Identifying uncertainties during the determination of VOCs in breath analysis



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Breath analysis is an emerging scientific field with promising medical applications. Volatile organic compounds (VOCs) of exhaled air are targeted and analysed with the use of various analytical instruments including GC-MS, PTR-MS, SIFT-MS, IMS, spectroscopic techniques, e-nose and sensors; potential differences in the emitted concentrations of specific or unspecific (pattern of compounds) biomarker(s) are related to health status or disease. Human breath is a clean, inexhaustible and non-invasive source of endogenous information; this makes it ideal for sampling especially vulnerable people and children. Despite its numerous and wide promising applications, it remains in an infant stage, as it suffers for standardised practices, presenting several qualitative and quantitative pitfalls. The sample, the size, the chemical diversity of exhaled breath volatiles, the role of confounders, the various sampling methodologies and techniques, sample treatment and data interpretation are considered among others, important factors of uncertainties and need of special attention. In this context, an overview of the critical issues employed in breath analysis are highlighted and reviewed, towards the vision of development personal care handheld monitoring devices. The understanding of these manifold issues assists researchers to validate their experimental design, towards adopting standard practices in the field of breath gas analysis. The ultimate goal, is sometime in the near future, breath analysis to become a reliable tool in the clinical setting, both for medical experts and human individuals for early diagnosis, clinical and personal monitoring.

1. Introduction

The last decade, there has been an increased interest on the determination of Volatile Organic Compounds (VOCs) for a number of challenging and novel applications, in the areas of:

a) Bio-analysis (disease prognosis and monitoring) [1]

b) Environment (indoor/outdoor/atmospheric pollution, plant/soil/waste emissions, etc.) [2, 3]

5. Uncertainties

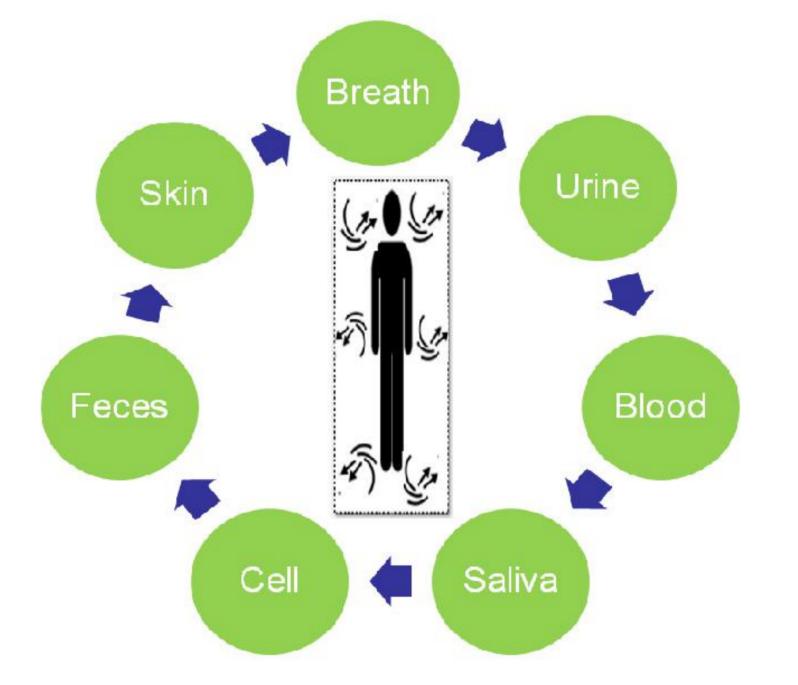
According to the Sampling and Standardization Interest Group of IABR [7], concentrations of VOCs in breath research are affected by the following factors [5-7]:

Storage, handling etc.)
Storage, handling etc.

c) Forensic and security (Urban Search and Rescue (USaR) applications) [4] The common core of these applications is the qualitative and quantitative determination of VOCs. These are defined as a large group of anthropogenic (xenobiotic) or biogenic organic compounds with relatively high vapor pressures. VOCs are emitted by all living organisms and can also be found in numerous domestic/industrial products. They can be potentially malodorous and hazardous for both human health (e.g. eye/nose irritation, headaches, nausea) and the environment (air, water, soil). The level of effect on humans and the environment varies greatly, depending on the substance, the amount and time of exposure, necessitating the identification and quantification of VOCs. The emission of VOCs creates a volatile chemical signature characterizing the source; this may lead to the detection of human presence and even in the noninvasive identification of human/plant diseases (e.g. microorganisms, infections).

2. The Human "Volatilome"

The study of the human volatilome includes the determination of VOCs released by body tissues and biological tissues, as presented in Figure 1. Their concentration is changing in pathological conditions enabling their use as novel medical diagnostic tools for various types of cancers, oxidative stress, asthma, diabetes, kidney-, liver- failure and many other medical disorders.



Storage transfer/transportation of breath samples (e.g. bags, containers, transfer lines, traps)

Pre-concentration/transfer of breath samples

Applied analytical method (e.g. selectivity and sensitivity of the method)

Physiology (e.g. hemodynamic effects, breathing pattern, paced breathing, breathing flow, distribution)

✤ Acute and previous exposure

- Nutrition, medication
- Pathology
- Genotype/phenotype, age and gender
- Effects/emissions of microorganisms
- Physicochemical properties of the VOC (in-out distribution)

Identification of potential VOC biomarkers (features, sum parameters, tentative, reference substance)

Quantitative assessment (rel. counts/areas vs. external calibration with reference substances)

✤ Units of VOC concentrations (nmol/L, nmol/m²BS, nmol/EtCO₂)

Data analyses/interpretation (e.g. in studies with a large number of potential VOC markers).

6. Conclusions [7]

• Although hundreds of VOCs have been described as potential biomarkers for diseases or disease states, however they have never been confirmed by independent studies and are therefore not viable

Figure 1: The human volatilome.

3. "Breathomics"

The early observations of Hippocrates, were confirmed with the analytical findings of Nobel Laureate Linus Pauling (1971) and therefore research on breath analysis ("breathomics") has been boosted; in parallel headspace analysis on the rest biological tissues followed. VOCs offer unique insights into on going biochemical processes in healthy and diseased humans [1]. Samples are non-invasively obtainable, as often as desired with acceptable discomfort, even during human dailies activities such as sleep, exercise, etc. It should be noted, that sampling procedure and methodology has been expanded with success to mammals, despite their size (e.g. mice, cow, dolphin). Nevertheless, the big challenge, remains that of development point of care medical diagnostics devices for personal monitoring.

4. Analytical Challenges

- Sample diversity
- Disease
- \blacktriangleright Analytical technique \rightarrow Inter-comparison

in the clinical setting.

 There are not "unique" or "exclusive" VOC breath markers for certain diseases.

The concentration changes of potential breath biomarkers may indicate pathological changes but may also reflect physiological effects, confounding variables (e.g. previous or acute exposure, medication, nutrition, microorganisms), methodological effects (sampling, storage, analytical set-up) and orgenetic/phenotype predisposition.

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