



GC×GC-MS AND BAYESIAN TESTING IN FORENSICS: TOWARDS THE IDENTIFICATION OF SUSPECTS THROUGH THEIR ODOR

Isabelle RIVALS¹, Vincent CUZUEL², Guillaume COGNON², Roman Lecon
Didier THIEBAUT³, Charles SAULEAU², Jérôme VIAL³

Équipe de Statistique Appliquée, UMRS 1158, ESPCI Paris, France

Institut de Recherche Criminelle de la Gendarmerie Nationale, Cergy-Pontoise, France

UMR CBI 8231 - Laboratoire Sciences Analytiques, Bioanalytiques et Miniaturisation – ESPCI Paris – PSL Research University

- GCxGC 2018, Riva Del Garda, Italy

CONTEXT

Popularization of the techniques used by the police



Criminals are more attentive and cautious!



Human odor



USE OF TRAINED DOGS

- Sufficient for identification of a person
- Limited probative value in courts of justice
- Need for corroborative evidence by analytical tools:
 - **Support the information** provided by dogs
 - **Probative value** to evidence in courts of justice



OBJECTIVES

- Develop a global strategy to characterize the olfactory fingerprints of individuals using **analytical and statistical tools**
- Volatile compounds **at trace levels**: preconcentration step required
- Complex mixtures**: multidimensional separation (GC×GC-MS)

Question to be answered

- Is the comparison of an “odor” reference chromatogram to a chromatogram obtained using an odor sample from a suspect (crime scene...) sufficient to prove that the odor belongs to the same person?



GLOBAL STRATEGY

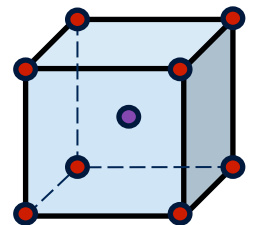
SAMPLING/PANEL

SEPARATION AND
DETECTION

DATA PROCESSING

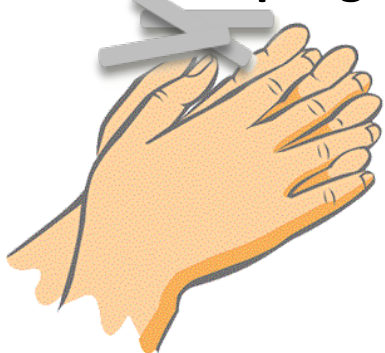


$$P(A|B) = \frac{P(B|A) P(A)}{P(B)}$$

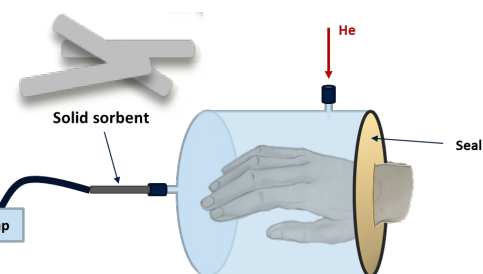


PRECONCENTRATION AND ANALYSIS: PURGE AND TRAP - GCxGC

Direct sampling



Indirect sampling



Thermodesorption coupled with **GCxGC-MS**

VSP4000, Action Europe (Sausheim,
France)

DB1MS-DB1701

2°C/min – modulation 8 s

Desorption optimization DC

- **synthetic mixture** of human odor (80 compounds ¹)
- **full factorial design** 2⁴



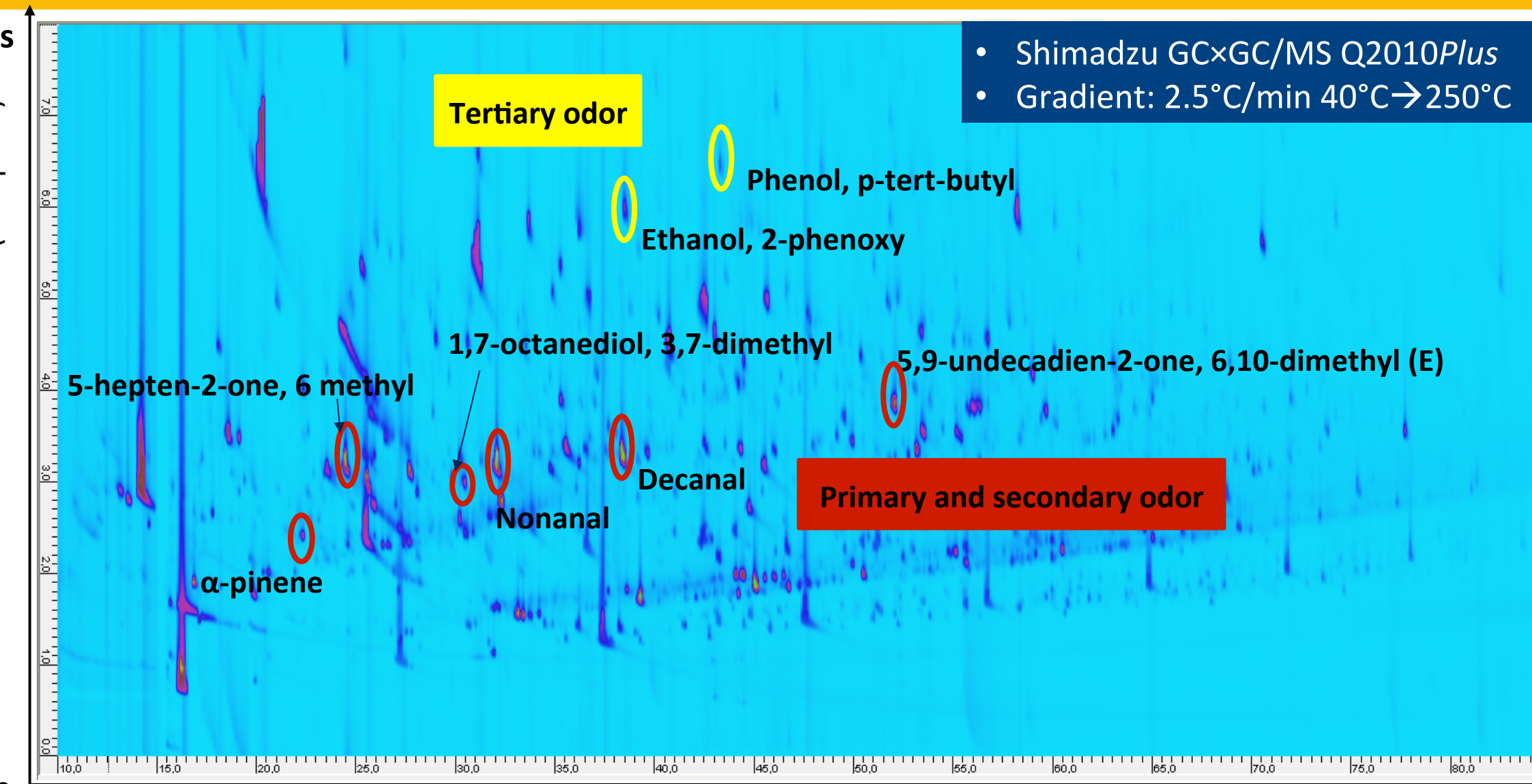
Sample temperature= 190°

Purge flow = 20 mL/min

Purge time= 20 min

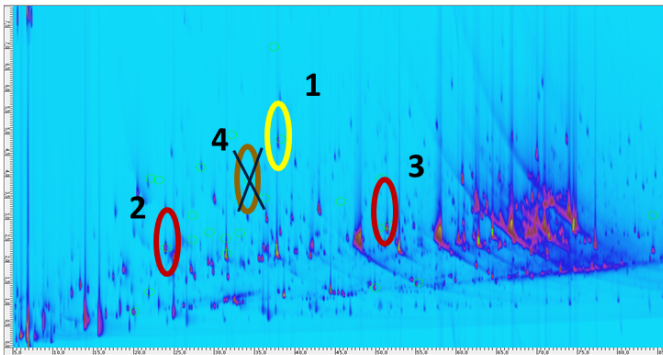
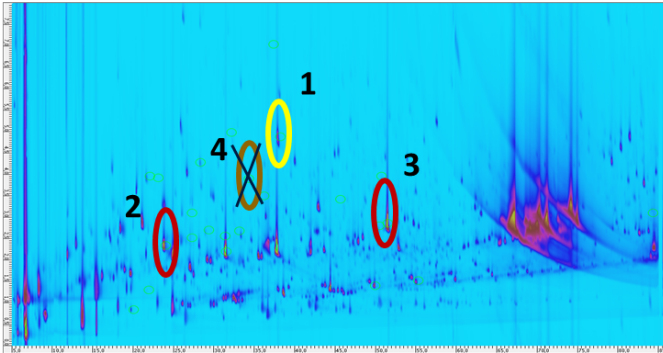
Split = 0 mL/min

CHROMATOGRAM OF A REAL SAMPLE

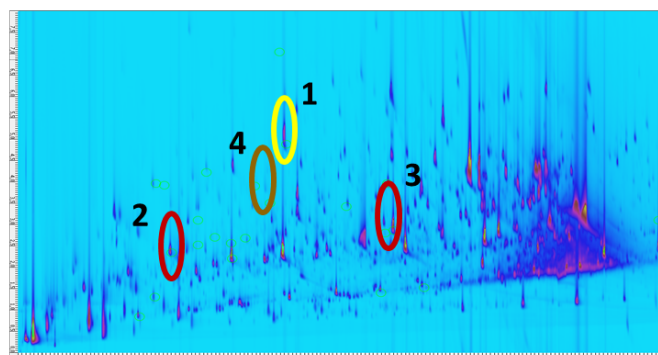
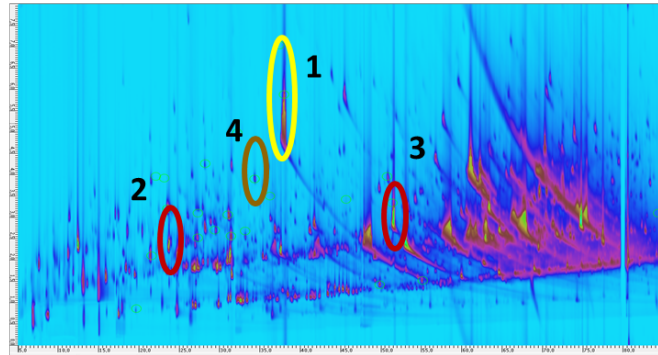


COMPARISON OF REAL SAMPLES?

Subject n°1 – Male – 25 y.o.



Subject n°2 – Female – 25 y.o.



- Complex samples
- **Comparison is not trivial**
- A lot of data to process

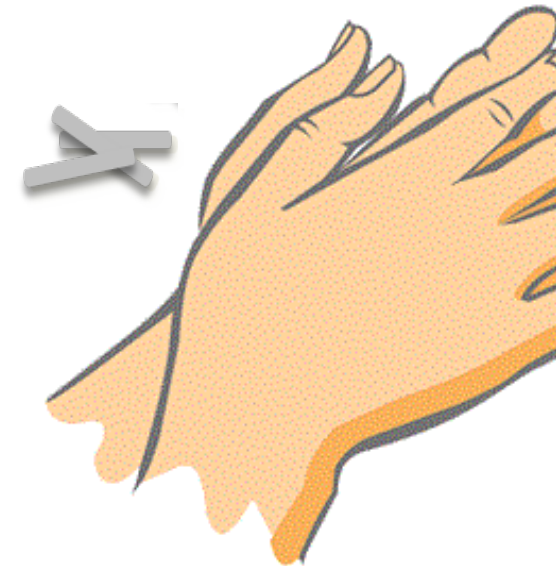
Need for an automated data processing to extract relevant information
Need for a panel of persons to evaluate the strategy

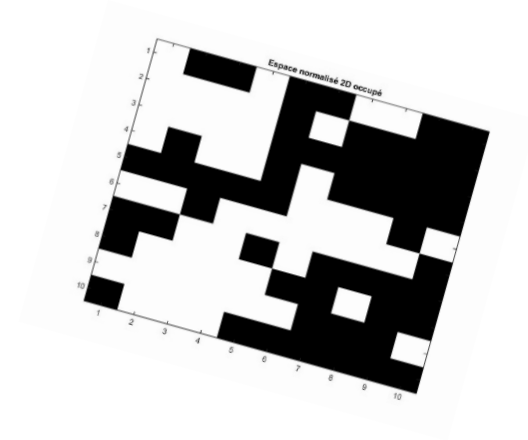
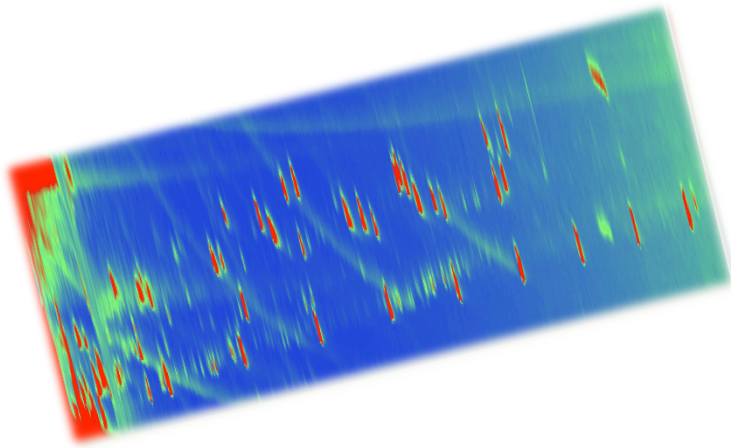
CHROMATOGRAMS OF REAL SAMPLES: PANEL

Panel of 119 persons

	gender		age (years)			phototype		
total	♂	♀	10-23	24-36	37-81	1	2	3
119	61	58	39	39	41	25	79	15

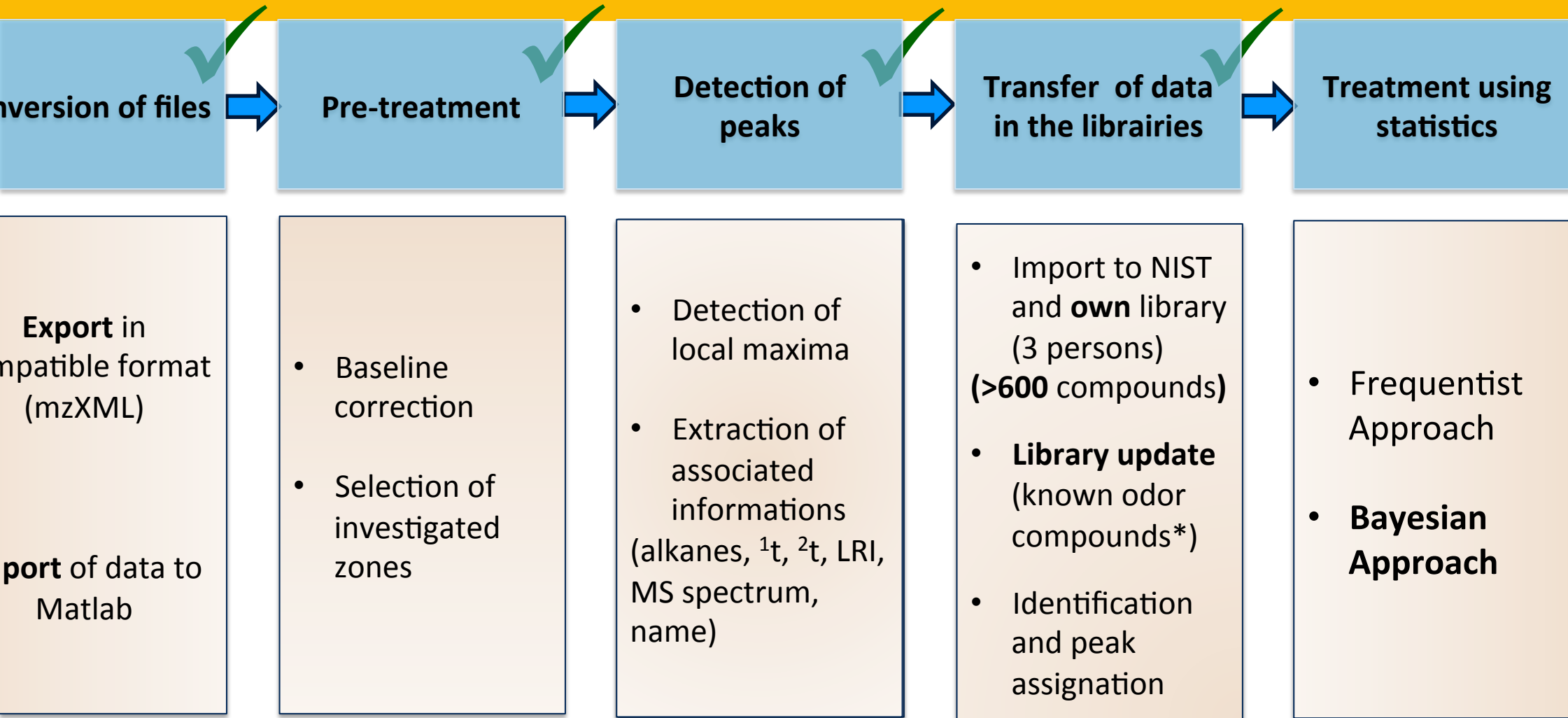
- Phototype 1 - skin is sun sensitive and does not burn
- Phototype 2 - intermediate skin
- Phototype 3 – well tanning skin
- 4 direct samplings of hands/person (Sorb-star®)
 - 15 minutes
 - Blank (sampling room)
- TD* - GC×GC-MS**
- **3 chromatograms/person**





DATA PROCESSING / BAYESIAN APPROACH

CHROMATOGRAPHIC DATA PROCESSING WITH MATLAB



Chromatogram ↔ 1 vector corresponding to 600 compounds peak intensities

* Cuzuel et al., *Origin, analytical characterization and use of human odor in forensics*, 2017, *J. Forensic Sci.*

BAYESIAN APPROACH (*A POSTERIORI*)

: the two chromatograms are obtained from the same person

: the two chromatograms are NOT obtained from the same person

D represent the observed data (the two chromatograms), Bayes formula gives :

$$P(H_0 | D) = \frac{f(D | H_0)P(H_0)}{f(D | H_0)P(H_0) + f(D | H_1)P(H_1)}$$

protocol:

Definition of a **distance d between 2 chromatograms** ($D \equiv d$)

Panel of chromatograms of individuals (119 persons sampled 4 times) splitted in
dependent calibration and test groups

Calibration group → estimation of **distributions of d** for couples of chromatograms from
the same person $f(d | H_0)$ and from different persons $f(d | H_1)$

Test group → estimation of **performance** (AUC, sensitivity, spécificity)

BAYESIAN APPROACH: CHOICE OF DISTANCE BETWEEN CHROM

Estimation of the statistical likelihood

Options :

d : distances between 600-vectors of intensities :

- a) **euclidian distance**
- b) **1 – Pearson correlation coefficient**
- c) **1 – Spearman correlation coefficient**

• intensities **normalized** / **binarized** (b=c)

Calibration group (260 chromatograms / 75 persons)

341 couples of chromatograms for H_0 (same person)

33 329 couples de chromatograms for H_1 (différent persons)

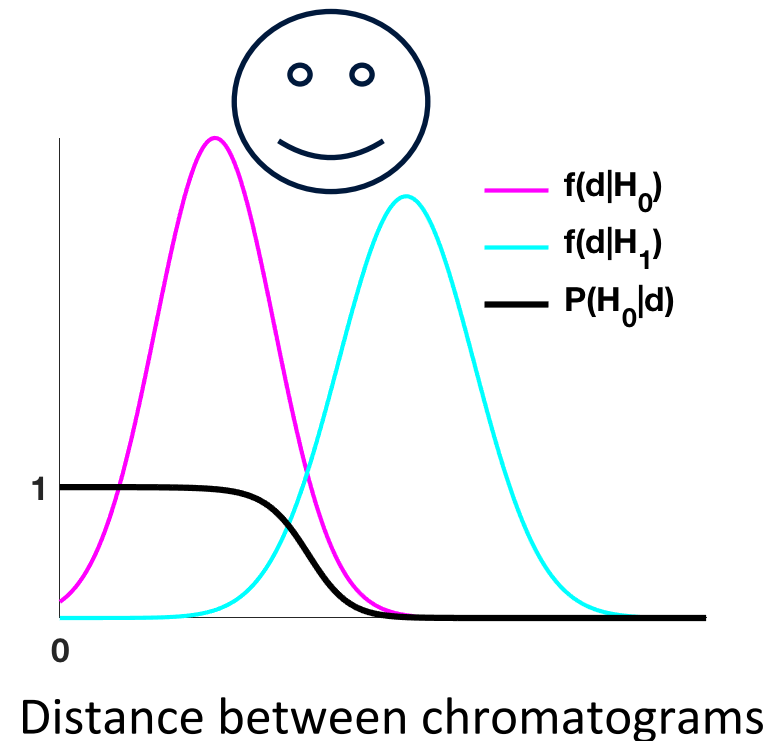
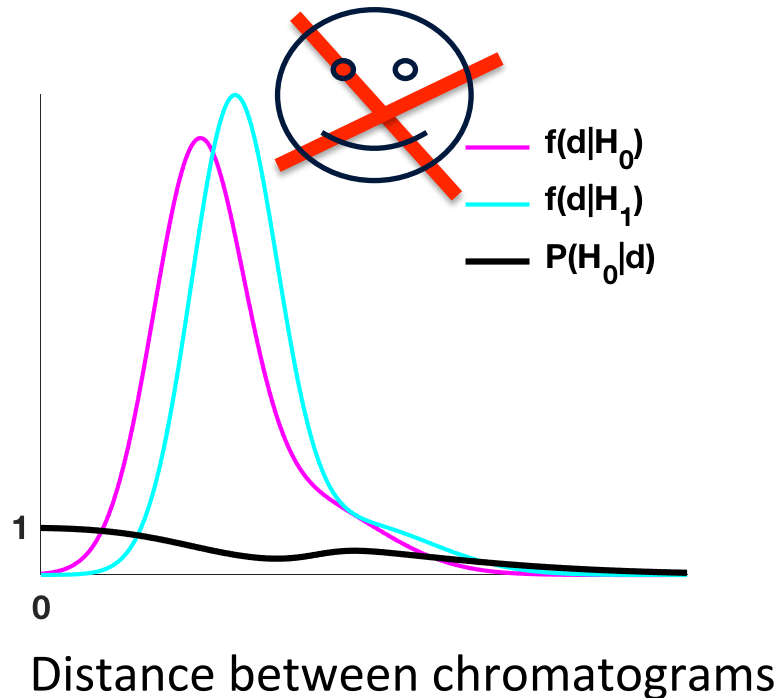
➔ **histograms** of d values for H_0 and H_1

Adjustment of histograms using several **gaussian curves**

➔ $f(d | H_0)$ and $f(d | H_1)$

BAYESIAN APPROACH: EXPECTED RESULTS

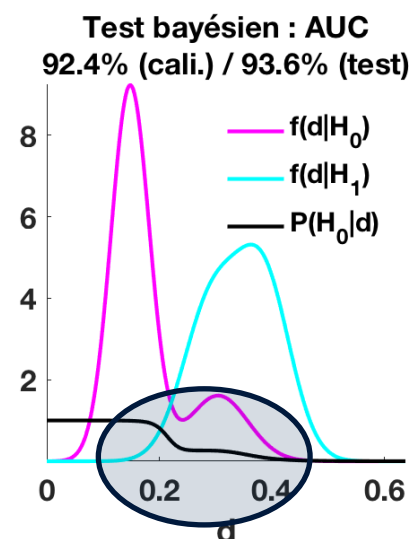
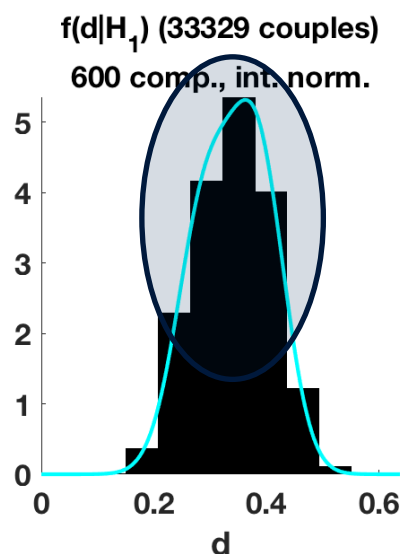
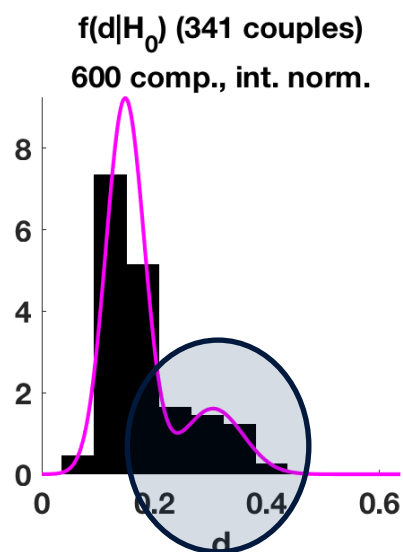
- Probabilities *a priori* : $P(H_0) = P(H_1) = 0.5$ $P(H_0 | d) = \frac{f(d | H_0)P(H_0)}{f(d | H_0)P(H_0) + f(d | H_1)P(H_1)}$
- Fictitious examples of statistical likelihood



BAYESIAN APPROACH: RESULTS USING 600 COMPOUNDS

(%AUC calibration / %AUC test)

distance \ intensities	euclidian	$1 - \rho_{\text{Pearson}}$	$1 - \rho_{\text{Spearman}}$
normalized	62.7% / 64.6%	74.6% / 74.7%	92.4% / 93.6%
binarized	88.4% / 91.6%	89.6% / 91.7%	



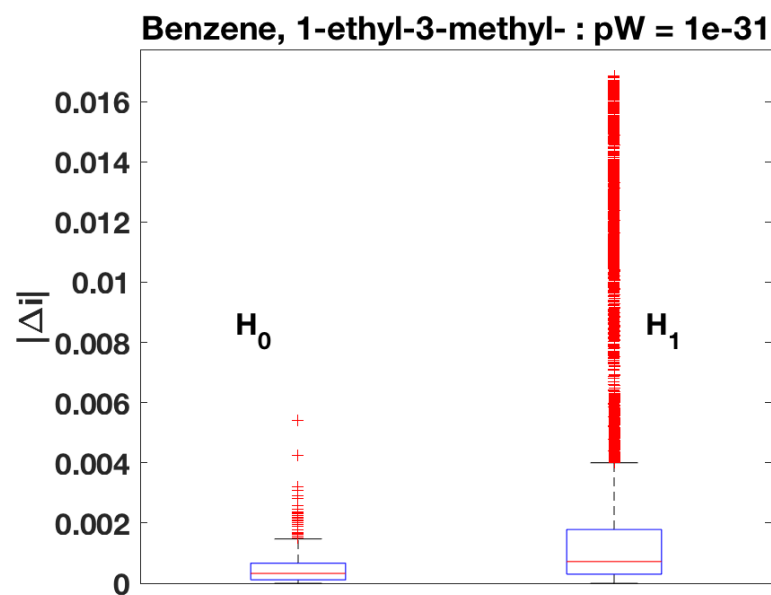
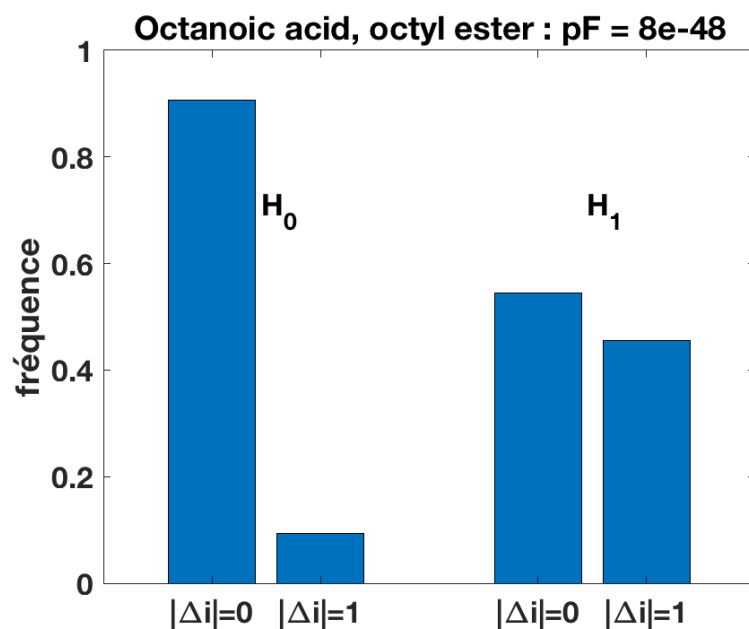
2 modes!

N.B. using the test group, there are 173 / 9 418 couples for H_0 / H_1 respectively

GCxGC-ISCC Riva del garda 2018

BAYESIAN APPROACH: DISCRIMINATING COMPOUNDS

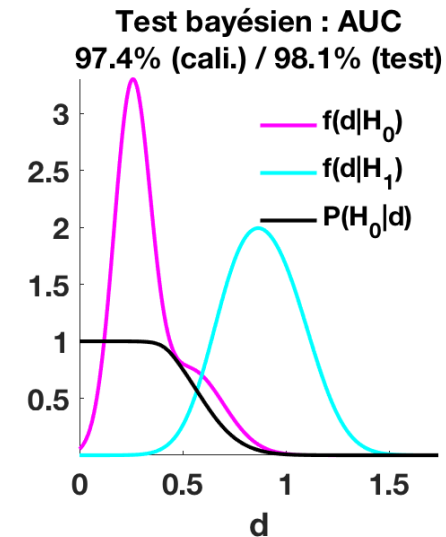
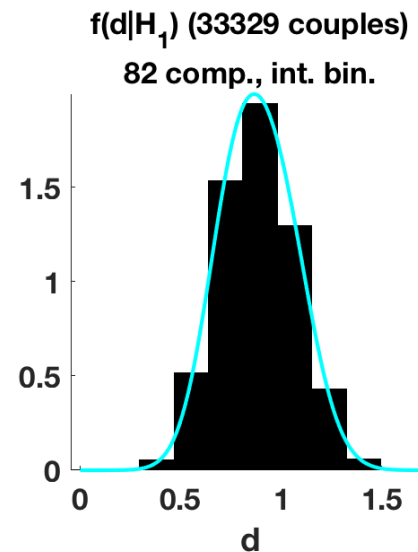
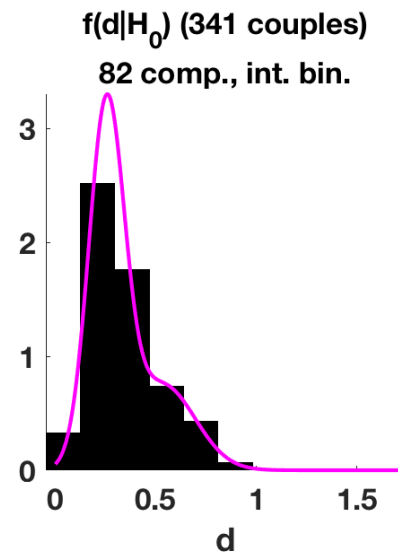
- Discriminating compounds for H_0 and H_1 : those which **intensity differences** $|\Delta i|$ are **significantly lower for H_0 than H_1**
- Quantification : **p-value** using **unilateral Fisher test** (binarized intensities) or **Wilcoxon** (normalized intensities) on $|\Delta i|$
- Examples :



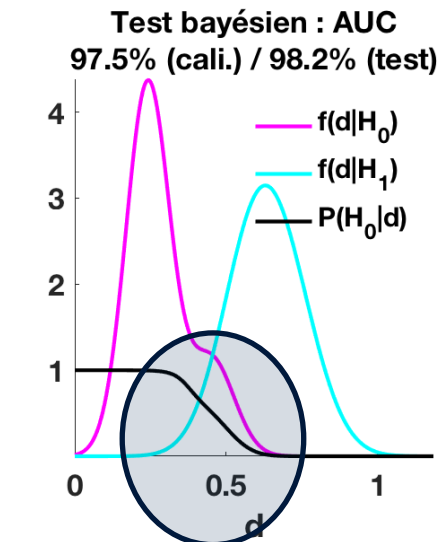
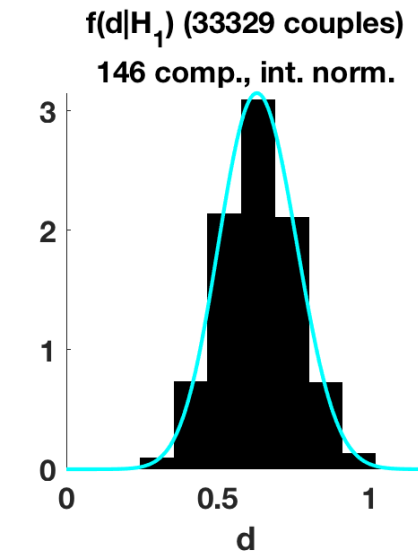
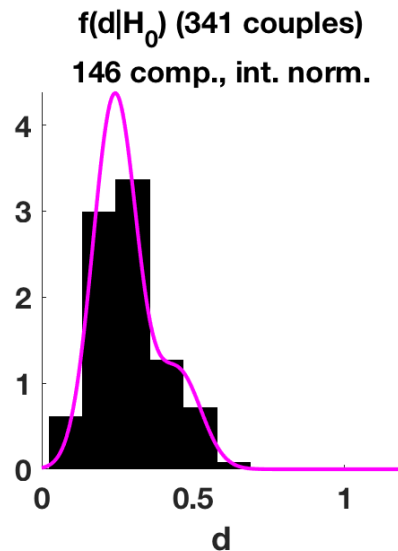
BAYESIAN APPROACH: RESULTS USING DISCRIMINATING

(%AUC calibration / %AUC test)

Binarized



Normalized



BAYESIAN APPROACH: RESULTS USING DISCRIMINATING COMPOUNDS

(%AUC calibration / %AUC test)

- **Threshold π value $-\log_{10}(p)$** of Fisher test (binarized intensities) or Wilcoxon (normalized intensities) : optimized value obtained using cross **validation** (K=3) on **calibration group**

distance intensities	euclidian	$1 - \rho_{\text{Pearson}}$	$1 - \rho_{\text{Spearman}}$
normalized	$\pi = 12 / 61$ comp. 76.2% / 73.9%	$\pi = 13 / 54$ comp. 78.1% / 75.2%	$\pi = 7 / 146$ comp. 97.5% / 98.2%
binarized	$\pi = 18 / 82$ comp. 93.1% / 94.8%	$\pi = 18 / 82$ comp. 97.4% / 98.1%	

Discussion

(%AUC calibration / %AUC test)

performances

intensities	AUC	sensitivity	specificity	nb. compounds
binarized	97.4% / 98.1 %	89.4% / 90.0%	94.9% / 92.5%	82
normalized	97.5% / 98.2 %	89.1% / 85.9%	93.7% / 95.0%	146

Adequate distance → **quantitative** exploitation of compounds intensities despite the analytical variability

Selection → second modes of $f(d|H_0)$ et $f(d|H_1)$ are strongly decreased → better results

Binarized : more **parsimonious (82/146 compounds to be used)**

57 common compounds for both classifiers

nota bene

same direct samples

no pollution by other odors

CONCLUSION AND PERSPECTIVES

Direct/non direct sampling procedures for human (hand) odor analysis

Comprehensive GC×GC-MS method and data (ToF)

Validation of procedures **in the field** with dog handlers

Large Panel of individuals to test the model

Storage of samples: standardized procedure

Data processing in progress for real application

✓ **Different samples** (direct or not...) and sampling conditions

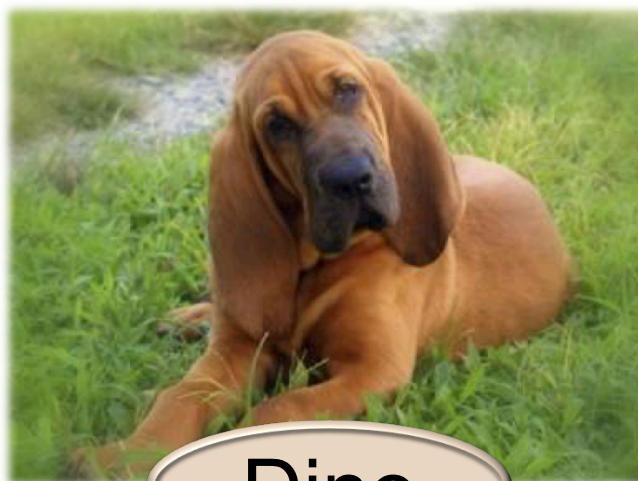
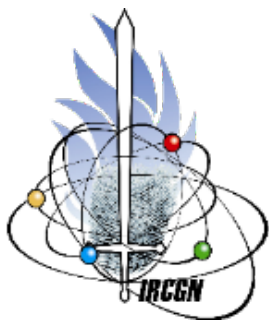
✓ Study of discriminating compounds

✓ Normalization on discriminating compounds, more complex distance...

The final answer to the question must be YES or NO not 98.2%



ACKNOWLEDGEMENTS



Dino



THANK YOU FOR YOUR ATTENTION!