

<u>G. Romoaldo^{1,2}, A.Blidar^{1,3}, S. Hambye¹, V. Tagliatti², J-M. Colet², B. Blankert¹</u>

¹ Laboratory of Pharmaceutical Analysis-² Laboratory of Human Biology and Toxicology – Umons Research Institute for Health Sciences and Technology- Place du Parc, 20 7000 Mons, (Belgium) ³ Erasmus + Leave from Iuliu Hatieganu University of Medicine and Pharmacy Cluj / Napoca (Romania)

Assessing emotional states : Metabonomic profiling and quantification of a salivary biomarker

O1 BACKGROUND

02 METHODS

01.FACTS

Did you know that 1st cause of plane crash remains human mistake?

In 2015, 16 fatal aircraft accidents occasioned 560 deaths. Because pilots are actively engaged during each stage of a flight, there are numerous opportunities for this to go wrong, from failing to program the vital flightmanagement computer to miscalculating the required fuel uplift.

03.INNOVATIVE STRATEGY

In this work, two strategies are proposed to assess physiological impact of emotional states. The first one involves a targeted analysis with the quantification of a biomarker named 3-methoxy-4-Hydroxyphénylglycol (MHPG). The second one uses a non-targeted analysis: metabonomic profiling. We

01.BIOMARKER QUANTIFICATION

MHPG, the final metabolite Norepinephrine, is known to increase in **stressful situations**.^{1,2} MHPG salivary concentration is correlated to blood concentration³ which makes it a good candidate as a salivary biomarker of stress.



That's why stress and fatigue management remains a constant preoccupation in aeronautics in order to keep aerial people transportation **safe**.

Fig. 1 : Plane Accidents in commercial aviation - 2015

02.BIOVOC PROJECT

Biovoc project intends to objectify cognitive overload and mental fatigue of subjects using noninvasive methods. The goal of the project is to detect in pilots' voice recordings whether they are able to take control of the aircraft.

Our work aims on correlating physiological changes due to stress and emotions to voice signal modulations.

We chose saliva as a biofluid because its sampling is convenient and therefore limits bias due to experimental conditions.





Its electroactive properties allow us to quantify it accurately with electrochemical methods such as amperometry.

A UPLC-ECD/FLR method is under development for the quantification of salivary free MHPG in physiological range (10 - 15ng/mL).^{4,5}

Separation	UPLC Acquity Waters
Detection	-Amperometry (Roxy reactor cell – Antec ©) -FLR Acquity detector
Column	Acquity CSH Phenyl-Hexyl (Waters) 1.7 μm x 2.1 x 100 mm – T° : 30° C.
Mobile phase	iso. 10 : 90 (v/v) ACN:H ₂ 0 15mM Ammonium acetate
Electrochemical conditions	Working potential +0.7V Magic Diamond working electrode Pd/H ₂ reference
Analysis time	6 min
Flow rate	LC : 0.400 mL /min – ECD : 50μL/min

02.METABONOMIC PROFILING

The **metabolome** depicts a view of all metabolites from a cell, a tissue, an organ or an organism. It relies on DNA expression but also on environment.

In order to assess the evolution of salivary metabolome when the subject is placed in a stressful environment, we decided to adapt the Trier social stress test and observed the evolution of the metabolome at different moments of the test.

Saliva is prepared by microcentrifuge filtration and analyzed by ¹**H-NMR** 128 scans with NOESY-presat sequence to get rid of proteins signal. Principal Component analysis (PCA) on SimcaP+ allows us to compare subjects.

The Trier Social Stress Test (TSST)

TSST was created in 1993 to induce psychosocial stress for research purposes. In this adaptation, we sample saliva at different moments along the tasks. Heart rate is monitored during a part of the test. The subject is asked to answer the STAI-Y (State-Trait Anxiety Inventory, a self assessment anxiety questionnaire) 3 times during the test. This will let us know how anxious the subject feels.



O3 RESULTS

04 CONCLUSION & OUTLOOKS

01.Method development

MHPG quantification

A calibration curve has been plotted with different aqueous standards of MHPG signal obtained by ECD and FLR detection. The results are comparable on both detection modes in terms of sensitivity in the 5-50 ng/mL linearity range with a LOD at 6,7 and 7,2 ng/mL for ECD and FLR respectively.



Fig. 2 : Overlay of chromatograms obtained on UPLC-ECD for aqueous standards of MHPG and calibration curve



Fig. 3 : Overlay of chromatograms obtained on UPLC-FLR for aqueous standards of MHPG and calibration curve

Sample preparation

Saliva is a complex matrix that needs a pretreatment before injection. Here we compared 2 SLE and 1 SPE method to extract and concentrate MHPG.

For all protocols : dilution of the saliva pooled and spiked samples with H_20 1:1 and elution with ethylacetate (EA).



Fig. 4: Comparison of different sample preparation methods. Each figure represents the overlay of chromatograms obtained on UPLC-FLR for a growing concentration of MHPG

02.TSST Study (on doind

10 subjects on 30 adult males took part to the TSST. Datas need to be completed for building a solid analysis of salivary metabolomes.

A preliminary study was realized on student to assess the evolution of salivary metabolome related to an academic exam.

It's interesting to notice (*fig.5*) that we have been able to discriminate 2 groups using salivary metabolome -after academic exam -resting condition In fig. 6, are listed metabolites that could explain this group separation.

01. PROJECT PROGRESS

The multi-faceted approach proposed in this project is very innovative in stress studies. Based on targeted biomarker quantification and non-targeted saliva metabolomic profiling, this on going study aims to build a better knowledge of the impact of stress and emotions on biochemical pathways.

A UPLC – ECD/FLR method has been developed for quantification. Specificity and sensitivity MHPG seem high enough for determination of salivary free MHPG in physiological range (10-15ng/mL)

A study is conducted to assess the impact of stress on salivary metabolome of male subjects. We need to recruit more subjects to raise the robustness of the comparison.

Final optimization of the LC and sample prep methods has to be achieved. Once settled, the method will be validated in accordance of the method of accuracy profiles.

03. FURTHER DEVELOPMENTS

To get a more detailed view of **physiological impact** of emotions, a second salivary biomarker will be targeted (salivary α -amylase or DHEA for instance).

According to the results of TSST study, we will consider the use of sweat as biofluid for assessing the metabolome evolution.

The methods will be thereafter used in a driving simulation context to control its efficiency before being adapted to plane simulator.

The final step of this project will consist of linking the voice signal changes to physiological expressions of emotions.



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Preliminary results showed that the cleaner signal is obtained with SLE cartridge. SLE protocols will be further optimized to achieve a higher recovery and a better reproducibility.

SLE 1 : Isolute + Biotage © (Diateoms)	SLE 2 : Novum Luna Phenomenex © (Polymer)	SPE HLB Oasis Waters © (polymer)
Load 400µL sample	Load 1mL sample	1mL EA
Wait 5'	Wait 5'	1mL MeOH conditioning
900µL EA elution	5mL EA elution	1mL Water equilibrating
Wait 5'	Wait 5'	Load sample
900µL EA elution	5mL EA elution	H ₂ 0 Wash
Wait 5'	Wait 5'	1mL EA elution



The TSST study will be continued to assess the impact of environmental stress on salivary metabolomic profile.



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Contact information email gilson.romoaldo@umons.ac.be web www.umons.ac.be 18-25 June 2017



