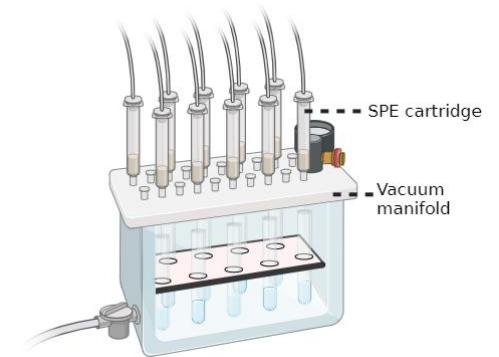
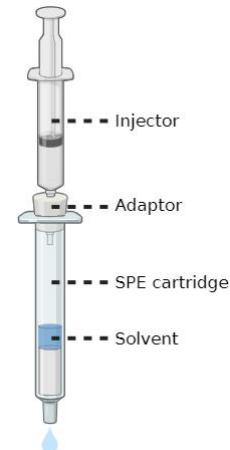
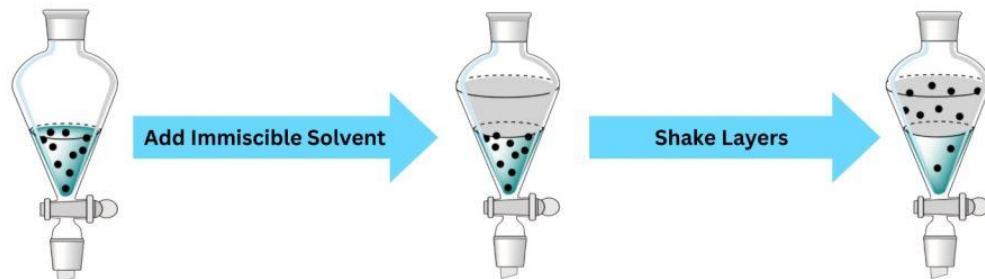


# **MODES D'INTRODUCTION DE L'ÉCHANTILLON**

- Analyse directe
- Couplage GC-MS
- Couplage LC-MS
- Couplage CE-MS
- Couplage SFC-MS

# Extraire (LLE, SLE, SPE)



Resolubiliser si nécessaire



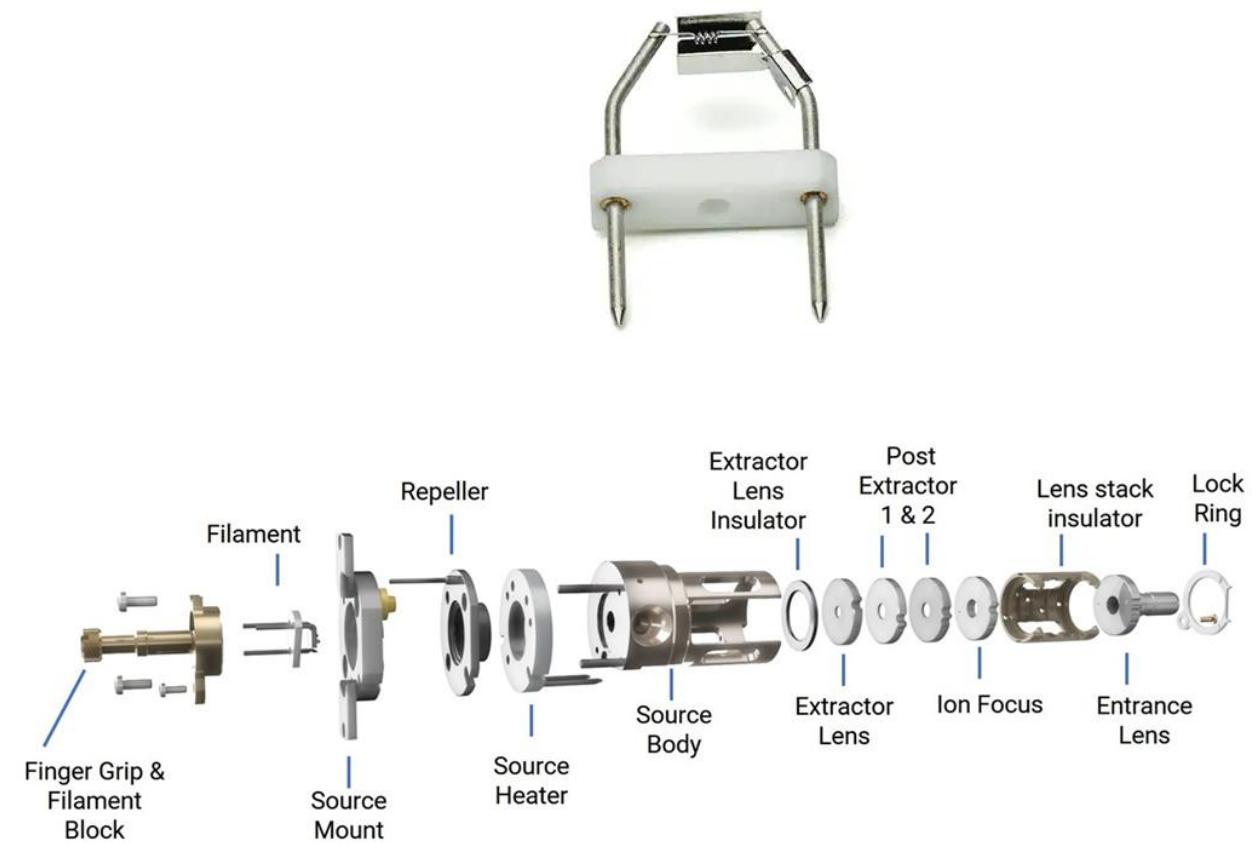
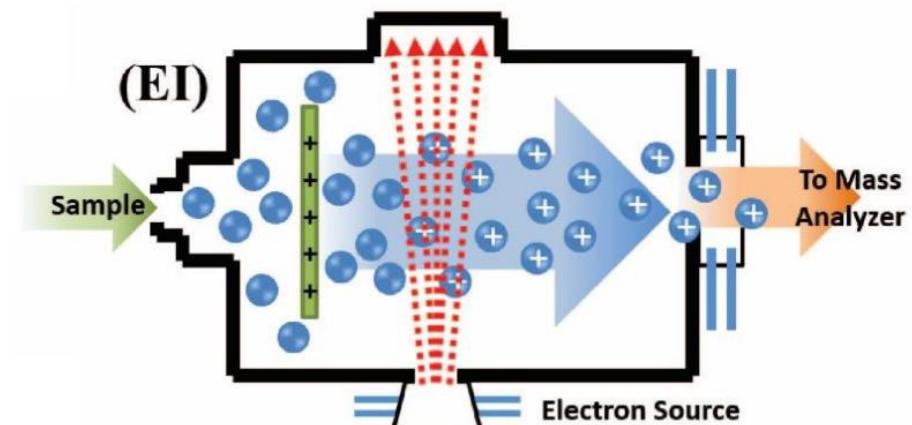
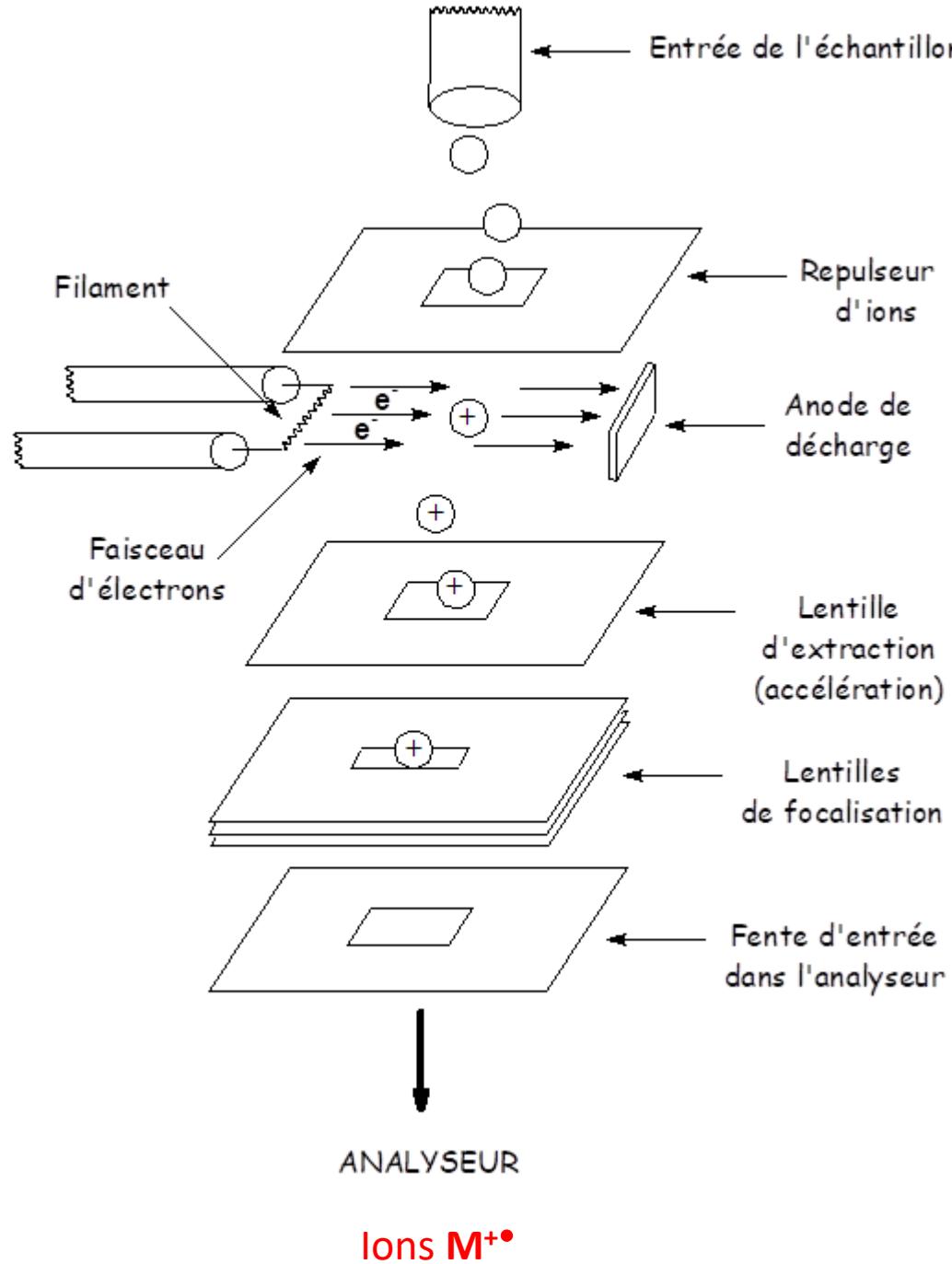
Dériver si nécessaire

# COUPLAGES GC-MS

## ANALYSE DES PETITES MOLÉCULES VOLATILES



# **Impact électronique (EI)**

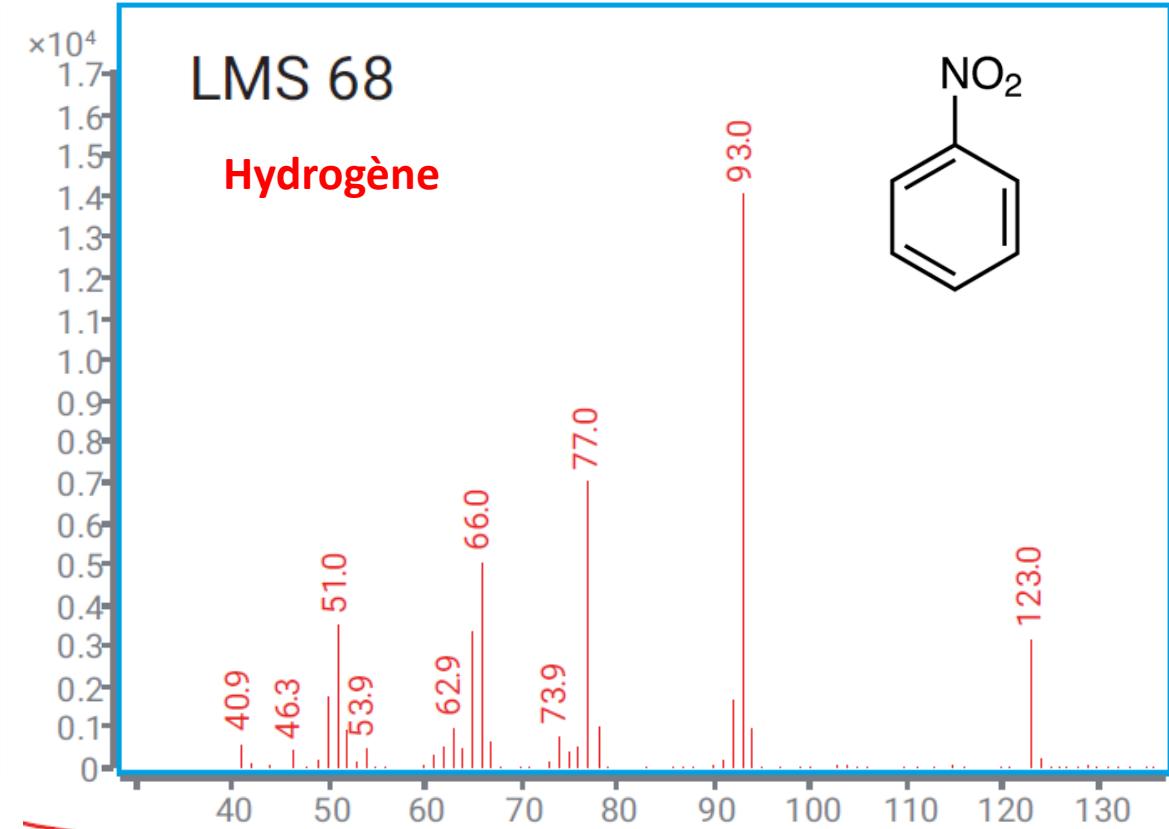
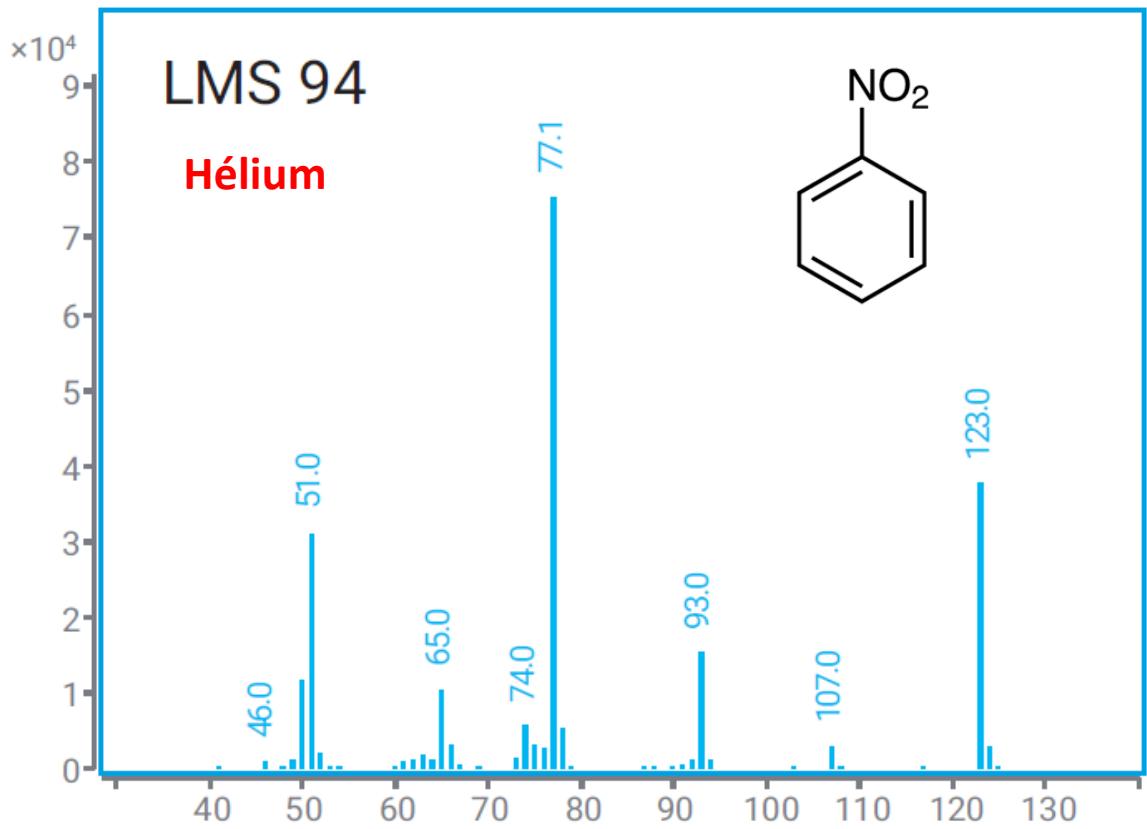


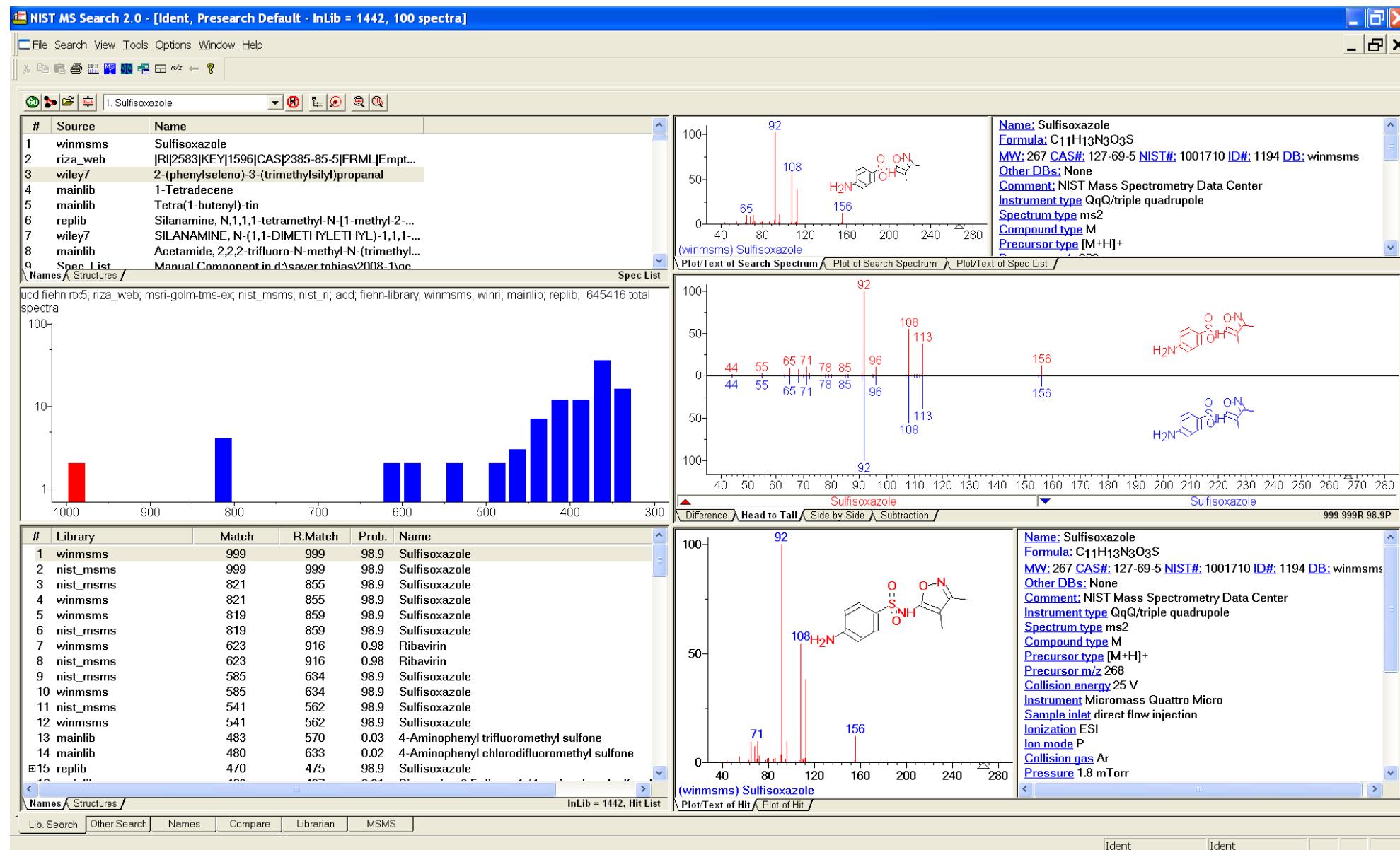
## **Principaux ions:**

Type d'analyte	Ionisation positive Principal // occasionnel	Ionisation négative
apolaire	$M^{+\bullet}$ // $M^{2+}$	-
polarité moyenne	$M^{+\bullet}$ // $[M+H]^+$ , $M^{2+}$	-
polaire	$M^{+\bullet}$ // $[M+H]^+$ , $M^{2+}$	-

## Intérêts et limitations:

- le composé doit être volatil ( $T_{\text{eb}} < 500^\circ\text{C}$ ; limite de masse  $\approx 10^3$ )
- ion moléculaire  $M^{+\bullet}$  (mais pas toujours) => **masse?**
- ions fragments => **informations structurales**
- source sous vide => **contraintes sur GC (dc, débits, gaz vecteur...)**





- NIST NIST/EPA/NIH Mass Spectral Library 2023: 394.000 spectres EI / 341,000 composés => **2500€**
  - WILEY Registry of Mass Spectral Data 12th Edition 860.000 spectres EI / 700.000 composés => **8500€**

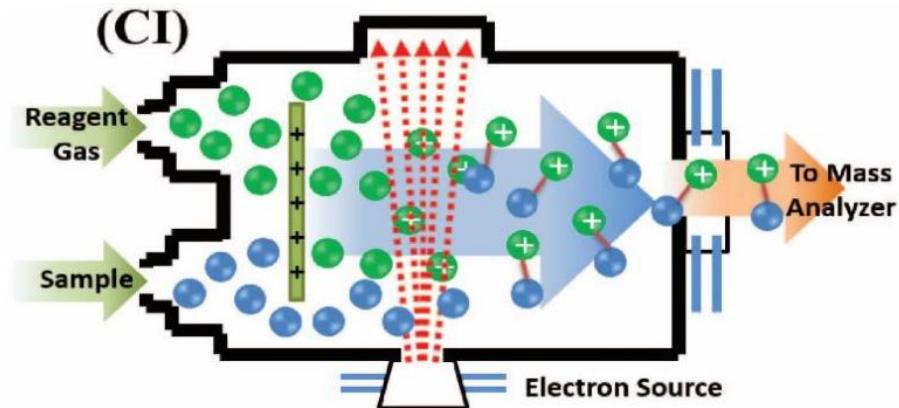
# **Ionisation Chimique (CI)**

## Principe:

Qté de Gaz Réactif >>> Qté de Produit Étudié  
 $10^3 \rightarrow 10^4$

ionisation préférentielle du gaz réactif (méthane, isobutane ou ammoniac)

Collision avec d'autres molécules du même gaz => plasma d'ionisation



Gaz réactif	AP (kcal.mol <sup>-1</sup> )	Ions réactifs	m/z	Abondance (%)
Méthane	127	$\text{CH}_5^+$ $\text{C}_2\text{H}_5^+$ $\text{C}_3\text{H}_5^+$	M+17 M+29 M+41	47 41 6
Isobutane	195	$\text{C}_4\text{H}_9^+$ $\text{C}_3\text{H}_7^+$ $\text{C}_3\text{H}_3^+$	M+57 M+43 M+39	93 4 3
Ammoniac	207	$\text{NH}_4^+$ $\text{NH}_3 + \text{NH}_4^+$	M+18 M+35	97 3

molécules organiques AP  $\approx 200$  Kcal.mol<sup>-1</sup>

=> ammoniac + sélectif que isobutane et méthane

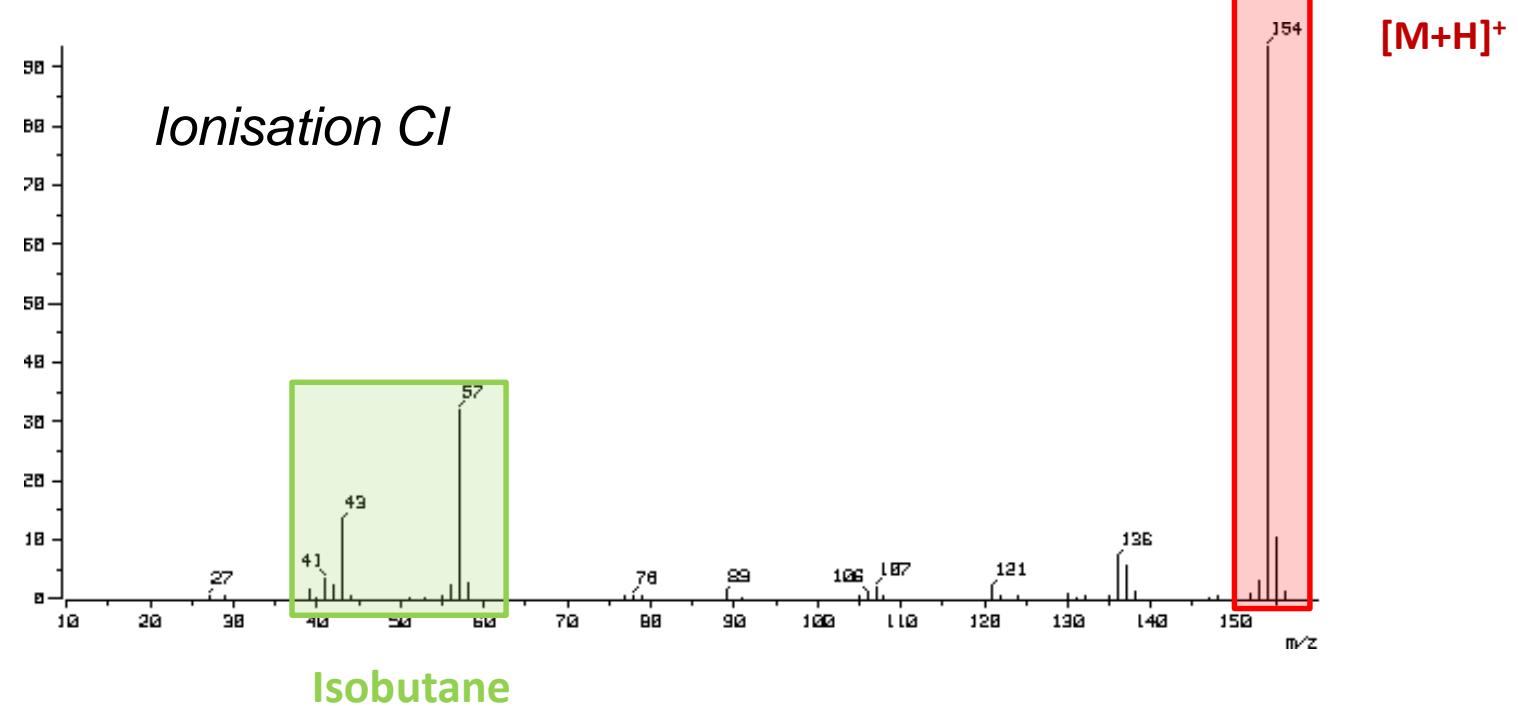
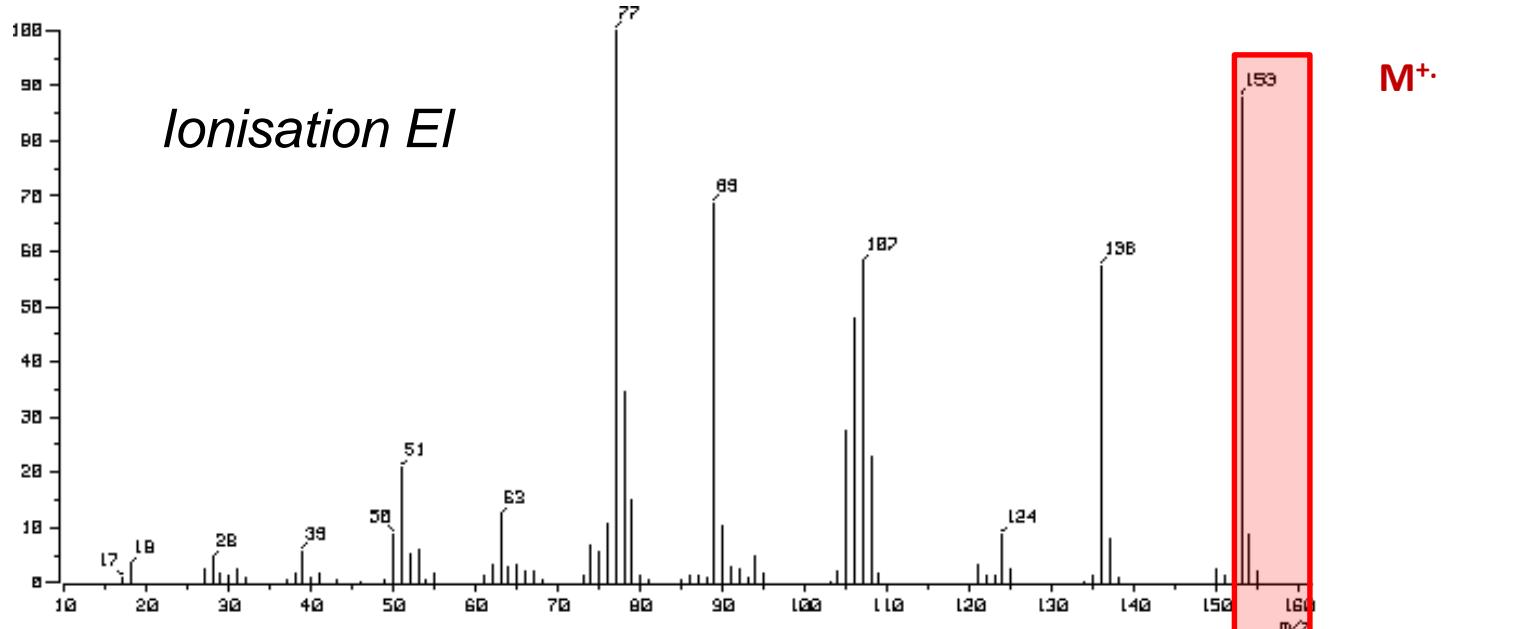
- énergétique que EI => favorise l'obtention de l'ion moléculaire  
(peu de fragmentation car énergie interne + faible que pour EI).

## Principaux ions:

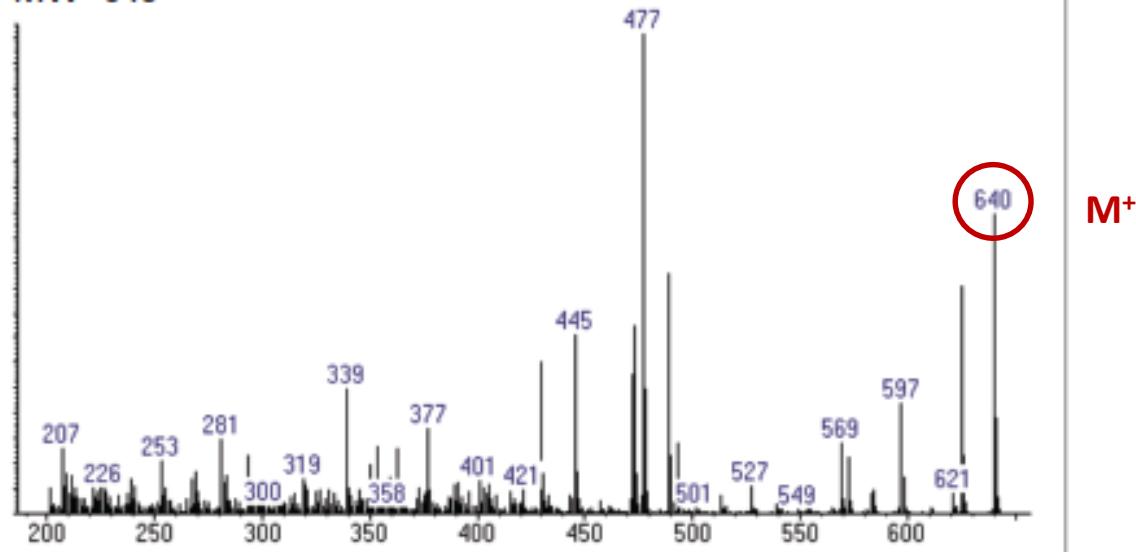
Type d'analyte	Ionisation positive principal // occasionnel	Ionisation négative principal // occasionnel
apolaire	$[M+H]^+$	$M^{\cdot-} // [M-H]^-$
polarité moyenne	$[M+H]^+ // [M+X]^+, [2M+H]^+$	$M^{\cdot-}, [M-H]^-$
polaire	$[M+H]^+, [M+X]^+ // [2M+H]^+$	$[M-H]^- // [M+R]^-$ , $M^-$

$X \Rightarrow$  selon les gaz (ex:  $[M+NH_4]^+$ ):       $M^{\cdot-}$  : capture d'électron  $M+e^-$

## Spectres de masse du 3-nitrobenzyl alcool

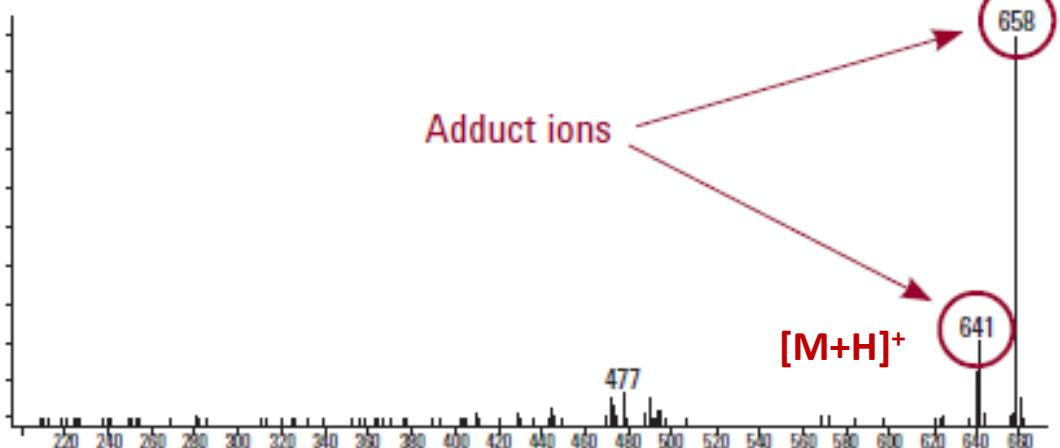


THCA-PFPA-HFIP **EI** Spectrum  
MW=640

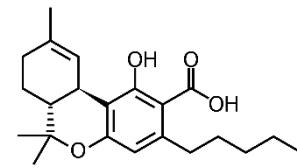


**M<sup>+</sup>.**

THCA-PFPA-HFIP **PCI NH<sub>3</sub>** Spectrum  
MW=640



**Comparison of  $\Delta 1$ -Tetrahydrocannabinolic acid (THCA).** The EI spectra show many fragments, but PCI and NCI show fewer. CI spectra enable easier identification of compounds by GC/MSD.



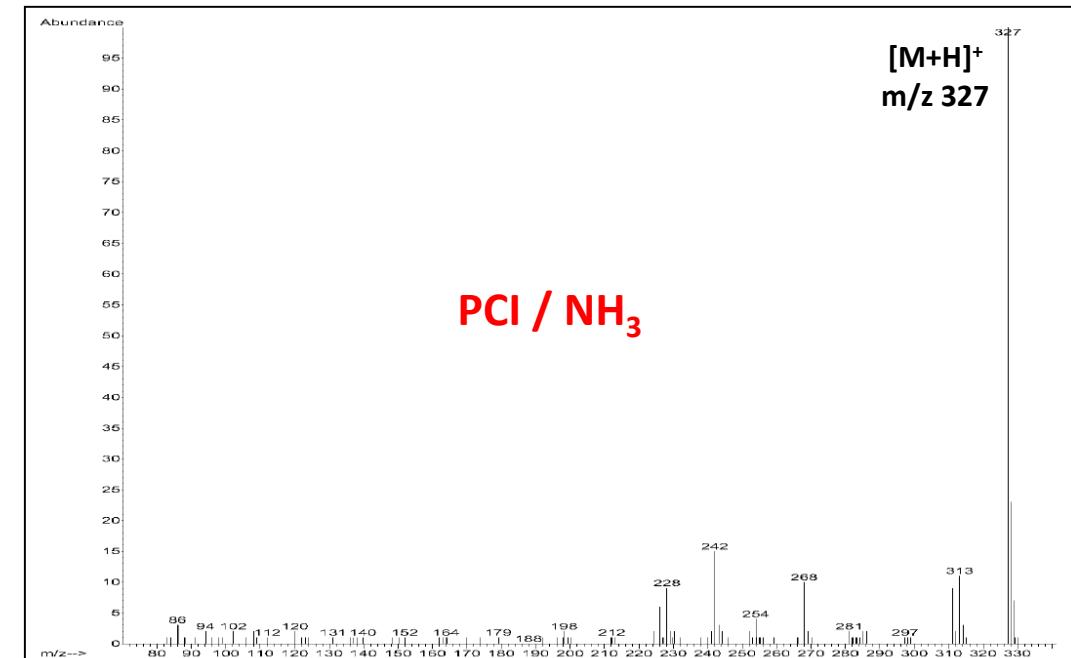
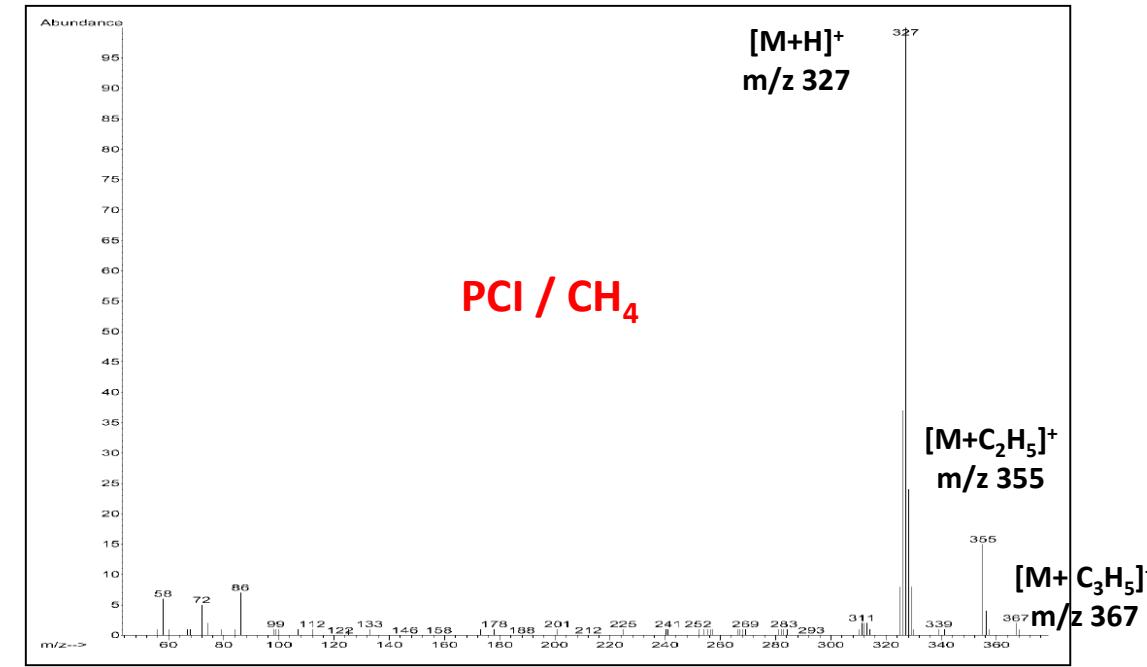
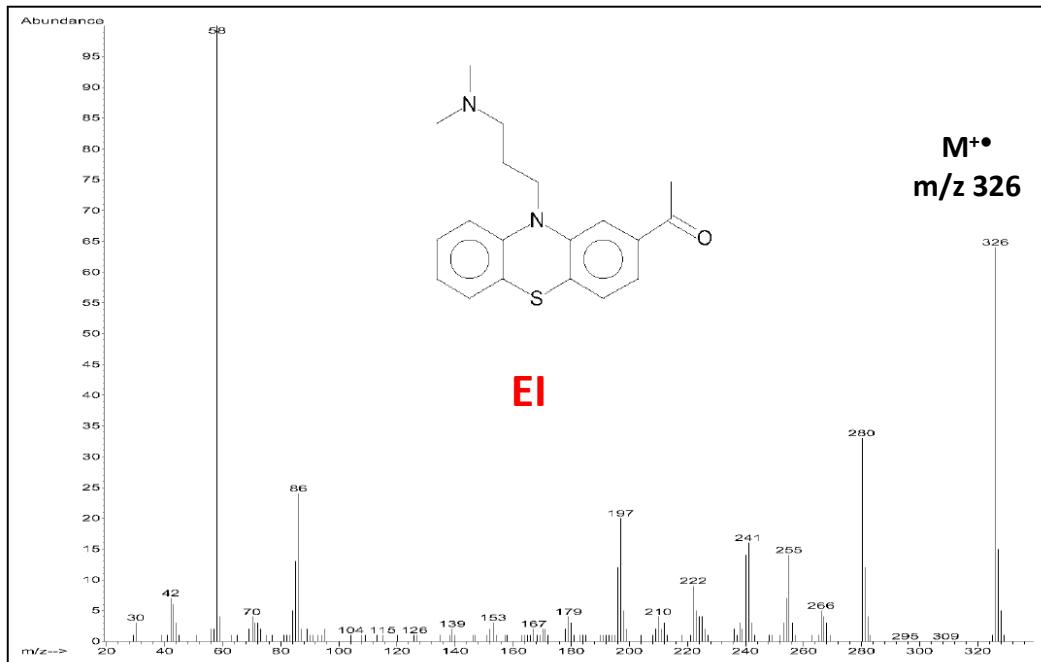
### **Intérêts et limitations:**

- on peut observer l'ion pseudo-moléculaire => **masse OK**
- c'est une technique **douce** : on obtient des ions fragments mais moins que pour EI  
=> **peu d'informations structurales**
- le composé doit être volatil ( $T_{eb} < 500^\circ\text{C}$ ; limite de masse  $10^3$ )
- source sous vide => **contraintes sur GC (dc, débits,...)**

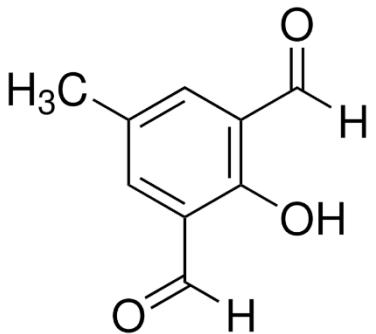
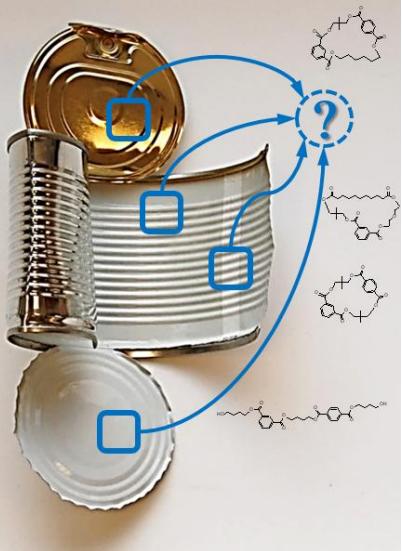
# Acepromazine

CAS-Nr. 61-00-7

Molecular formula:  $C_{19}H_{22}N_2OS$

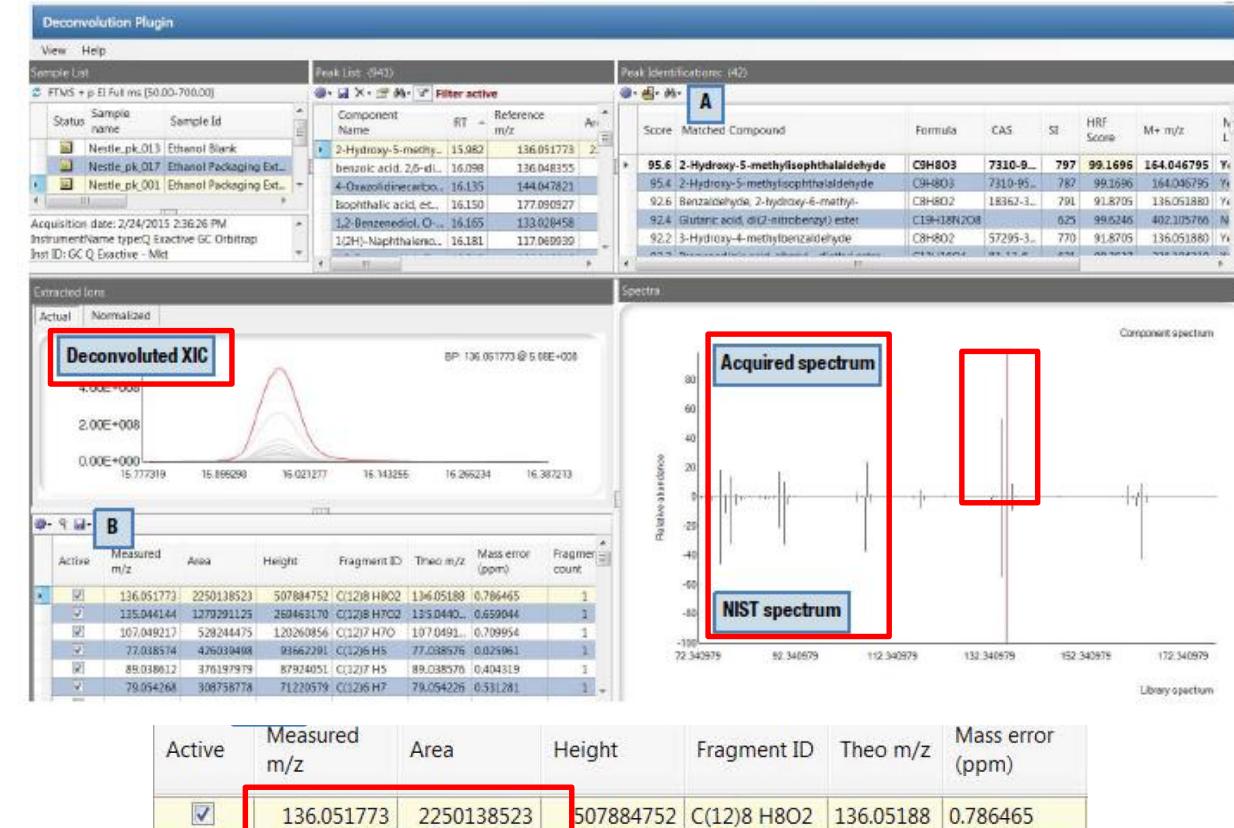
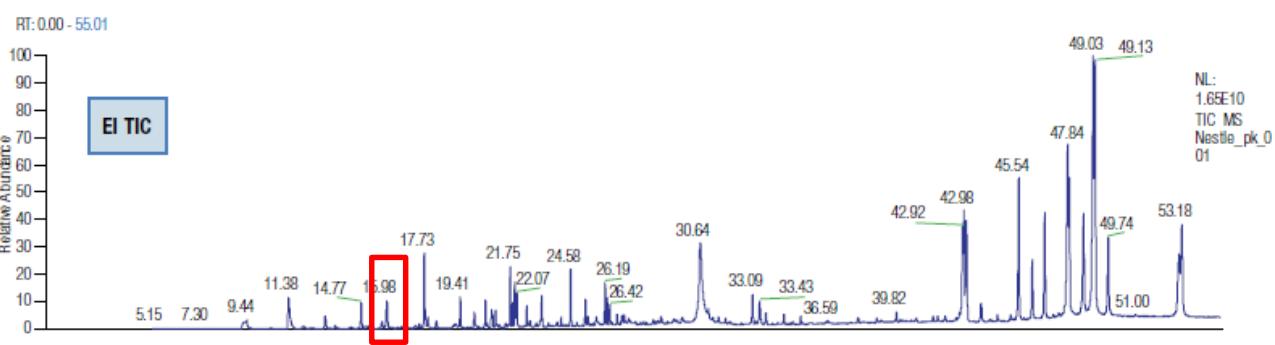


# **Applications GC-MS**



Monoisotopic mass: 164.047344

2-Hydroxy-5-methyl-1,3-benzeneddicarboxaldehyde



# GC-HRMS

Characterizing unknowns in food packaging using GC Orbitrap Mass Spectrometry  
 Dominic Roberts, Jesus Varela, Yves-Alexis Hammel and Paul Silcock  
 Thermo AN 10527 (2016)

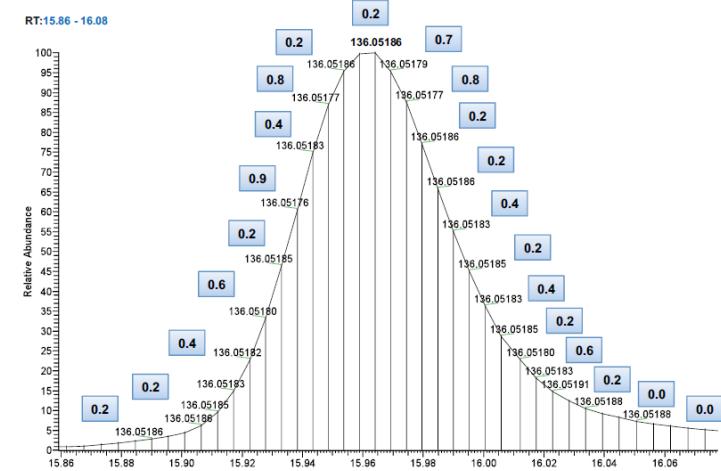
## TRACE 1310 GC System Parameters

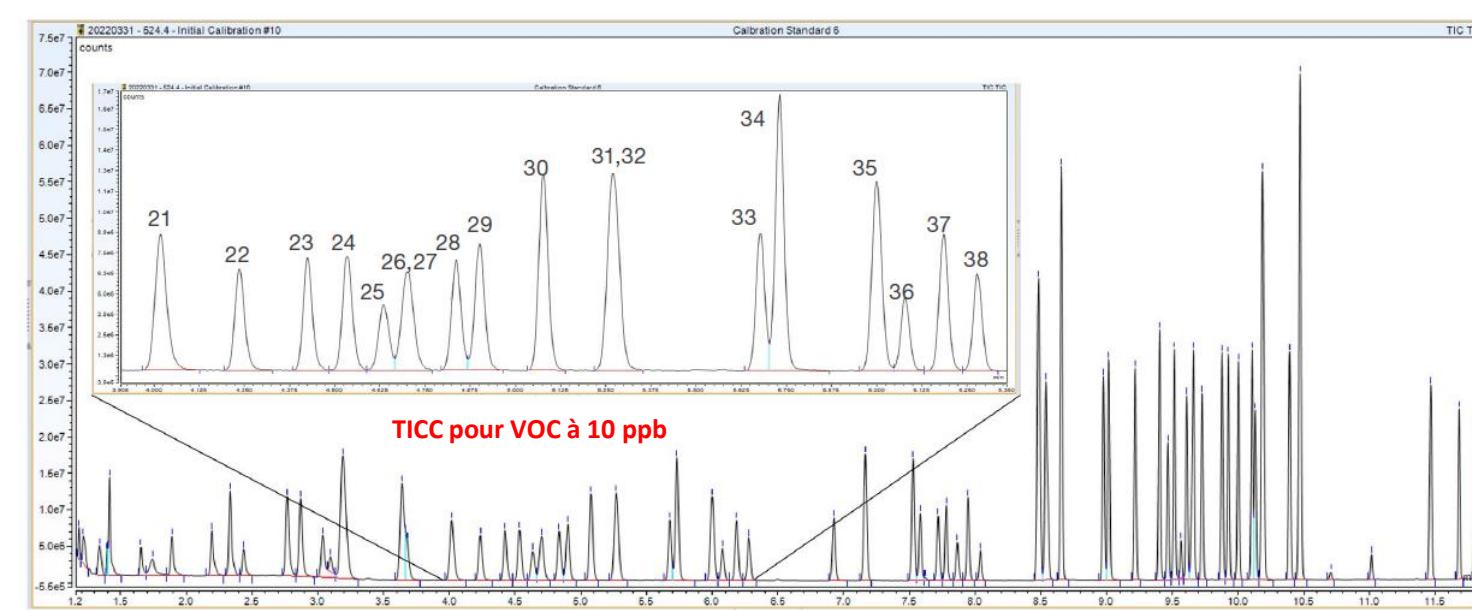
Injection Volume (μL)	1
Liner	Single gooseneck P/N 453A0344-UI
Inlet (°C)	SSL 280
Carrier Gas, (mL/min)	He, 1.3
<b>Oven Temperature Program</b>	
Temperature 1 (°C)	40
Hold Time (min)	0.5
Temperature 2 (°C)	325
Rate (°C/min)	5.5
Hold Time (min)	12

Table 2. Mass spectrometer conditions.

## Q Exactive GC Mass Spectrometer Parameters

Transfer line (°C)	280
Ionization type	EI/PCI
Ion source (°C)	230 EI / 190 CI
Electron energy (eV)	70
Acquisition mode	Full-scan
Mass range (Da)	50–700
Resolving power (FWHM at <i>m/z</i> 200)	120,000
Lockmass, column bleed ( <i>m/z</i> )	207.03235



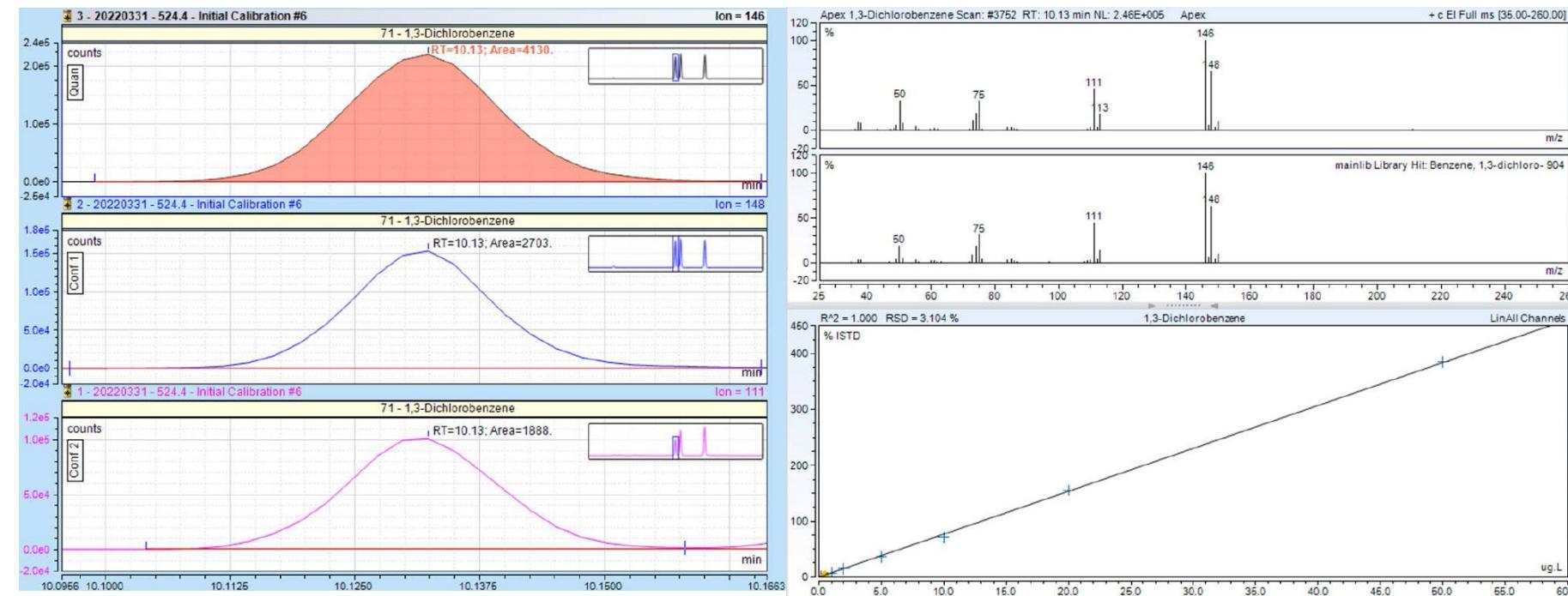


21. t-butyl ethyl ether  
22. *cis*-1,2-dichloroethene  
23. Bromochloromethane  
24. Chloroform  
25. Carbon tetrachloride  
26. Tetrahydrofuran

27. 1,1,1-trichloroethane  
28. 1,1-dichloropropene  
29. 1-chlorobutane  
30. Benzene  
31. *t*-amyl methyl ether  
32. 1,2-dichloroethane

33. Trichloroethylene  
34. 1,4-dichlorobenzene (IS)  
35. *t*-amyl ethyl ether  
36. Dibromomethane  
37. 1,2-dichloropropane  
38. Bromodichloromethane

# GC-MS



**EIC pour 1,3-dichlorobenzene à 0.5 ppb**  
**Ion de quantification m/z 146**  
**Ions de confirmation m/z 148 et 111**

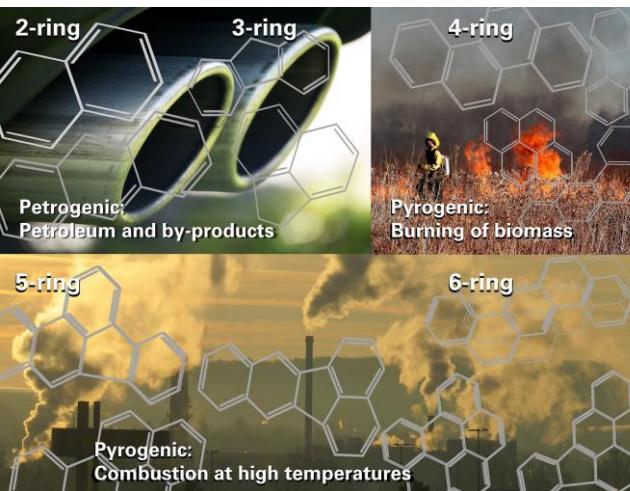
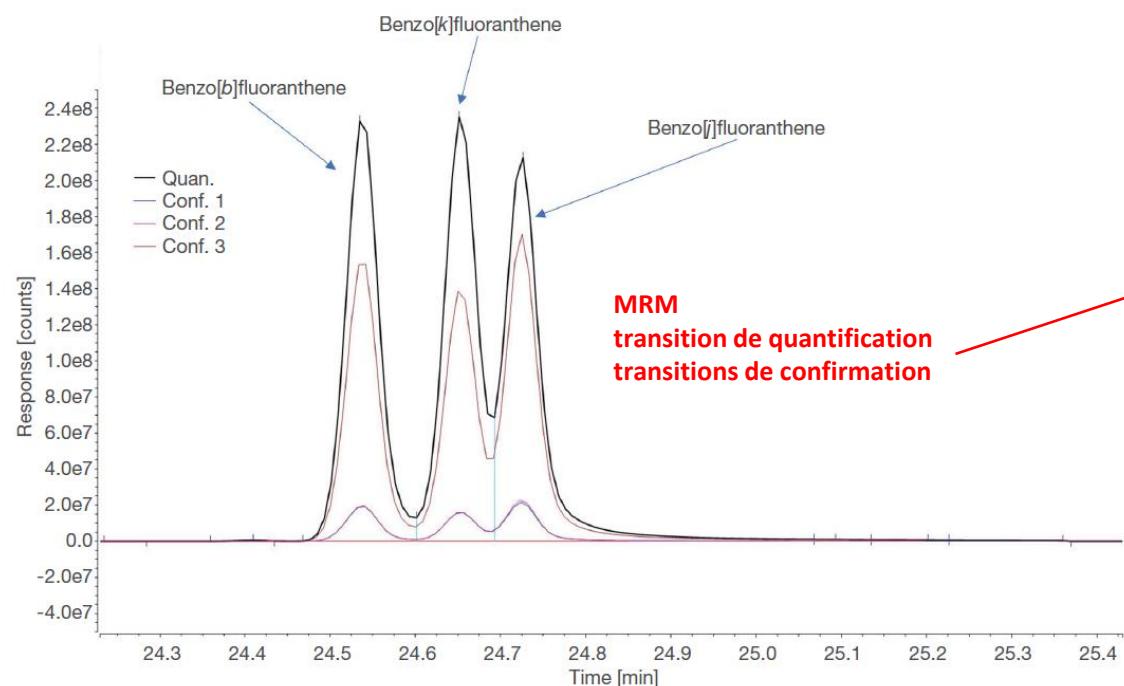
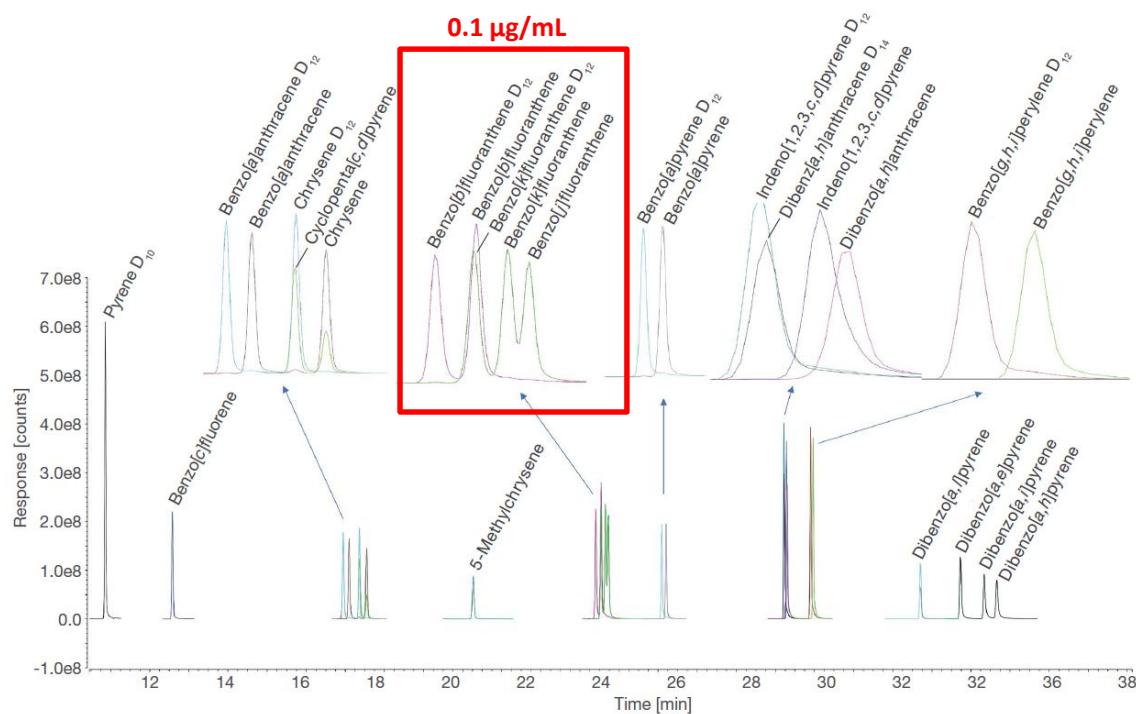
**gamme : 0.2 à 50 ppb**

Analysis of volatile organic compounds in drinking water according to  
U.S. EPA Method 524.4  
Thermo AN 001236 (2022)



# GC-MS/MS

sub ppb



Compound	R <sup>2</sup>	Range [µg/mL]	IDL [µg/mL]
Benzo[ <i>b</i> ]fluoranthene	0.9995	0.0005–1.0000	0.0001
Benzo[ <i>c</i> ]fluorene	0.9992	0.0005–1.0000	0.0002
Benzo[ <i>g,h,i</i> ]perylene	0.9996	0.0005–1.0000	0.0002
Benzo[ <i>jj</i> ]fluoranthene	0.9996	0.0005–1.0000	0.0001
Benzo[ <i>k</i> ]fluoranthene	0.9996	0.0005–1.0000	0.0001

Compound	Retention time (min)	Precursor ion (m/z)	Product ion (m/z)	Collision energy (V)
Benzo[ <i>b</i> ] fluoranthene	24.69	252	226	30
	24.69	252	250	30
	24.69	252	252	20
Benzo[ <i>k</i> ] fluoranthene	24.83	252	226	30
	24.83	252	226	20
	24.83	252	250	30
Benzo[ <i>j</i> ] fluoranthene	24.83	252	252	20
	24.87	252	226	30
	24.87	252	250	30
	24.87	252	252	20

Sensitive and reproducible analysis of 16 polycyclic aromatic hydrocarbons (PAHs) using gas chromatography – triple quadrupole mass spectrometry (GC-MS/MS). Thermo AN 10566 (2022)