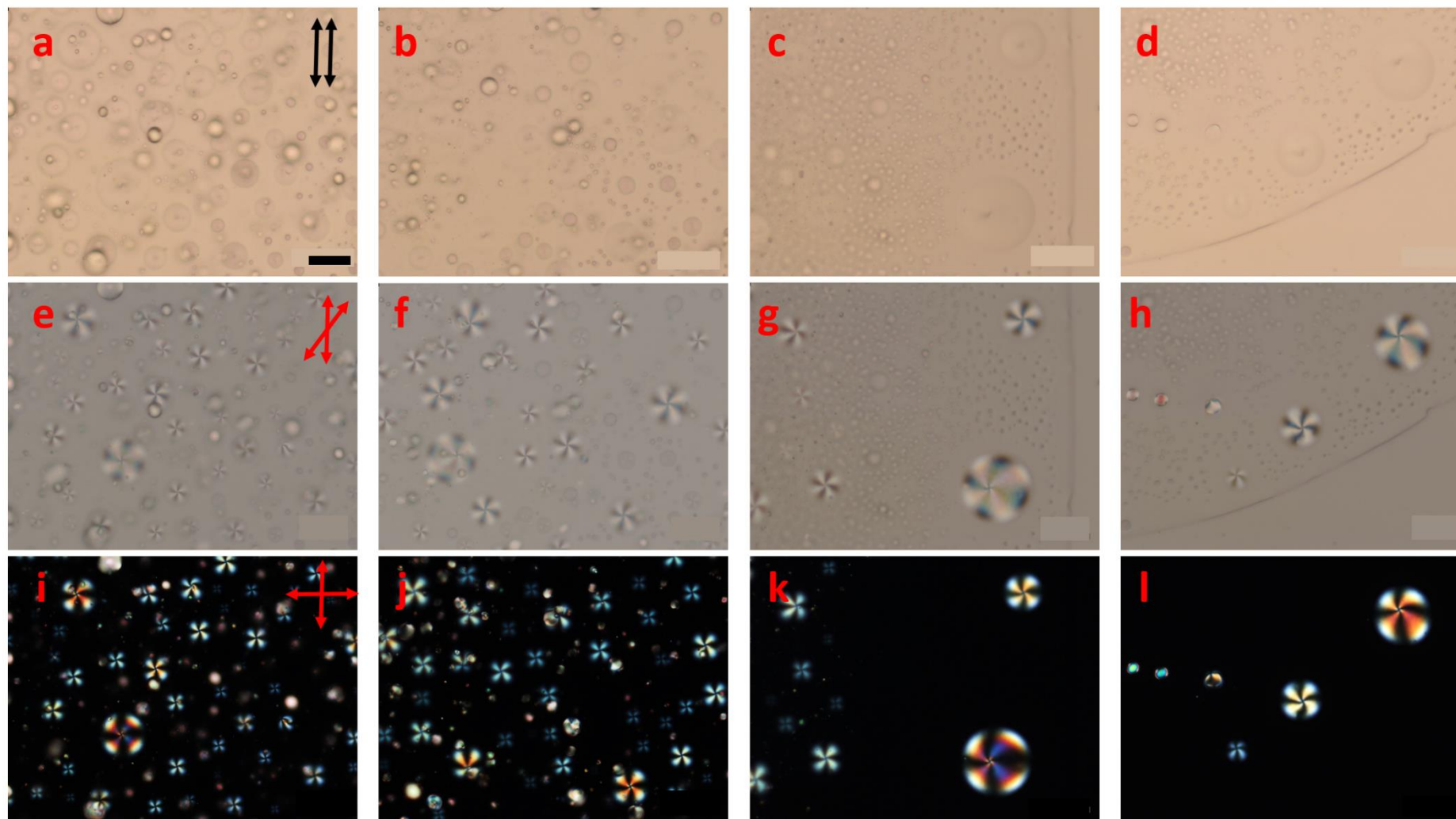


**Figure A13-** POM images of P1 gelatin biogels ([BMIM][DCA]/5CB/P1), in 5 different fields of view. a to d: uncrossed polarizers. e to h: semi-crossed polarizers. i to l: crossed polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .





**Figure A14-** POM images of P2 gelatin biogels ([BMIM][DCA]/5CB/P2), in 5 different fields of view. a to d: uncrossed polarizers. e to h: semi-crossed polarizers. i to l: crossed polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .



**Figure A15-** POM images of P3 gelatin biogels ([BMIM][DCA]/5CB/P3), in 5 different fields of view. a to d: uncrossed polarizers. e to h: semi-crossed polarizers. i to l: crossed polarizers. All images have the same scale. The scale bar corresponds to 50  $\mu\text{m}$ .

## **Appendix 2- Pathogen- VOCs Database**

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T		
Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range	Unit	Reference	Journal Code	Year		
Aspergillus fumigatus	fungus	2-pentylfuran	19602	SPME-GC-MS analysis				*				unknown	unknown	unknown	unknown	unknown	Sethi, S., Nanda, B., & Chakraborty, T. (2013). Chippendale, T. W., Gilchrist, F. J., Spaniel, P., Alcock, A., Lenney, W., & Smith, D. (2014). Quantification by SIFT-MS of volatile compounds emitted by Aspergillus fumigatus cultures and in co-culture with Pseudomonas aeruginosa, Staphylococcus aureus and Streptococcus pneumoniae. <i>Analytical Methods</i> , 6 (20), 8154-8164. DOI: 10.1039/C3AN00076G	Clin. Microbiol. Rev.	2013		
		ammonia (↑)	222																		
		methanol (↑)	887																		
		ethanol (↑)	702																		
		1-propanol (↑)	1031																		
		acetone (↑)	180	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	clinical isolate	cultured overnight on agar plates and identified	The cultures were incubated at 37 °C for 72 h prior to the headspace analysis	unknown	ppb		Anal. Methods	2014		
		methanethiol (↑)	878																		
		dimethyl sulfide (↑)	1068																		
		acetaldehyde (↓)	177																		
		butanal (↓)	261																		
		pentanal (↓)	8063																		
		3-octanone	246728	MCC-IMS analysis	not used	not used	not used	not used	not used	not used	not used	not used	DSM 21023	grown on columbia sheep blood agar for 24h 37°C	unknown	unknown	unknown	unknown	M., Borg-von Zeppelin, M., & Quintel, M. (2011). Detection of characteristic metabolites of Aspergillus fumigatus and Candida species	Mycoses	2011
		isopentanol	31260																		
		ethanol	702																		
		cyclohexanone	7967																		
		8-nonen-2-one	21108																		
		2,3-dihydro-1,1,3-trimethyl-3-phenyl-1-H-indole	19793																		
		2-tridecanone	11622																		
		1-methylethylpyrazine	62897	TD-GC-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	clinical isolate	stored in 10% glycerol broth at -80 °C, and revived by subculturing on Sabouraud dextrose agar (SDA) supplemented with 0.02% chloramphenicol, for 2 × 7 d at 37 °C.	16h, 24h and 48h	unknown	unknown	Neerincx, A. H., Geurts, B. P., Habets, M. F. J., Boon, J. A., van Loon, J., Jansen, J. J., ... & Wevers, R. A. (2016). Identification of Pseudomonas aeruginosa and Aspergillus fumigatus mono- and co-cultures based on volatile biomarker combinations. <i>Journal of breath research</i> , 10 (1), 016002.	J. Breath Res.	2016	
		azacyclotridecan-2-one	13690																		
		4-ethyl-5-methylpyrazine	26335																		
		2-undecanone	8163																		
		2-nonanone	13187																		
		1-hydroxy-2-propanone	8299																		
		2-acetylthiazole	520108																		
Campylobacter jejuni	Gram negative bacterium	butanoic acid (↓)	264																		
		2-methylbutanoic acid (↓)	8314																		
		carbon disulfide (↓)	6348																		
		dimethyl sulfide (↓)	1068																		
		dimethyl disulfide (↓)	12232																		
		indole (↓)	798																		
		4-methylphenol (↓)	2879																		
		3-methylfuran (↓)	13587																		
		3-methylbutanoic acid (↓)	10430																		
		dimethyl trisulfide (↓)	19310																		
		limonene (↓)	22311																		
		methanethiol (↓)	878																		
		propanal (↓)	527																		
		methyl acetate (↓)	6584																		
		2-hexanone (↓)	11583																		
		cymol (↓)	7463																		
		2-phenylacetaldehyde (↓)	998																		
		methyl butyrate (↓)	12180																		
		methyl propionate (↓)	11124																		
		2,3-pentanedione (↑)	11747																		
		phenol (↑)	996																		
		2-propanol (↓)	3776																		
		2-methylpropanoic acid (↓)	6590																		
		3-methyl-1H-indole (↓)	6736																		
		isobutanol (↓)	6560																		
		3-carene (↓)	26049																		
		heptanoic acid (↑)	8094																		
		methyl isobutyl ketone (↓)	7909																		
		isopentanol (↓)	31260																		
		2-phenylethanol (↑)	6054																		
		1-propanol (↑)	1031																		
		acetophenone (↑)	7410																		
		2,3-hexanedione (↓)	19707																		
		caryophyllene (↓)	5322111																		
		n-propyl acetate (↑)	7997																		
		styrene (↑)	7501																		
		toluene (↑)	11440																		
		pentanal (↑)	8063																		
		2-(2-ethoxyethoxy)-ethanol (↑)	8146																		
		2-butanol (↓)	6568																		
		n-pentane (↓)	8003																		
		2-pentylfuran (↑)	19602																		
		ethyl pentanoate (↓)	10882																		
		1-octanol (↑)	957																		
		2-ethylhexanoic acid (↑)	8697																		
		methanol (↑)	887																		
		3-penten-2-one (↑)	12248																		
		benzoic acid (↑)	243																		
		ethylbenzene (↑)	7500																		
		linolol (↑)	6549																		
		heptanal (↑)	8130																		
		phenylmethanol (↑)	244																		
		octanal (↑)	454																		
		3-heptanone (↓)	7802																		
		methyl salicylate (↓)	4133																		
		ethenyl hexanoate (↑)	76451																		
		2-ethyl-1-hexanol (↑)	7720																		
		1-nonanol (↓)	8914																		
		2-butenal (↑)	447466																		
		5-methyl-2-hexanone (↓)	8034																		
		2-piperidone (↑)	12665																		
		6-methyl-3,5-heptadien-2-one (↑)	5370101																		
		allyl isothiocyanate (↓)	5971																		
		dodecane (↑)	8182	SPME-GC-MS analysis								x	unknown	unknown	unknown	unknown	unknown	unknown	Garner, C. E., Smith, S., de Lacy Costello, B., White, P., Spencer, R., Probert, C. S., & Ratcliffe, N. M. (2007). Volatile organic compounds from feces and their potential for diagnosis of gastrointestinal disease. The FASEB Journal, 21(8), 1675-1688.	FASEB J.	2007
		nonanal (↑)	31289																		
		propyl propanoate (↑)	7805																		
		p-xylene (↑)	7809																		
		acetamide (↓)	178																		
		3-methyl-2-butenal (↑)	61020																		
		2-butyfuran (↑)	20534																		
		decanal (↑)	8175																		
ethyl hexanoate (↑)	31265																				
1-octen-3-ol (↑)	18837																				
2-acetyl-5-methylfuran (↑)	14514																				
3-octen-2-one (↑)	15475																				
cyclohexanecarboxylic acid (↑)	7413																				
5-ethylidihydro-2(3H)-furanone (↑)	12756																				
isobutyl acetate (↓)	8038																				
acetonitrile (↑)	6342																				
4-methylbenzaldehyde (↑)	7725																				
isoamyl butyrate (↑)	7795																				
propyl hexanoate (↑)	12293																				
methacrolein (↑)	6562																				
naphthalene (↑)	931																				
2,6-dimethylpyridine (↑)	7937																				
trichloromethane (↑)	6212																				
benzonitrile (↑)	7505																				
3-methylhexanal (↑)	140511																				
hexanenitrile (↑)	123																				

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				1-butoxy-2-propanol (↑)		21210																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
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				5H-1-Pyridine (↑)		575987																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				benzyl acetate (↑)		8785																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				alpha-farnesene (↑)		5281516																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				1,2,3-trimethylbenzene (↑)		10686																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				propylbenzene (↑)		7668																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				ethyl isopropyl ketone (↑)		11265																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
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				cis-1-p-menthanol (↑)		89437																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				2-butoxyethanol (↑)		8133																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				2-ethoxyethanol (↑)		8076																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				2-(formyl-oxo)-1-phenyl-ethanone (↑)		569595																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				2,5-dimethylfuran (↑)		12266																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				2-ethylhexanal (↑)		31241																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				methyl isocyanide (↑)		11646																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				methyl tiglate (↑)		532652																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				methyl vinyl ketone (↑)		6570																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				2-methylnaphthalene (↑)		7055																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				1-nitro-pentane (↑)		220639																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				ethyl pentanoate (↑)		10882																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				propyl phenylacetate (↑)		221641																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				propyl octanoate (↑)		69351																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				2,5-dimethylpyrazine (↑)		31252																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				isoamyl isovalerate (↑)		12613																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				trimethylamine (↑)		1146																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				phenylacetic acid (↑)		999																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				ethanol		702																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																															
	Candida albicans	Fungus		formaldehyde		712		SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown									CDC S-24	sterile urine (20 ml) from healthy males inoculated to a concentration of between 10^7 and 10^9 cfu/mL	37°C for 6h	unknown	ppb		Storer, M. K., Hibbard-Melies, K., Davis, B., & Scotter, J. (2011). Detection of volatile compounds produced by microbial growth in																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																						

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T
Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range	Unit	Reference	Journal Code	Year
		1-propanol (↓)	1031	SPME-GC-MS analysis								unknown	unknown	fresh or 7 days	unknown	unknown	Garner, C. E., Smith, S., de Lacy Costello, B., White, P., Spencer, R., Probert, C. S., & Ratcliffe, N. M. (2007). Volatile organic compounds from feces and their potential for diagnosis of gastrointestinal disease. The FASEB Journal, 21(8), 1675-1688.	FASEB J.	2007
		2-acetylfuran (↓)	14505																
		styrene (↑)	7501																
		toluene (↓)	1140																
		pentanal (↓)	8063																
		n-butyl acetate (↓)	31272																
		propyl butyrate (↓)	7770																
		methyl pentanoate (↓)	12206																
		2-butanol (↓)	6568																
		ethyl propionate (↓)	7749																
		2-octanone (↓)	8093																
		methyl thioacetate (↓)	73750																
		2-pentylfuran (↓)	19602																
		methyl hexanoate (↓)	7824																
		ethyl pentanoate (↓)	10882																
		2-ethylhexanoic acid (↓)	8697																
		methanol (↓)	887																
		butyl propionate (↓)	11529																
		3-penten-2-one (↑)	12248																
		benzoic acid (↑)	243																
		phenylmethanol (↑)	244																
		undecane (↓)	14257																
		2,3,5-trithiahexane (↓)	93236																
		2-ethyl-3-hexanol (↑)	7720																
		dodecane (↓)	8182																
		2-butenal (↓)	447466																
		acetamide (↓)	178																
		2-furaldehyde (↓)	7962																
		2-nonanone (↓)	13187																
		3-methyl-2-butenal (↑)	61020																
		1-octen-3-ol (↑)	18827																
		4-heptanone (↑)	31246																
		methacrolein (↓)	6562																
		ethyl isobutyrate (↑)	7342																
		benzonitrile (↑)	7505																
		isomyl acetate (↑)	31276																
		4-methyl-1-pentanol (↑)	12296																
		1-penten-3-ol (↑)	12020																
		3-methyl-2-butenol (↑)	11173																
		3-methyl-2-cyclohexenone (↑)	14511																
		6-methyl-2-heptanone (↑)	13572																
		(Z)-2-heptenal (↑)	5362616																
		2-hexenal (↑)	5281168																
		(E)-3-octenal (↑)	5283325																
		(Z)-2-pentenol (↑)	5364919																
		acrolein (↑)	7847																
		5H-1-Pyridine (↑)	575987																
		4-methylphenyl acetate (↑)	8797																
		(dimethylamino)acetoneitrile (↑)	61237																
		methyl isovalerate (↑)	11160																
		propyl isovalerate (↑)	11176																
		cis-1-p-menthanol (↑)	89437																
		cyclohexanone (↑)	7967																
		cyclopentane (↑)	9253																
		2-(methylthio)-ethanol (↑)	78925																
		4-methylpentanoic acid methyl ester (↑)	17008																
		propyl pentanoate (↑)	67328																
		2-methoxyphenol (↑)	460																
		2-methoxy-5-methylphenol (↑)	7144																
		2,5-dimethylpyrazine (↑)	31252																
		acetone	180	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	NTCC 775	sterile urine (20 mL) from healthy males inoculated to a concentration of between 10 <sup>4</sup> and 10 <sup>9</sup> cfu/mL	37°C for 6h	unknown	ppb	Storer, M. K., Hibbard-Melles, K., Davis, B., & Scotter, J. (2011). Detection of volatile compounds produced by microbial growth in urine by selected ion flow tube mass spectrometry (SIFT-MS). Journal of microbiological methods, 87(1), 111-113. doi: 10.1016/j.mimet.2011.06.012.	J. Microbiol. Methods	2011
		2-butanone	6569																
		2-pentanone	7895																
		formaldehyde	712																
		2-methylbutanal	7284																
		ethyl butanoate	7762																
		n-propyl acetate	7997																
		hydrogen sulfide	402																
		dimethyl sulfide	1068																
		dimethyl disulfide	12232																
		methanethiol	878	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	Bos, L. D., Sterk, P. J., & Schultz, M. J. (2013). Volatile metabolites of pathogens: a systematic review. doi:10.1371/journal.ppat.1003311	PLOS	2013	
		ammonia	222																
		propene	8252																
		1-butanol	263																
		1-propanol	1031																
		1-pentanol	6276																
		phenylacetic acid	999																
		formaldehyde	712																
		2-butanone	6569																
		2-pentanone	7895																
		acetone	180	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	clinical isolate	blood agar	48h	146.13 14.06 1.34 55.07 17.72 365.59 4.13 335.86 148.46 6.58 334.37 41.93 0.8 44.28	ppb	Thorn, R., Reynolds, D. M. and Greenman, J. (2011) Multivariate analysis of bacterial volatile compound profiles for discrimination between selected species and strains in vitro. Journal of Microbiological Methods, 84 (2), pp. 258-264. ISSN 0167-7012 DOI: 10.1016/j.mimet.2010.12.001	J. Microbiol. Methods	2011	
		ethanol	702																
		ethyl butanoate	7762																
		formaldehyde	712																
		hydrogen sulfide	402																
		isoprene	6557																
		methanethiol	878																
		phenylacetic acid	999																
		pyrrole	8027																
		trimethylamine	1146																
		acetone	180	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	W310	sterile urine (20 mL) from healthy males inoculated to a concentration of between 10 <sup>4</sup> and 10 <sup>9</sup> cfu/mL	37°C for 6h	unknown	ppb	Storer, M. K., Hibbard-Melles, K., Davis, B., & Scotter, J. (2011). Detection of volatile compounds produced by microbial growth in urine by selected ion flow tube mass spectrometry (SIFT-MS). Journal of microbiological methods, 87 (1), 111-113. doi: 10.1016/j.mimet.2011.06.012.	J. Microbiol. Methods	2011	
		acetic acid	176																
		methanol	887																
		ethanol	702																
		formaldehyde	712																
		ethyl acetate	8857																
		ethyl butanoate	7762																
		n-propyl acetate	7997																
		hydrogen sulfide	402																
		dimethyl disulfide	12232																

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T											
Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range		Reference	Journal Code	Year											
															Value	Unit														
Escherichia coli	Gram negative bacteria	trimethylamine	1146	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	ATCC 25922	blood culture bottles	24 h	unknown	unknown	Allardcyce, Randall A., et al. "Detection of volatile metabolites produced by bacterial growth in blood culture media by selected ion flow tube mass spectrometry (SIFT-MS)." <i>Journal of microbiological methods</i> 65.2 (2006): 361-365.	J. Microbiol. Methods	2006											
		2-aminoacetophenone	11952																											
		indole	798		IMR-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown									unknown	DH5 5678	blood agar plates	24 h	unknown	unknown	Dolch, M. E., et al. "Volatile compound profiling for the identification of Gram-negative bacteria by ion-molecule reaction-mass spectrometry." <i>Journal of</i>	J. Appl. Microbiol.	2012		
		ethanol	702																											
		1-pentanol	6276																											
		formaldehyde	712																											
		acetaldehyde	177																											
		acetic acid	176																											
		hydrogen sulfide	402																											
		methanethiol	878																											
		dimethyl sulfide	1068																											
		dimethyl disulfide	12232																											
		trimethylamine	1146																											
		indole	798																											
		1-propanol	1031																											
		2-aminoacetophenone	11952																											
		hexanal	6184																											
		carbon dioxide	280																											
		ammonia	222																											
		methanethiol	878																											
		indole	798																											
		1-butanol	263																											
		1-pentanol	6276																											
		acetoin	179																											
		butanoic acid	264																											
		ethanol	702																											
		ethyl acetate	8857																											
		ethyl butanoate	7762																											
		formaldehyde	712																											
		hydrogen sulfide	402																											
		indole	798																											
		isoprene	6557																											
		methanethiol	878																											
		phenylacetic acid	999																											
		trimethylamine	1146																											
		1-butanol	263																											
		1-pentanol	6276																											
		2-aminoacetophenone	11952																											
		acetoin	179																											
		butanoic acid	264																											
		ethanol	702																											
		ethyl acetate	8857																											
		formaldehyde	712																											
		hydrogen sulfide	402																											
		indole	798																											
		methanethiol	878																											
		phenylacetic acid	999																											
		trimethylamine	1146																											
		isoprene	6557																											
		1-propanol	1031																											
		3-methylbutanal	11552																											
		2-methylbutanal	7284																											
		2,3,3-trimethylpentane	11215																											
		benzaldehyde	240																											
		acetic acid	176																											
		2,3-butanedione	650																											
		n-propyl acetate	7997																											
		indole	798																											
		3-methyl-4-(1-methylethenyl)cyclohexane	14299																											
		acetonitrile	6342																											
		ethanol	702																											
		indole	798																											
		ethanol	702																											
		acetone	180																											
		2-nonanone	13187																											
		2-heptanone	8051																											
		1-octanol	957																											
		1-decanol	8174																											
		1-dodecanol	8193																											
		2-undecanone	8163																											
		tridecen-2-one	53427438																											
		ethanol	702																											
		1-propanol	1031																											
		isopentanol	31260																											
		1-octanol	957																											
		9-decenol	25612																											
		1-decanol	8174																											
		indole	798																											
		1-dodecanol	8193																											
		(Z)-7-tetradecen-1-ol	5362795																											
		1-tetradecanol	8209																											
		dimethyl disulfide	12232																											
		ethanol	702																											
		2-nonanone	13187																											
		2-heptanone	8051																											
		pentyl cyclopropane	75640																											
		indole	798																											
		ethanol (↑)	702																											
		indole (↑)	798																											
		2-(methylthio)-ethanol (↑)	78925																											
		3-methylbutanal (↑)	11552																											
		dimethyl disulfide (↑)	12232																											
		methypyrazine (↑)	7976																											
		2-(methylthio)-ethanol (↑)	78925																											
		phenol (↑)	996																											
		dimethyl trisulfide (↑)	19310																											
		benzonitrile (↑)	7505																											
		2,3,5-trimethylpyrazine (↑)	26808																											
		N-(phenylmethylene)-methanamine (↑)	72954																											
		2-nonanone (↑)	13187																											
N,N'-dibenzylidenedimethylenediamine (↑)	66033																													
2-decanone (↑)	12741																													
N-(phenylmethylene)-1-propanamine (↑)	250250																													
ethyl phenylacetate (↑)	7590																													
N-(phenylmethylene)-1-butanamine (↑)	296031																													
indole (↑)	798																													
1-methyl-naphthalene (↑)	7002																													
1-decanol	8174																													
indole	798																													
1-dodecanol	8193																													
acetic acid	176																													
1-decanol	8174																													
Escherichia coli	Gram negative bacteria	trimethylamine	1146	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	24 h	unknown	unknown	Thorn, R., Reynolds, D. M. and Greenman, J. (2011) Multivariate analysis of bacterial volatile compound profiles for discrimination between selected species and strains in vitro.Journal of Microbiological Methods, 84 (2), pp. 258-264. ISSN 0167-7012 DOI: 10.1016/j.jmimet.2010.12.001.	J. Microbiol. Methods	2011											
		2-aminoacetophenone	11952																											
		indole	798		GC-MS analysis	not used	not used	not used	not used	not used	not used									not used	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	unknown	unknown	unknown	Boots AW, Smolinska A, van Berkel JJ, Fijten RR, Stobberingh EE, et al. (2014) Identification of microorganisms based on headspace analysis of volatile organic compounds by gas chromatography-mass spectrometry. J Breath Res 8: 027106. doi:10.1088/1752-7155/8/2/027106.	J. Breath Res.	2014		
		ethanol	702																											
		acetone	180																											
		2-nonanone	13187																											
		2-heptanone	8051																											
		1-octanol	957																											
		1-decanol	8174																											
		1-dodecanol	8193																											
		2-undecanone	8163																											
		tridecen-2-one	53427438																											
		ethanol	702																											
		1-propanol	1031																											
		isopentanol	31260																											
		1-octanol	957																											
		9-decenol	25612																											
		1-decanol	8174																											
		indole	798																											
		1-dodecanol	8193																											
		(Z)-7-tetradecen-1-ol	5362795																											
		1-tetradecanol	8209																											
		dimethyl disulfide	12232																											
		ethanol	702																											
		2-nonanone	13187																											
		2-heptanone	8051																											
		pentyl cyclopropane	75640																											
		indole	798																											
		ethanol (↑)	702																											
		indole (↑)	798																											
		2-(methylthio)-ethanol (↑)	78925																											
		3-methylbutanal (↑)	11552																											
		dimethyl disulfide (↑)	12232																											
		methypyrazine (↑)	7976																											
		2-(methylthio)-ethanol (↑)	78925																											
		phenol (↑)	996																											
		dimethyl trisulfide (↑)	19310																											
		benzonitrile (↑)	7505																											
		2,3,5-trimethylpyrazine (↑)	26808																											
		N-(phenylmethylene)-methanamine (↑)	72954																											
		2-nonanone (↑)	13187																											
		N,N'-dibenzylidenedimethylenediamine (↑)	66033																											
		2-decanone (↑)	12741																											
		N-(phenylmethylene)-1-propanamine (↑)	250250																											
		ethyl phenylacetate (↑)	7590																											
		N-(phenylmethylene)-1-butanamine (↑)	296031																											
		indole (↑)	798																											
		1-methyl-naphthalene (↑)	7002																											
		1-decanol	8174																											
		indole	798																											
		1-dodecanol	8193																											
		acetic acid	176																											
		1-decanol	8174																											
		Escherichia coli	Gram negative bacteria	trimethylamine	1146	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	24 h	unknown	unknown	Thorn, R., Reynolds, D. M. and Greenman, J. (2011) Multivariate analysis of bacterial volatile compound profiles for discrimination between selected species and strains in vitro.Journal of Microbiological Methods, 84 (2), pp. 258-264. ISSN 0167-7012 DOI: 10.1016/j.jmimet.2010.12.001.	J. Microbiol. Methods	2011									
				2-aminoacetophenone	11952																									
				indole	798		GC-MS analysis	not used	not used	not used	not used	not used	not used									not used	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	unknown	unknown	unknown	Boots AW, Smolinska A, van Berkel JJ, Fijten RR, Stobberingh EE, et al. (2014) Identification of microorganisms based on headspace analysis of volatile organic compounds by gas chromatography-mass spectrometry. J Breath Res 8: 027106. doi:10.1088/1752-7155/8/2/027106.	J. Breath Res.	2014
				ethanol	702																									
				acetone	180																									
				2-nonanone	13187																									
				2-heptanone	8051																									
				1-octanol	957																									
				1-decanol	8174																									
				1-dodecanol	8193																									
				2-undecanone	8163																									
				tridecen-2-one	53427438																									
				ethanol	702																									
				1-propanol	1031																									
				isopentanol	31260																									
				1-octanol	957																									
				9-decenol	25612																									
				1-decanol	8174																									
				indole	798																									
				1-dodecanol	8193																									
				(Z)-7-tetradecen-1-ol	5362795																									
				1-tetradecanol	8209																									
				dimethyl disulfide	12232																									
				ethanol	702																									
				2-nonanone	13187																									
				2-heptanone	8051																									
				pentyl cyclopropane	75640																									
				indole	798																									
				ethanol (↑)	702																									
				indole (↑)	798																									
				2-(methylthio)-ethanol (↑)	78925																									
				3-methylbutanal (↑)	11552																									
				dimethyl disulfide (↑)	12232																									
				methypyrazine (↑)	7976																									
				2-(methylthio)-ethanol (↑)	78925																									
				phenol (↑)	996																									
				dimethyl trisulfide (↑)	19310																									
				benzonitrile (↑)	7505																									
				2,3,5-trimethylpyrazine (↑)	26808																									
				N-(phenylmethylene)-methanamine (↑)	72954																									
				2-nonanone (↑)	13187																									
				N,N'-dibenzylidenedimethylenediamine (↑)	66033																									
				2-decanone (↑)	12741																									
				N-(phenylmethylene)-1-propanamine (↑)	250250																									
				ethyl phenylacetate (↑)	7590																									
				N-(phenylmethylene)-1-butanamine (↑)	296031																									
				indole (↑)	798																									
1-methyl-naphthalene (↑)	7002																													
1-decanol	8174																													
indole	798																													
1-dodecanol	8193																													
acetic acid	176																													
1-decanol	8174																													
Escherichia coli	Gram negative bacteria			trimethylamine	1146	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	24 h	unknown	unknown	Thorn, R., Reynolds, D. M. and Greenman, J. (2011) Multivariate analysis of bacterial volatile compound profiles for discrimination between selected species and strains in vitro.Journal of Microbiological Methods, 84 (2), pp. 258-264. ISSN 0167-7012 DOI: 10.1016/j.jmimet.2010.12.001.	J. Microbiol. Methods	2011									
				2-aminoacetophenone	11952																									
				indole	798		GC-MS analysis	not used	not used	not used	not used	not used	not used									not used	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	unknown	unknown	unknown	Boots AW, Smolinska A, van Berkel JJ, Fijten RR, Stobberingh EE, et al. (2014) Identification of microorganisms based on headspace analysis of volatile organic compounds by gas chromatography-mass spectrometry. J Breath Res 8: 027106. doi:10.1088/1752-7155/8/2/027106.	J. Breath Res.	2014
				ethanol	702																									
				acetone	180																									
				2-nonanone	13187																									
				2-heptanone	8051																									
				1-octanol	957																									
				1-decanol	8174																									
				1-dodecanol	8193																									
				2-undecanone	8163																									
				tridecen-2-one	53427438																									
				ethanol	702																									
				1-propanol	1031																									
				isopentanol	31260																									
				1-octanol	957																									
				9-decenol	25612																									
				1-decanol	8174																									
				indole	798																									
				1-dodecanol	8193																									
				(Z)-7-tetradecen-1-ol	5362795																									
				1-tetradecanol	8209																									
				dimethyl disulfide	12232																									
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				2-nonanone	13187																									
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				pentyl cyclopropane	75640																									
				indole	798																									
				ethanol (↑)	702																									
				indole (↑)	798																									
				2-(methylthio)-ethanol (↑)	78925																									
				3-methylbutanal (↑)	11552																									
				dimethyl disulfide (↑)	12232																									
				methypyrazine (↑)	7976																									
				2-(methylthio)-ethanol (↑)	78925																									
				phenol (↑)	996																									
				dimethyl trisulfide (↑)	19310																									
				benzonitrile (↑)	7505																									
				2,3,5-trimethylpyrazine (↑)	26808																									
				N-(phenylmethylene)-methanamine (↑)	72954																									
				2-nonanone (↑)	13187																									
				N,N'-dibenzylidenedimethylenediamine (↑)	66033																									
				2-decanone (↑)	12741																									
				N-(phenylmethylene)-1-propanamine (↑)	250250																									
				ethyl phenylacetate (↑)	7590																									
				N-(phenylmethylene)-1-butanamine (↑)	296031																									
				indole (↑)	798																									
		1-methyl-naphthalene (↑)	7002																											
		1-decanol	8174																											
		indole	798																											
		1-dodecanol	8193																											
		acetic acid	176																											
		1-decanol	8174																											
		Escherichia coli	Gram negative bacteria	trimethylamine	1146	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	24 h	unknown	unknown	Thorn, R., Reynolds, D. M. and Greenman, J. (2011) Multivariate analysis of bacterial volatile compound profiles for discrimination between selected species and strains in vitro.Journal of Microbiological Methods, 84 (2), pp. 258-264. ISSN 0167-7012 DOI: 10.1016/j.jmimet.2010.12.001.	J. Microbiol. Methods	2011									
				2-aminoacetophenone	11952																									
				indole	798		GC-MS analysis	not used	not used	not used	not used	not used	not used									not used	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	unknown	unknown	unknown	Boots AW, Smolinska A, van Berkel JJ, Fijten RR, Stobberingh EE, et al. (2014) Identification of microorganisms based on headspace analysis of volatile organic compounds by gas chromatography-mass spectrometry. J Breath Res 8: 027106. doi:10.1088/1752-7155/8/2/027106.	J. Breath Res.	2014
				ethanol	702																									
				acetone	180																									
				2-nonanone	13187																									
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				1-dodecanol	8193																									
				2-undecanone	8163																									
				tridecen-2-one	53427438																									
				ethanol	702																									
				1-propanol	1031																									
				isopentanol	31260																									
				1-octanol	957																									
				9-decenol	25612																									
				1-decanol	8174																									
				indole	798																									
				1-dodecanol	8193																									
				(Z)-7-tetradecen-1-ol	5362795																									
				1-tetradecanol	8209																									
				dimethyl disulfide	12232																									
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				2-nonanone	13187																									
				2-heptanone	8051																									
				pentyl cyclopropane	75640																									
				indole	798																									
				ethanol (↑)	702																									
				indole (↑)	798																									
				2-(methylthio)-ethanol (↑)	78925																									
				3-methylbutanal (↑)	11552																									
				dimethyl disulfide (↑)	12232																									
				methypyrazine (↑)	7976																									
				2-(methylthio)-ethanol (↑)	78925																									
				phenol (↑)	996																									
				dimethyl trisulfide (↑)	19310																									
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				2,3,5-trimethylpyrazine (↑)	26808																									
				N-(phenylmethylene)-methanamine (↑)	72954																									
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1-dodecanol	8193																													
acetic acid	176																													
1-decanol	8174																													
Escherichia coli	Gram negative bacteria			trimethylamine	1146	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	24 h	unknown	unknown	Thorn, R., Reynolds, D. M. and Greenman, J. (2011) Multivariate analysis of bacterial volatile compound profiles for discrimination between selected species and strains in vitro.Journal of Microbiological Methods, 84 (2), pp. 258-264. ISSN 0167-7012 DOI: 10.1016/j.jmimet.2010.12.001.	J. Microbiol. Methods	2011									
				2-aminoacetophenone	11952																									
				indole	798		GC-MS analysis	not used	not used	not used	not used	not used	not used									not used	ATCC 25922	blood agar plates;incubated overnight at 37°C; transfer to sterile Brain Heart Infusion broth, growth for 4h with constant agitation at 37°C	unknown	unknown	unknown	Boots AW, Smolinska A, van Berkel JJ, Fijten RR, Stobberingh EE, et al. (2014) Identification of microorganisms based on headspace analysis of volatile organic compounds by gas chromatography-mass spectrometry. J Breath Res 8: 027106. doi:10.1088/1752-7155/8/2/027106.	J. Breath Res.	2014
				ethanol	702																									
				acetone	180																									
				2-nonanone	13187																									
				2-heptanone	8051																									
				1-octanol	957																									
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				1-dodecanol	8193																									
				2-undecanone	8163																									
				tridecen-2-one	53427438																									
				ethanol	702																									
				1-propanol	1031																									
				isopentanol	31260																									
				1-octanol	957																									
				9-decenol	25612																									
				1-decanol	8174																									
				indole	798																									
				1-dodecanol	8193																									
				(Z)-7-tetradecen-1-ol	5362795																									
				1-tetradecanol	8209																									
				dimethyl disulfide	12232																									
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				2-nonanone	13187																									
				2-heptanone	8051																									
				pentyl cyclopropane	75640																									
				indole	798																									
				ethanol (↑)	702																									
				indole (↑)	798																									
				2-(methylthio)-ethanol (↑)	78925																									
				3-methylbutanal (↑)	11552																									
				dimethyl disulfide (↑)	12232																									
				methypyrazine (↑)	7976																									
				2-(methylthio)-ethanol (↑)	78925																									
				phenol (↑)	996																									
				dimethyl trisulfide (↑)	19310																									
				benzonitrile (↑)	7505																									
				2,3,5-trimethylpyrazine (↑)	26808																									
				N-(phenylmethylene)-methanamine (↑)	72954																									
				2-nonanone (↑																										

Pathogen	Classification	VOCs	PubChem ID	Methods	F	G	H	I	J	K	L	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range		Reference	Journal Code	Year	
					Saliva	Blood	Breath	Skin	Urine	Faeces	Milk				Value	Unit				
		Indole	798	MCC-IMS analysis								DSM 25944 and 12 clinical isolates	1% glucose enteric fermentation broth	unknown	unknown	unknown	Ohsaeki, T. and Teraoka, H. (2005). <i>Journal of Chromatography B</i> , 829, 117-125.			
		1-dodecanol	8193																	
		1-tetradecanol	8209																	
		1-decanol	8174																	
		5-methylheptan-3-one	7822																	
		2-phenylacetaldehyde	998																	
		ethanol	702																	
		nonanal	31289																	
		ammonia	222																	
		Indole	798																	
		1-octanol	957																	
		1-octanol (dimer)	not available																	
		methanol	887																	
		acetaldehyde	177																	
		ethanol	702																	
		methanethiol	878																	
		acetone	180																	
		acetic acid	176																	
		Indole	798																	
		dimethyl disulfide	12232																	
		dimethyl disulfide	12232																	
		dimethyl disulfide	12232																	
		dimethyl disulfide	12232																	
		dimethyl disulfide	12232																	
		dimethyl disulfide	12232																	
		1-propanol	1031																	
		ethanol	702																	
		ethanol	702																	
		dimethyl disulfide	12232																	
		methanethiol	878																	
trimethylamine	1146																			
ammonia	222																			
		hexane	8058	HS-GLC analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	clinical isolate 1 clinical isolate 2 clinical isolate 3 clinical isolate 4 clinical isolate 5 clinical isolate 6	YE8 medium	18h	unknown	unknown	Hayward, N. J., et al. "Development of specific tests for rapid detection of <i>Escherichia coli</i> and all species of <i>Proteus</i> in urine." <i>Journal of clinical microbiology</i> 6.3 (1977): 195-201.	J. Clin. Microbiol.	1977	
		2-methyl-1-butanol	8723																	
		1-butanol	263																	
		1-decanol	8174																	
		1-dodecanol	8193																	
		ethanol	702																	
		methanol	887																	
		1-propanol	1031																	
		octanol	957																	
		1-pentanol	6276																	
		phenylacetic acid	999																	
		propanoic acid	1032																	
		3-methylbutanal	11552																	
		acetaldehyde	177																	
		benzaldehyde	240																	
		formaldehyde	712																	
		hexanal	6184																	
		dodecane	8182																	
		2-heptanone	8051																	
		acetoin	179																	
		acetone	180																	
		1-methyl-naphthalene	7002																	
		2-methylnaphthalene	7055																	
		2-methylphenol	335																	
		phenol	996																	
		ethyl acetate	8857																	
		ethyl butanoate	7762																	
		ethyl phenylacetate	7590																	
		n-propyl acetate	7997																	
		propyl phenylacetate	221641																	
2-(methylthio)-ethanol	78925																			
dimethyl disulfide	12232																			
dimethyl trisulfide	19310																			
hydrogen sulfide	402																			
methanethiol	878																			
2,3,5-trimethylpyrazine	26808																			
2-aminoacetophenone	11952																			
3-methyl-1H-indole	6736																			
4-chloro-1H-indole	91345																			
acetonitrile	6342																			
benzonitrile	7505																			
indole	798																			
methylpyrazine	7976																			
N,N'-dibenzylideneethylenediamine	66033																			
N-butyl-1-phenylmethanamine	296011																			
N-phenylmethylene-1-propanamine	250250																			
N-phenyl(methylene-methanamine	73954																			
trimethylamine	1146																			
		2,2,4,4-tetramethyloctane (↑)	182333	GC-MS analysis								unknown	unknown	unknown	unknown	unknown	Bond, A., Vernon, A., Reade, S., Mayor, A., Wastling, J., Minetti, C., ... & Probert, C. (2015). PWE-173 Investigation of volatile organic compounds emitted from faeces for the diagnosis of giardiasis. <i>Gut</i> , 64(Suppl 1), A268-A268. DOI: http://dx.doi.org/10.15403/gut.2014.1121.243.abo	Gut	2015	
		acetic acid (↑)	176																	
		2,2,4,6,6-pentamethylheptane (↑)	26058																	
		cyclopentane (↑)	9253																	
		2-pentanone (↑)	7895																	
		2,6,6-trimethyldecane (↑)	545605																	
		1-propanol (↓)	1031																	
		3-methylfuran (↓)	13587																	
		1,3-bis (1,1-dimethylethyl) benzene (↑)	136810																	
		ethanol (↓)	702																	
		2,5-dimethylpyrazine (↑)	31252																	
		propanoic acid (↑)	1032																	
		pentanal (↑)	8063																	
		4-pentanone (↑)	7921																	
		2-hydroxy-3-pentanone (↑)	521790																	
		2,2,3,3-tetramethylpentane (↑)	92723																	
		5-ethylcyclopent-1-enecarboxaldehyde (↑)	580057																	
		(E)-2-octenal (↑)	528324																	
		ethyl isobutyrate (↓)	7342																	
		o-xylene (↓)	7237																	
		terpinolene (↓)	11463																	
		Indole	798																	
		benzaldehyde	240																	
		acetic acid	176																	
		phenylmethanol	244																	
		acetaldehyde (↑)	177																	
		butanal (↑)	261																	
		propanal (↑)	527																	
		1-butanol (↑)	263																	
		methanol (↑)	887																	
2,3-butanedione (↑)	650																			
2-pentanone (↑)	7895																			
		Indole	798	SPME-GC-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	clinical isolate	blood agar or chocolate blood agar	48h	unknown	unknown	Preti, George, et al. "Volatile compounds characteristic of sinus-related bacteria and infected sinus mucus: analysis by solid-phase microextraction and gas	Clin. Microbiol. Rev.	2009	
		benzaldehyde	240																	
		acetic acid	176																	
		phenylmethanol	244																	
		acetaldehyde (↑)	177																	
		butanal (↑)	261																	
		propanal (↑)	527																	
		1-butanol (↑)	263																	
		methanol (↑)	887																	
		2,3-butanedione (↑)	650																	
2-pentanone (↑)	7895																			

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A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T				
Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	cubation time before analy	Concentration	range	Reference	Journal Code	Year				
								In vivo Sample							Value	Unit							
Haemophilus influenzae	Gram negative bacterium	4-heptanone (↑)	31246	GC-MS analysis				x				clinical isolate	chocolate agar plates; liquid cultures: tryptic soy broth	3, 4.5, 6 and 7.5 h after inoculation	unknown	ppt-pptm	W. Filipiak, A. Sponring, M. M. Baur, C. Ager, A. Filipiak, H. Wiesenhofer, M. Nagl, J. Troppmair, A. Amann. Characterization of volatile metabolites taken up by or released from Streptococcus pneumoniae and Haemophilus influenzae by using GC-MS. Microbiology 2012, 158, 3044. doi: 10.1099/mic.0.062687-0.	Microbiol.	2012				
		acetic acid (↑)	176																				
		ethyl acetate (↑)	8857																				
		methyl methacrylate (↑)	6658																				
		vinyl butyrate (↑)	31247																				
		methyl propionate (↑)	11124																				
		3-(ethythio)propanal (↑)	229467																				
		dimethyl sulfide (↑)	1068																				
		methanethiol (↑)	878																				
		dimethyl disulfide (↑)	12232																				
		carbon disulfide (↑)	6348																				
		methyl thioacetate (↑)	519840																				
		ethyl methyl sulfide (↑)	12230																				
		dimethyl trisulfide (↑)	19310																				
		2-methyl-2-butene (↑)	10553																				
		isoprene (↑)	6557																				
		3-methyl-1-butene (↑)	11239																				
		o-hydroxybenzaldehyde (↑)	6998																				
		furan (↑)	8029																				
		2-acetyl-1,4,5,6-tetrahydropyridine (↑)	520194																				
		gamma-butyrolactone (↑)	7302																				
Helicobacter pylori	Gram negative bacterium	3-ethyl-6-pentamethyldisilyloctane	590048	GC-MS analysis								unknown	blood culture test tube over 48 h	one aliquot was first cultivated in SP4 broth (dilution of 1:10 for 24 h at 37°C); further diluted 1:50 (24 h), when the log phase was observed	unknown	unknown	Abd El Gader, A., Lieberman, D., Shemer Ami, Y., Svetodin, N., Lazavitch, T., Sagi, O., & Zeiri, Y. (2015). Volatile organic compounds generated by cultures of bacteria and viruses associated with respiratory infections. Biomedical Chromatography. DOI: 10.1002/bmc.3494.	Biomed. Chromatogr.	2015				
		heptane	8900																				
		5-[tri-t-butoxysilyl]-2-mercaptoethylamine	6058																				
		methylcyclohexane	7962																				
		4-fluorohistamine	541569																				
		isopentanol	31260																				
		7-methyl-1,8-naphthyridin-2-amine	594420																				
		ethylpentamethyldisiloxane	20667832																				
		5-methylthienol(3,2-b)pyridine	591057																				
		2,5-bis(trimethylsilyloxy)-benzaldehyde	622536																				
		1,2-bis(trimethylsilyl)benzene	519794																				
		decamethyl tetrasiloxane	8852																				
		isobutane	6360		SPME-GC-MS analysis				x					unknown	unknown	unknown	unknown	Sethi, S., Nanda, R., & Chakraborty, T. (2013). Clinical application of volatile organic compound analysis for detecting infectious diseases. Clinical microbiology reviews , 26 (3), 462-475. doi: 10.1128/CMR.00020-13.	Clin. Microbiol. Rev.	2013			
		2-butanone	6569																				
		ethyl acetate	8857																				
		hydrogen cyanide	768			PTR-MS analysis				x					NCTC 11637	unknown	unknown	unknown					
		hydrogen nitrate	944																				
		dimethylether	8254																				
		2,3-butadiene	not available																				
		acetaldehyde	177																				
		ethanol	702																				
isobutane	6360																						
acetonitrile	6342																						
n-butane	7843																						
acetone	180																						
2-propanol	3776																						
ethylether	3283																						
isoprene	6557																						
n-pentane	8003																						
2-methylfuran	10797	SPME-GC-MS analysis						x					clinical isolate	air sample transferred from a Tedlar bag to a glass vial; SPME fiber was inserted into the vial and exposed to the gaseous mixture	15 min	unknown	unknown	Ulanowska, A., Kowalkowski, T., Hryniewicz, K., Jackowski, M., & Buszewski, B. (2011). Determination of volatile organic compounds in human breath for Helicobacter pylori detection by SPME-GC/MS. Biomedical Chromatography , 25 (3), 391-397. DOI 10.1002/bmc.1460.	Biomed. Chromatogr.	2011			
2-butanone	6569																						
ethyl acetate	8857																						
2-methylpentane	7892																						
3-methylpentane	7282																						
benzene	241																						
methylcyclopentane	7296																						
hexane	8058																						
toluene	1140																						
2-hexanone	11583																						
ethylbenzene	7500																						
p-xylene	7809																						
styrene	7501																						
benzaldehyde	240																						
nonane	8141																						
propane	6334		SPME-GC-MS analysis										unknown	isolation from patient stomach mucous membrane biopsies; selective medium BD BBLTM Stacker Plates; culture at 37°C in microaerophilic conditions 5-6 days; suspension of isolated bacteria in sterile water for analysis	1 h	unknown	unknown						
acetaldehyde	177																						
ethanol	702																						
methanethiol	878																						
(E)-2-butene	62695																						
isobutane	6360																						
2-methylpropene	8255																						
acetonitrile	6342																						
n-butane	7843																						
pentafluoroethane	9633																						
acetone	180																						
carbon disulfide	6348																						
2-propanol	3776																						
ethylether	3283																						
methyl acetate	6584																						
dichlorofluoroethane	15586																						
2-methylbutane	6556																						
2-pentene	12585																						
n-pentane	8003																						
cyclopentane	9253																						
2-methylpropanal	6561																						
trichloromethane	6212																						
2-butanone	6569																						
ethyl acetate	8857																						
2-methyl-1-pentene	12986																						
isobutanol	6560																						
2-methylpentane	7892																						
3-methylpentane	7282																						
benzene	241																						
methylcyclopentane	7296																						
hexane	8058																						
cyclohexane	8078																						
toluene	1140																						
mercaptoacetone	520144																						
3-methylbutanal	11552																						
2-ethoxy-2-methylpropane	12512																						
dimethyl disulfide	12232																						
1-pentanol	6276																						
4-methylpentane	not available																						
methylcyclohexane	7962																						
tetrahydro-2,2,4,4-tetramethylfuran	27010																						
ethylbenzene	7500																						
styrene	7501																						
2,4-dimethyl-1-heptene	123385																						
octane	356																						
3,5-dimethyloctane	139989																						

[illegible]

A		B		C		D		E		F		G		H		I		J		K		L		M		N		O		P		Q		R		S		T																																																																																																																																																																																																																																																																																																																																																																																																																																																															
Pathogen		Classification		VOCs		PubChem ID		Methods		Saliva		Blood		Breath		Skin		Urine		Faeces		Milk		Bacterial strain		Culture conditions/Growth medium		Incubation time before analysis		Concentration range		Reference		Journal Code		Year																																																																																																																																																																																																																																																																																																																																																																																																																																																																	
										Value		Unit														Value		Unit																																																																																																																																																																																																																																																																																																																																																																																																																																																																									
Morganella morganii		Gram negative bacterium		dimethyl trisulfide		139310		SPME-GC-MS analysis		unknown		unknown		unknown		unknown		unknown		unknown		unknown		WILD 10257		brain-heart-infusion broth		overnight incubation 37°C		unknown		unknown		diagnostic tool for the detection of pathogenic bacteria. TrAC Trends in Analytical Chemistry, 53, 117-125.		TrAC, Trends Anal. Chem.		2014																																																																																																																																																																																																																																																																																																																																																																																																																																																															
				1-decanol		8174				1-dodecanol		8193		phenol		996		1-methyl-naphthalene		7002		1,4-dimethylcyclohexane		11523		1,3-nubenzofurandione		6881		2,3-dimethylpentane		11260		acetaldehyde		177		phenylmethanol		244		1,1,3-trimethyl-3-phenyl-cyclohexane		not available		tridecane		12388		3,7-dimethyldodecane		28468		5-ethyl-2-methyl-heptane		26056		1,3,5-trimethylbenzene		7947		4,6,8-trimethyl-1-nonene		41077		hexacyclohexane		20283		4-methyl-1-hexene		19589		bis-(3,5,5-trimethylhexyl) phthalate		34277		4-methyldodecane		521958		3-(1-methylethyl)oxetane		543882		3-(1-methylethyl)oxetane		543882		4-methyldodecane		521958		hexacyclohexane		20283		bis-(3,5,5-trimethylhexyl) phthalate		34277		1,3,5-trimethylbenzene		7947		3,7-dimethyldodecane		28468		tridecane		12388		4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		1-methyl-naphthalene		7002		3-heptanone		7802		methyldichlorododecane		524446		2,2,4,6,6-pentamethylheptane		26558		cymol		7463		1,4-dimethylcyclohexane		11523		methyl nicotinate		7151		methyl 4-anisate		8499		2-phenylanisol		6835		4-methylanisol		7731		ethyl 4-anisate		60979		trimethylxazole		30215		methyl 2-aminobenzoate		8635		benzothiazole		7222		4-hydroxy-4-methyl-2-pentanone		21256		3-methyl-4-pentanone		248934		4-methyl-5-hexanolide		544628		dimethylpentanolide		not available		cyclic proline-glycine		456653		2-phenylethanol		6054		methyl benzoate		7150		4-pentanone		7921		methyl phenylacetate		7559		methyl 2-furoate		11902		methyl salicylate		4133		camphor		2537		methylbutenolide		not available		methyl dimethylbenzoate		32786		phenylmethanol		244		ethyl benzoate		7165		methyl phenylacetate		7559		methyl 4-anisate		8499		methyl nicotinate		7151		2-phenylanisol		6835		o-xylene (↑)		7237		isopropyl acetate (↑)		7935		3-pentanol (↓)		11428		dimethylstyrene (↓)		62385		cymol (↓)		7463		o-xylene		7237		isopropyl acetate		7935		3-pentanol		11428		dimethylstyrene		62385		cymol		7463		camphene		6616		beta-pinene		14896		1,3,5-trimethylbenzene		7947		1-methyl-naphthalene		7002		tridecane		12388		2-butyloctanol		19800		4-methyldodecane		521958		methyl nicotinate		7151		methyl phenylacetate		7559		methyl 4-anisate		8499		2-phenylanisol		6835		3-(1-methylethyl)oxetane		543882		4-methyldodecane		521958		hexacyclohexane		20283		bis-(3,5,5-trimethylhexyl) phthalate		34277		1,3,5-trimethylbenzene		7947		3,7-dimethyldodecane		28468		tridecane		12388		4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane		19041		styrene		7501		1,2-dimethylcyclopropane		102832		2-methylpropanal		6561		methacrolein		6562		N-2-dimethyl-1-propanamine		12249		3-methylbutanal		11552		cyclohexanone		280		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane			
				1,3,5-trimethylbenzene		7947				4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		1-methyl-naphthalene		7002		3-heptanone		7802		methyldichlorododecane		524446		2,2,4,6,6-pentamethylheptane		26558		cymol		7463		1,4-dimethylcyclohexane		11523		methyl nicotinate		7151		methyl 4-anisate		8499		2-phenylanisol		6835		4-methylanisol		7731		ethyl 4-anisate		60979		trimethylxazole		30215		methyl 2-aminobenzoate		8635		benzothiazole		7222		4-hydroxy-4-methyl-2-pentanone		21256		3-methyl-4-pentanone		248934		4-methyl-5-hexanolide		544628		dimethylpentanolide		not available		cyclic proline-glycine		456653		2-phenylethanol		6054		methyl benzoate		7150		4-pentanone		7921		methyl phenylacetate		7559		methyl 2-furoate		11902		methyl salicylate		4133		camphor		2537		methylbutenolide		not available		methyl dimethylbenzoate		32786		phenylmethanol		244		ethyl benzoate		7165		methyl phenylacetate		7559		methyl 4-anisate		8499		methyl nicotinate		7151		2-phenylanisol		6835		o-xylene (↑)		7237		isopropyl acetate (↑)		7935		3-pentanol (↓)		11428		dimethylstyrene (↓)		62385		cymol (↓)		7463		o-xylene		7237		isopropyl acetate		7935		3-pentanol		11428		dimethylstyrene		62385		cymol		7463		camphene		6616		beta-pinene		14896		1,3,5-trimethylbenzene		7947		1-methyl-naphthalene		7002		tridecane		12388		2-butyloctanol		19800		4-methyldodecane		521958		methyl nicotinate		7151		methyl phenylacetate		7559		methyl 4-anisate		8499		2-phenylanisol		6835		3-(1-methylethyl)oxetane		543882		4-methyldodecane		521958		hexacyclohexane		20283		bis-(3,5,5-trimethylhexyl) phthalate		34277		1,3,5-trimethylbenzene		7947		3,7-dimethyldodecane		28468		tridecane		12388		4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane		19041		styrene		7501		1,2-dimethylcyclopropane		102832		2-methylpropanal		6561		methacrolein		6562		N-2-dimethyl-1-propanamine		12249		3-methylbutanal		11552		cyclohexanone		280		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane																																																																																																											
				1,3,5-trimethylbenzene		7947				4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		1-methyl-naphthalene		7002		3-heptanone		7802		methyldichlorododecane		524446		2,2,4,6,6-pentamethylheptane		26558		cymol		7463		1,4-dimethylcyclohexane		11523		methyl nicotinate		7151		methyl 4-anisate		8499		2-phenylanisol		6835		4-methylanisol		7731		ethyl 4-anisate		60979		trimethylxazole		30215		methyl 2-aminobenzoate		8635		benzothiazole		7222		4-hydroxy-4-methyl-2-pentanone		21256		3-methyl-4-pentanone		248934		4-methyl-5-hexanolide		544628		dimethylpentanolide		not available		cyclic proline-glycine		456653		2-phenylethanol		6054		methyl benzoate		7150		4-pentanone		7921		methyl phenylacetate		7559		methyl 2-furoate		11902		methyl salicylate		4133		camphor		2537		methylbutenolide		not available		methyl dimethylbenzoate		32786		phenylmethanol		244		ethyl benzoate		7165		methyl phenylacetate		7559		methyl 4-anisate		8499		methyl nicotinate		7151		2-phenylanisol		6835		o-xylene (↑)		7237		isopropyl acetate (↑)		7935		3-pentanol (↓)		11428		dimethylstyrene (↓)		62385		cymol (↓)		7463		o-xylene		7237		isopropyl acetate		7935		3-pentanol		11428		dimethylstyrene		62385		cymol		7463		camphene		6616		beta-pinene		14896		1,3,5-trimethylbenzene		7947		1-methyl-naphthalene		7002		tridecane		12388		2-butyloctanol		19800		4-methyldodecane		521958		methyl nicotinate		7151		methyl phenylacetate		7559		methyl 4-anisate		8499		2-phenylanisol		6835		3-(1-methylethyl)oxetane		543882		4-methyldodecane		521958		hexacyclohexane		20283		bis-(3,5,5-trimethylhexyl) phthalate		34277		1,3,5-trimethylbenzene		7947		3,7-dimethyldodecane		28468		tridecane		12388		4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane		19041		styrene		7501		1,2-dimethylcyclopropane		102832		2-methylpropanal		6561		methacrolein		6562		N-2-dimethyl-1-propanamine		12249		3-methylbutanal		11552		cyclohexanone		280		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane																																																																																																											
				1,3,5-trimethylbenzene		7947				4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		1-methyl-naphthalene		7002		3-heptanone		7802		methyldichlorododecane		524446		2,2,4,6,6-pentamethylheptane		26558		cymol		7463		1,4-dimethylcyclohexane		11523		methyl nicotinate		7151		methyl 4-anisate		8499		2-phenylanisol		6835		4-methylanisol		7731		ethyl 4-anisate		60979		trimethylxazole		30215		methyl 2-aminobenzoate		8635		benzothiazole		7222		4-hydroxy-4-methyl-2-pentanone		21256		3-methyl-4-pentanone		248934		4-methyl-5-hexanolide		544628		dimethylpentanolide		not available		cyclic proline-glycine		456653		2-phenylethanol		6054		methyl benzoate		7150		4-pentanone		7921		methyl phenylacetate		7559		methyl 2-furoate		11902		methyl salicylate		4133		camphor		2537		methylbutenolide		not available		methyl dimethylbenzoate		32786		phenylmethanol		244		ethyl benzoate		7165		methyl phenylacetate		7559		methyl 4-anisate		8499		methyl nicotinate		7151		2-phenylanisol		6835		o-xylene (↑)		7237		isopropyl acetate (↑)		7935		3-pentanol (↓)		11428		dimethylstyrene (↓)		62385		cymol (↓)		7463		o-xylene		7237		isopropyl acetate		7935		3-pentanol		11428		dimethylstyrene		62385		cymol		7463		camphene		6616		beta-pinene		14896		1,3,5-trimethylbenzene		7947		1-methyl-naphthalene		7002		tridecane		12388		2-butyloctanol		19800		4-methyldodecane		521958		methyl nicotinate		7151		methyl phenylacetate		7559		methyl 4-anisate		8499		2-phenylanisol		6835		3-(1-methylethyl)oxetane		543882		4-methyldodecane		521958		hexacyclohexane		20283		bis-(3,5,5-trimethylhexyl) phthalate		34277		1,3,5-trimethylbenzene		7947		3,7-dimethyldodecane		28468		tridecane		12388		4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane		19041		styrene		7501		1,2-dimethylcyclopropane		102832		2-methylpropanal		6561		methacrolein		6562		N-2-dimethyl-1-propanamine		12249		3-methylbutanal		11552		cyclohexanone		280		ethanol		702		formaldehyde		712		methanethiol		878		indole		798		ethanol		702		dimethyl sulfide		1068		1,3,5-trifluorobenzene		9745		2,3,4-trimethylpentane		11269		2,2,5-trimethylcyclohexane																																																																																																											
				1,3,5-trimethylbenzene		7947				4,6,8-trimethyl-1-nonene		41077		5-ethyl-2-methyl-heptane		26056		4-methyl-1-hexene		19589		1-methyl-naphthalene		7002		3-heptanone		7802		methyldichlorododecane		524446		2,2,4,6,6-pentamethylheptane		26558		cymol		7463		1,4-dimethylcyclohexane		11523		methyl nicotinate		7151		methyl 4-anisate		8499		2-phenylanisol		6835		4-methylanisol		7731		ethyl 4-anisate		60979		trimethylxazole		30215		methyl 2-aminobenzoate		8635		benzothiazole		7222		4-hydroxy-4-methyl-2-pentanone		21256		3-methyl-4-pentanone		248934		4-methyl-5-hexanolide		544628		dimethylpentanolide		not available		cyclic proline-glycine		456653		2-phenylethanol		6054		methyl benzoate		7150																																																																																																																																																																																																																																																																																																																																																																																													

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	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T
	Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	in vivo Sample	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range	Reference	Journal Code	Year
																	Value	Unit		
1001			isoprene	6557																
1002			10-methyl-1-undecene	519941																
1003			pyrrole	8027																
1004			3-methyl-1H-pyrrole	12023																
1005			1-vinyl aziridine	21843																
1006			2,3-butanedione (↓)	650																
1007			benzaldehyde (↓)	240																
1008			acetaldehyde (↓)	177																
1009			methacrolein (↓)	6562																
1010			3-methylbutanal (↓)	11552																
1011			nonanal (↓)	31289																
1012			propanal (↓)	527																
1013			3-methyl-2-butenal (↓)	61020																
1014			acrolein (↓)	7847																
1015			butanal (↓)	261																
1016			2-methylpropanal (↓)	6561																
1017			octanal (↓)	454																
1018			isoprene	6557																
1019			3-methylbutanal (↓)	11552																
1020			2-methylbutanal (↓)	7284																
1021			2,3,3-trimethylpentane (↓)	11215																
1022			benzaldehyde (↓)	240																
1023			1-undecene	13190																
1024			2-pentene	12585																
1025			2,3-butanedione (↓)	650																
1026			2-butanone	6569																
1027			2-heptanone	8051																
1028			2-nonanone	13187																
1029			1-methyl-4-(1-methylethenyl)cyclohexane (↓)	14299																
1030			ethanol	702																
1031			acetone	180																
1032			2-butanone	6569																
1033			2-pentanone	7895																
1034			isoprene	6557																
1035			2-aminoacetophenone	11952																
1036			dimethyl sulfide	1068																
1037			dimethyl disulfide	12232																
1038			dimethyl trisulfide	19310																
1039			methyl thiocyanate	11168																
1040			methyl isopropyl ketone	11251																
1041			acetophenone	7410																
1042			methyl thioacetate	519840																
1043			methyl thiobutanoate	62444																
1044			hydrogen cyanide	768																
1045			acetonitrile	6342																
1046			ethanol	702																
1047			acetone	180																
1048			acetic acid	176																
1049			ethylene glycol	174																
1050			2-pentanone	7895																
1051			4-methylphenol	2879																
1052			indole	798																
1053			2-aminoacetophenone	11952																
1054			2-nonanone	13187																
1055			2-undecanone	8163																
1056			2-aminoacetophenone	11952																
1057			acetic acid	176																
1058			acetone	180																
1059			acetonitrile	6342																
1060			ammonia	222																
1061			2-butanone	6569																
1062			dimethyl sulfide	1068																
1063			dimethyl disulfide	12232																
1064			ethanol	702																
1065			hydrogen cyanide	768																
1066			isoprene	6557																
1067			methanol	887																
1068			methanethiol	878																
1069			hydrogen cyanide (↑)	768																
1070			methyl thiocyanate	11168																
1071			2-aminoacetophenone	11952																
1072																				
1073																				
1074			ethanol (↑)	702																
1075			3-methylbutanal (↑)	11552																
1076			dimethyl disulfide (↑)	12232																
1077			isopentanol (↑)	31260																
1078			benzonitrile (↑)	7505																
1079			1-undecene (↑)	13190																
1080			2-nonanone (↑)	13187																
1081			acetone	180																
1082			2-phenylacetaldehyde	998																
1083			ammonia	222																
1084			5-methylheptan-3-one	7822																
1085			nonanal	31289																
1086			ammonia (dimer)	not available																
1087			dodecane	8182																
1088			2-ethyl-1-hexanol	7720																
1089			acetone	180																
1090			2-phenylacetaldehyde	998																
1091			ammonia	222																
1092			5-methylheptan-3-one	7822																
1093			nonanal	31289																
1094			ammonia (dimer)	not available																
1095			2-nonanone	13187																
1096			2,4-dimethyl-1-heptene	123385																
1097			1-heptene	11610																
1098			isopentanol	31260																
1099			limonene	22311																
1100			ethane (↑)	6324																
1101			propane	6334																
1102			n-pentane (↑)	8023																
1103			methanol (↓)	887																
1104			ethanol (↓)	702																
1105			2-propanol (↓)	3776																
1106			acetone (↓)	180																
1107			isoprene (↓)	6557																
1108			benzene (↑)	241																
1109			toluene (↑)	1140																
1110			dimethyl sulfide (↓)	1068																
1111			limonene (↑)	22311																
1112			alpha-pinene	6654																
1113			2,2,6-trimethyloctane	522006																
1114																				

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T
Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range	Unit	Reference	Journal Code	Year
		dodecane	8182	GC-MS analysis				x				unknown	Blood agar; Mannitol Salt agar; Mac-Conkey agar	24h at 37°C	unknown	unknown	Goeminne, P. C., Vandendriessche, T., Van Eldere, J., Nicolai, B. M., Hertog, M. L., & Dupont, L. J. (2012). Detection of <i>Pseudomonas aeruginosa</i> in sputum headspace through volatile organic compound analysis. <i>Respir Res</i> , 13, 87. doi: 10.1186/1465-9921-13-87.	Respir. Res.	2012
		4-terpinenol	11230																
		1-undecene	13190																
		linalool	6549																
		2,6,7-trimethyldecane	43924																
		indole	798																
		toluene	1140																
		ethanol	702																
		acetoin	179																
		acetic acid	176																
		amylene hydrate	6405																
		carvophyllene	5322111																
		cis-1-p-menthanol	89437																
		2,5-dimethyl-2,5-hexanediol	8031																
		2-nonanone	13187																
		acetone	180																
		2-ethyl-1-hexanol	7720																
		2-heptanone	8051																
		2-ethoxy-2-methylpropane	12512																
		2-phenylethanol	6054																
		1-octen-3-ol	18827																
		4-methyloctane	16665																
		isoamyl acetate	31276																
		limonene	22311																
		eucalyptol	2758																
		6-methyl-2-heptanone	13572																
		thymol	6989																
		2-phenylacetaldehyde	998																
		2-hexanone	11583																
		2,4-dimethyl-1-heptene	123385																
		5-methyl-2-(1-methylethyl)-cyclohexanone	6986																
		2,4-dimethylheptane	16656																
		pyrrolidine	31268																
		2,6-dimethyl-7-octen-2-ol	29096																
		methyl thiocyanate	11168																
		hydrogen cyanide	768																
		methyl thiocyanate	11168																
		hydrogen cyanide	768																
		methyl thiocyanate	11168																
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Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	cubation time before analysis	Concentration range	Unit	Reference	Journal Code	Year
		methlyl thioacetate	519840	TD-GC-MS analysis	not used	not used	not used	not used	not used	not used	not used	ATCC 27853	Brain Heart Infusion (BHI)	16h, 24h and 48h	unknown	unknown	Neerincx, A. H., Geurts, B. P., Habets, M. F. J., Booi, J. A., van Loon, J., Jansen, J. J., ... & Wevers, R. A. (2016). Identification of <i>Pseudomonas aeruginosa</i> and <i>Aspergillus fumigatus</i> mono- and co-cultures based on volatile biomarker combinations. <i>Journal of breath research</i> , 10 (1), 016002.	J. Breath Res.	2016
		2-furaldehyde	7362																
		dimethyl trisulfide	19310																
		tetradecane	12389																
		1-undecene	13190																
		hexanal	6184																
		6-tridecane	142600																
		dimethyl disulfide	12232																
		butanal	261																
		3-methyl-1H-pyrrole	12023																
		2-methylbutanal	7284	SIFT-MS analysis	not used	not used	not used	not used	not used	not used	not used	ATCC 27853	blood culture bottles	24 h	unknown	unknown	Allardyce, Randall A., et al. "Detection of volatile metabolites produced by bacterial growth in blood culture media by selected ion	J. Microbiol. Methods	2006
		hydrogen sulfide	402																
		methanethiol	878																
		dimethyl sulfide	1068																
		carbon dioxide	280																
		ammonia	222	IMR-MS analysis	not used	not used	not used	not used	not used	not used	not used	ATCC 27853	blood agar plates	24 h	unknown	unknown	Dolch, M. E., et al. "Volatile compound profiling for the identification of Gram-negative bacteria by ion-molecule reaction-mass spectrometry." <i>Journal of</i>	J. Appl. Microbiol.	2012
		methanethiol	878																
		indole	798																
		2-aminoacetophenone	11952	SPME-GC-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	clinical isolate	blood agar or chocolate blood agar	48h	unknown	unknown	Pretl, George, et al. "Volatile compounds characteristic of sinus-related bacteria and infected sinus mucus: analysis by solid-phase microextraction and gas chromatography-mass spectrometry." <i>Journal</i>	Clin. Microbiol. Rev.	2009
		dimethyl disulfide	12232																
		1-undecene	13190																
		2,5-dimethylpyrazine	31252																
		dimethyl sulfide	1068																
		isoprene	6557	CG-FID analysis	not used	not used	not used	not used	not used	not used	not used	ATCC 10145	minimal salt AB medium+ 1% citrate	overnight	unknown	unknown	Schöller, Charlotte, Søren Molin, and Ken Wilkins. "Volatile metabolites from some gram-negative bacteria." <i>Chemosphere</i> 35.7 (1997): 1487-1495.	Chemosphere	1997
		dimethyl disulfide	12232																
		dimethyl trisulfide	19310																
		1-undecene	13190																
		hydrogen cyanide	768																
		ammonia	222	SIFT-MS analysis				x				clinical isolate	blood agar (BA) and <i>Pseudomonas</i> selective media (PSM)	48h at 37°C	unknown	ppb	Carroll, Will, et al. "Detection of volatile compounds emitted by <i>Pseudomonas aeruginosa</i> using selected ion flow tube mass spectrometry." <i>Pediatric pulmonology</i> 39.5 (2005): 452-456.	Pediatr. Pulmonol.	2005
		acetonitrile	6342																
		dimethyl disulfide	12232																
		ethanol	702																
		2-aminoacetophenone	11952		GC-MS/ Colorimetric analysis	not used	not used	not used	not used	not used	not used	not used	ATCC 15692	blood agar plates	20h	unknown	unknown	Cox, Charles D., and J. Parker. "Use of 2-aminoacetophenone production in	J. Clin. Microbiol.
		dimethyl disulfide	12232		unknown	unknown	unknown	unknown	unknown	unknown	unknown	10 clinical isolates							
		dimethyl trisulfide	19310																
		2-nonanone	13187																
		2-undecanone	8163																
		methanethiol	878																
		2-aminoacetophenone	11952																
		1-butanol	263																
		toluene	1140																
		2-butanone	6569																
		1-undecene	13190																
		isopentanol	31260																
		dimethyl disulfide	12232																
		dimethyl trisulfide	19310																
		2-nonanone	13187																
		2-undecanone	8163																
		methanethiol	878																
		2-aminoacetophenone	11952																
		1-butanol	263																
		toluene	1140																
		2-butanone	6569																
		1-undecene	13190																
		isopentanol	31260																
		dimethyl disulfide	12232																
		dimethyl trisulfide	19310																
		2-nonanone	13187																
		2-undecanone	8163																
		methanethiol	878																
		2-aminoacetophenone	11952																
		1-butanol	263																
		toluene	1140																
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		1-undecene	13190																
		isopentanol	31260																
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		1-undecene	13190																
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		methanethiol	878																
		2-aminoacetophenone	11952																
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		toluene	1140																
		2-butanone	6569																
		1-undecene	13190																
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		dimethyl disulfide	12232																
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		2-nonanone	13187																
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		methanethiol	878																
		2-aminoacetophenone	11952																
		1-butanol	263																
		toluene	1140																
		2-butanone	6569																
		1-undecene	13190																
		isopentanol	31260																
		dimethyl disulfide	12232																
		dimethyl trisulfide	19310																
		2-nonanone	13187																
		2-undecanone	8163																
		methanethiol	878																
		2-aminoacetophenone	11952																
		1-butanol	263																
		toluene	1140																
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		1-undecene	13190																
		isopentanol	31260																
		dimethyl disulfide	12232																
		dimethyl trisulfide	19310																
		2-nonanone	13187																
		2-undecanone	8163																
		methanethiol	878																
		2-aminoacetophenone	11952																
		1-butanol	263																
		toluene	1140																
		2-butanone	6569																
		1-undecene	13190																
		isopentanol	31260																

A		B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	
Pathogen		Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	In vivo Sample	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range	Reference	Journal Code	Year	
																	Value	Unit			
1300			dimethyl trisulfide	19310										ATCC 17423							
1301			2-nonanone	13187																	
1302			2-undecanone	8163																	
1303			methanethiol	878										ATCC 7701							
1304			2-aminoacetophenone	11952																	
1305			1-butanol	263																	
1306			toluene	1140										CDC 9104							
1307			2-butanone	6569																	
1308			1-undecene	13190																	
1309			isopentanol	31260										CDC 9171							
1310			dimethyl disulfide	12232																	
1311			dimethyl trisulfide	19310																	
1312			2-nonanone	13187																	
1313			2-undecanone	8163																	
1314			methanethiol	878																	
1315			2-aminoacetophenone	11952																	
1316			1-butanol	263																	
1317			toluene	1140																	
1318			2-butanone	6569																	
1319			1-undecene	13190																	
1320			isopentanol	31260																	
1321			dimethyl disulfide	12232																	
1322			dimethyl trisulfide	19310																	
1323			2-nonanone	13187																	
1324			2-undecanone	8163																	
1325			methanethiol	878																	
1326			2-aminoacetophenone	11952																	
1327			1-butanol	263																	
1328			toluene	1140																	
1329			2-butanone	6569																	
1330			1-undecene	13190																	
1331			isopentanol	31260																	
1332			dimethyl disulfide	12232																	
1333			dimethyl trisulfide	19310																	
1334			2-nonanone	13187																	
1335			2-undecanone	8163																	
1336			methanethiol	878																	
1337			2-aminoacetophenone	11952																	
1338			1-butanol	263																	
1339			toluene	1140																	
1340			2-butanone	6569																	
1341			1-undecene	13190																	
1342			isopentanol	31260																	
1343			dimethyl disulfide	12232																	
1344			dimethyl trisulfide	19310																	
1345			2-nonanone	13187																	
1346			2-undecanone	8163																	
1347			methanethiol	878																	
1348			2-aminoacetophenone	11952																	
1349			1-butanol	263																	
1350			toluene	1140																	
1351			2-butanone	6569																	
1352			1-undecene	13190																	
1353			isopentanol	31260																	
1354			dimethyl disulfide	12232																	
1355			dimethyl trisulfide	19310																	
1356			2-nonanone	13187																	
1357			2-undecanone	8163																	
1358			methanethiol	878																	
1359			2-aminoacetophenone	11952																	
1360			1-butanol	263																	
1361			toluene	1140																	
1362			2-butanone	6569																	
1363			1-undecene	13190																	
1364			isopentanol	31260																	
1365			dimethyl disulfide	12232																	
1366			dimethyl trisulfide	19310																	
1367			2-nonanone	13187																	
1368			2-undecanone	8163																	
1369			methanethiol	878																	
1370			2-aminoacetophenone	11952																	
1371			1-butanol	263																	
1372			toluene	1140																	
1373			2-butanone	6569																	
1374			1-undecene	13190																	
1375			isopentanol	31260																	
1376			dimethyl disulfide	12232																	
1377			dimethyl trisulfide	19310																	
1378			2-nonanone	13187																	
1379			2-undecanone	8163																	
1380			methanethiol	878																	
1381			2-aminoacetophenone	11952																	



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2	Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	Skin	Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	incubation time before analysis	Concentration range	Unit	Reference	Journal Code	Year																											
1377	Staphylococcus aureus	Gram positive bacterium	ethyl isovalerate	7945																																											
1378			isoamyl acetate	31276																																											
1379			ethyl formate	8025																																											
1380			methyl methacrylate	6658																																											
1381			methanethiol	878																																											
1382			dimethyl disulfide	12232																																											
1383			1,3-butadiene	7845																																											
1384			2-methylpropene	8255																																											
1385			n-butane	7843																																											
1386			(Z)-2-butene	5287573																																											
1387			(E)-2-butene	62695																																											
1388			propane	6334																																											
1389			acetonitrile	6342																																											
1390			ethanol	702																																											
1391			1-butanol	263																																											
1392			acetone	180																																											
1393			acetic acid	176																																											
1394			ethylene glycol	174																																											
1395			isopentanol	31260																																											
1396			pyrimidine	9260																																											
1397			2-pentanone	7895																																											
1398			4-methylphenol	2879																																											
1399			2-nonanone	13187																																											
1399			acetaldehyde (↑)	177																	SESI-MS analysis	not used	not used	not used	not used	not used	not used	not used	unknown	unknown	unknown	unknown	Sohrabi M, Zhang L, Zhang K, Ahmetagic A, Wei MQ (2014) Volatile Organic Compounds as Novel Markers for the Detection of Bacterial Infections. Clin Microbiol 3: 151. doi:10.4172/2327-5073.1000151	J. Clin. Microbiol.	2014												
1400			2-methylpropene (↑)	8255																																											
1401			n-butane (↑)	7843																																											
1402			2-pentanone	7895																																											
1403			propanal (↑)	527																																											
1404			ethyl acetate (↑)	8857																																											
1405			methyl vinyl ketone	6570																																											
1406			hexanal	6184																																											
1407			1,3-butadiene (↑)	7845																																											
1408			benzaldehyde (↓)	240																																											
1409			4-heptanone	31246																																											
1410			dimethyl sulfide	1068																																											
1411			ethanol (↑)	702																																											
1412			propane (↑)	6334																																											
1413			3-methylbutanal (↑)	11552																																											
1414			methacrolein (↑)	6562																																											
1415			(Z)-2-butene (↑)	5287573																																											
1416			acetic acid (↑)	176																																											
1417			(E)-2-butene (↑)	62695																																											
1418			2,3-butanedione (↑)	650																																											
1419			carbon disulfide	6348																																											
1420			2-methylpropanal (↑)	6561																																											
1421			3-methyl-2-butenal (↑)	61020																																											
1422			butanal	261																																											
1423			n-propyl acetate	7997																																											
1424			(E)-2-pentene	5326161																																											
1425			acetoin (↑)	179																																											
1426			(E)-2-methyl-2-butenal (↑)	5321950																																											
1427			(Z)-2-methyl-2-butenal (↑)	10336																																											
1428			1-butanol (↑)	263																																											
1429			1-propanol	1031																																											
1430			2-ethylacrolein (↑)	70203																																											
1431			2-methylbutyl acetate	12209																																											
1432			isobutanol (↑)	6560																																											
1433			2-methyl-2-butene	10553																																											
1434			isopentanol (↑)	31260																																											
1435			dimethyl disulfide	12232																																											
1436	ethyl formate (↑)	8025																																													
1437	ethyl isovalerate (↑)	7945																																													
1438	ethyl butanoate	7762																																													
1439	1-hydroxy-2-propanone (↑)	8299																																													
1440	isoamyl butyrate	7795																																													
1441	isoamyl propionate	7772																																													
1442	isobutyl acetate	8038																																													
1443	isoamyl acetate (↑)	31276																																													
1444	3-methylbutanoic acid (↑)	10430																																													
1445	methanethiol (↑)	878																																													
1446	methyl methacrylate (↑)	6658																																													
1447	n-butyl acetate (↑)	31272																																													
1448	ethyl vinyl ether	8023																																													
1449	formaldehyde	712																																													
1450	2-methylbutanal	7284																																													
1451	methanethiol	878																																													
1452	ammonia	222																																													
1453	(E)-2-butene	62695	TD-GC-MS analysis	x							clinical sample	blood agar, chocolate agar and macConkey agar plates; overnight incubation	unknown	unknown	ppb	Filipiak, W., Beer, R., Sponring, A., Filipiak, A., Ager, C., Schiefecker, A., ... & Amann, A. (2015). Breath analysis for in vivo detection of pathogens related to ventilator-associated pneumonia in intensive care patients: a prospective pilot study. Journal of breath research , 9 (1), 016004. DOI: 10.1088/1752-7155/9/1/016004	J. Breath Res.	2015																													
1454	2-methylpropene	8255																																													
1455	n-butane	7843																																													
1456	propane	6334																																													
1457	ethylene glycol	174																																													
1458	2-butanol	6568																																													
1459	isobutane	6360																																													
1460	isopentanol	31260																																													
1461	ethanol	702																																													
1462	isobutanol	6560																																													
1463	2-methyl-1-butanol	8723																																													
1464	acetic acid	176																																													
1465	3-methylbutanoic acid	10430																																													
1466	phenylacetic acid	999																																													
1467	2-ethylacrolein	70203																																													
1468	(E)-2-methyl-2-butenal	5321950																																													
1469	2-methylbutanal	7284																																													
1470	methacrolein	6562																																													
1471	3-methylbutanal	11552																																													
1472	acetaldehyde	177																																													
1473	benzaldehyde	240																																													
1474	hexanal	6184																																													
1475	2,3-butanedione	650																																													
1476	2-nonanone	13187																																													
1477	2-heptanone	8051																																													
1478	1-hydroxy-2-propanone	8299																																													
1479	toluene	1140																																													
1480	n-butyl acetate	31272																																													
1481	ethyl formate	8025																																													
1482	ethyl isovalerate	7945																																													
1483	methyl methacrylate	6658																																													
1484	2-(methylthio)-ethanol	78925																																													
1485	acetonitrile	6342																																													
1486	ammonia	222																																													
1487	pyrimidine	9260																																													
1488	acetone	180																	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	NCTC 7447	sterile urine (20 ml) from healthy males inoculated to a concentration of between 10*7 and 10*9 cfu/mL	37°C for 6h	unknown	ppb	Storer, M. K., Hibbard-Melies, K., Davis, B., & Scotter, J. (2011). Detection of volatile compounds produced by microbial growth in urine by selected ion flow tube mass	J. Microbiol. Methods	2011														
1489	2-methylbutanal	7284																																													
1490	methacrolein	6562																																													
1491	3-methylbutanal	11552																																													
1492	acetaldehyde	177																																													
1493	benzaldehyde	240																																													
1494	hexanal	6184																																													
1495	2,3-butanedione	650																																													
1496	2-nonanone	13187																																													
1497	2-heptanone	8051																																													
1498	1-hydroxy-2-propanone	8299																																													
1499	toluene	1140																																													
1500	n-butyl acetate	31272																																													
1501	ethyl formate	8025																																													
1502	ethyl isovalerate	7945																																													
1503	methyl methacrylate	6658																																													
1504	2-(methylthio)-ethanol	78925																																													
1505	acetonitrile	6342																																													
1506	ammonia	222																																													
1507	pyrimidine	9260																																													
1508	acetone	180																																unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	Bos, L. D., Sterk, P. J., & Schultz, M. J. (2013). Volatile metabolites of pathogens: a systematic review. doi:10.1371/journal.ppat.1003311	PLOS	2013
1509	2-methylbutanal	7284																																													
1510	methacrolein	6562																																													
1511	3-methylbutanal	11552																																													
1512	acetaldehyde	177																																													
1513	benzaldehyde	240																																													
1514	hexanal	6184																																													
1515	2,3-butanedione	650																																													
1516	2-nonanone	13187																																													
1517	2-heptanone	8051																																													
1518	1-hydroxy-2-propanone	8299																																													
1519	toluene	1140																																													
1520	n-butyl acetate	31272																																													
1521	ethyl formate	8025																																													
1522	ethyl isovalerate	7945																																													
1523	methyl methacrylate	6658																																													
1524	2-(methylthio)-ethanol	78925																																													
1525	acetonitrile	6342																																													
1526	ammonia	222																																													
1527	pyrimidine	9260																																													
1528	acetone	180																																													

A		B		C		D		E		F		G	H	I		J	K	L	M		N		O		P		Q	R		S	T
Pathogen	Classification	VOCs	PubChem ID	Methods	Saliva	Blood	Breath	In vivo Sample		Urine	Faeces	Milk	Bacterial strain	Culture conditions/Growth medium	Incubation time before analysis	Concentration range		Reference	Journal Code	Year											
																Value	Unit														
Staphylococcus epidermidis	Gram positive bacterium	2-butanone	6569	SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	ATCC 14990	sterile urine (20 mL) from healthy males inoculated to a concentration of between 10 <sup>7</sup> and 10 <sup>9</sup> cfu/mL	37°C for 6h	unknown	ppb	Storer, M. K., Hibbard-Melles, K., Davis, B., & Scotter, J. (2011). Detection of volatile compounds produced by microbial growth in urine by selected ion flow tube mass spectrometry (SIFT-MS). <i>Journal of microbiological methods</i> , 87(1), 111-113. doi: 10.1016/j.jmimet.2011.06.012.	J. Microbiol. Methods	2011											
		2-pentanone	7895																												
		2-hexanone	11583																												
		acetaldehyde	177																												
		2-methylbutanal	7284																												
		ethyl acetate	8857																												
		n-propyl acetate	7997																												
		hydrogen sulfide	402																												
		dimethyl disulfide	12232																												
		dimethyl sulfide	1068																												
		ammonia	222																												
		indole	798																												
		benzaldehyde	240	SPME-GC-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	clinical isolate	blood agar or chocolate blood agar	48h	unknown	unknown	Preti, George, et al. "Volatile compounds characteristic of sinus-related bacteria and infected sinus mucus: analysis by solid-phase microextraction and gas chromatography-mass spectrometry." <i>Journal</i>	Clin. Microbiol. Rev.	2009											
		phenylmethanol	244																												
		2-phenylethanol	6054																												
		acetic acid	176																												
		methanethiol	878																												
		acetaldehyde (↑)	177																												
		2-butanal (↑)	447466																												
		propanal (↑)	527																												
		2-methylpropanal (↑)	6561																												
		3-methyl-2-butanal (↑)	61020																												
		butanal (↑)	261																												
		1-butanol (↑)	263																												
Streptococcus pneumoniae	Gram positive bacterium	ethanol (↑)	702	GC-MS analysis									clinical isolate (later identified as ATCC 4963)	blood agar plates and tryptic soy broth	3, 3.75, 4.5, 6 and 7.5 h after inoculation	unknown	ppt-ppm	W. Filiplak, A. Sponring, M. M. Baur, C. Ager, A. Filiplak, H. Wiesenhofer, M. Nagl, J. Troppmair, A. Amann. Characterization of volatile metabolites taken up by or released from <i>Streptococcus pneumoniae</i> and <i>Haemophilus influenzae</i> by using GC-MS. <i>Microbiology</i> 2012, 158, 3044. doi: 10.1099/mic.0.062687-0.	Microbiol.	2012											
		2,3-butanedione (↑)	650																												
		2-pentanone (↑)	7895																												
		2-nonanone (↑)	13187																												
		2-butanone (↑)	6569																												
		acetone (↑)	180																												
		acetic acid (↑)	176																												
		methoxy methacrylate (↑)	6658																												
		ethyl acetate (↑)	8857																												
		methanethiol (↑)	878																												
		dimethyl disulfide (↑)	12232																												
		carbon disulfide (↑)	6348																												
		dimethyl trisulfide (↑)	19310																												
		dimethyl sulfide (↑)	1068																												
		1,3-butadiene (↑)	7845																												
		2-methylpropene (↑)	8255																												
		(E)-2-butene (↑)	62695																												
		(Z)-2-butene (↑)	5287573																												
		2-methyl-1-butene (↑)	11240																												
		furan (↑)	8029																												
		3-(methylthio)propanal (↑)	18635																												
		ethylbenzene (↑)	7500																												
		3-phenylfuran (↑)	518802																												
		3-methylbutanal (↓)	11552																												
		hexanal (↓)	6184																												
		ethanol	702																												
		formaldehyde	712		SIFT-MS analysis	unknown	unknown	unknown	unknown	unknown	unknown	unknown									unknown	ATCC 49619	blood culture bottles	24 h	unknown	unknown	Allardyce, Randall A., et al. "Detection of volatile metabolites produced by bacterial growth in blood culture media by selected ion flow tube mass spectrometry (SIFT-MS)." <i>Journal of microbiological methods</i> 65.2 (2006): 361-365.	J. Microbiol. Methods	2006		
		acetaldehyde	177																												
		dimethyl sulfide	1068																												
		trimethylamine	1146																												
		indole	798																												
		2-aminoacetophenone	11952																												
		hexanal	6184																												
		ethanol	702																												
		3-methylbutanal	11552																												
		acetaldehyde	177																												
formaldehyde	712																														
hexanal	6184																														
unknown	unknown	acetone	180	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	unknown	Bos, L. D., Sterk, P. J., & Schultz, M. J. (2013). Volatile metabolites of pathogens: a systematic review. doi:10.1371/journal.ppat.1003311	PLOS	2013												
		2-pentylfuran	19602																												
		dimethyl sulfide	1068																												
		2-aminoacetophenone	11952																												
		benzonitrile	7505																												
		trimethylamine	1146																												