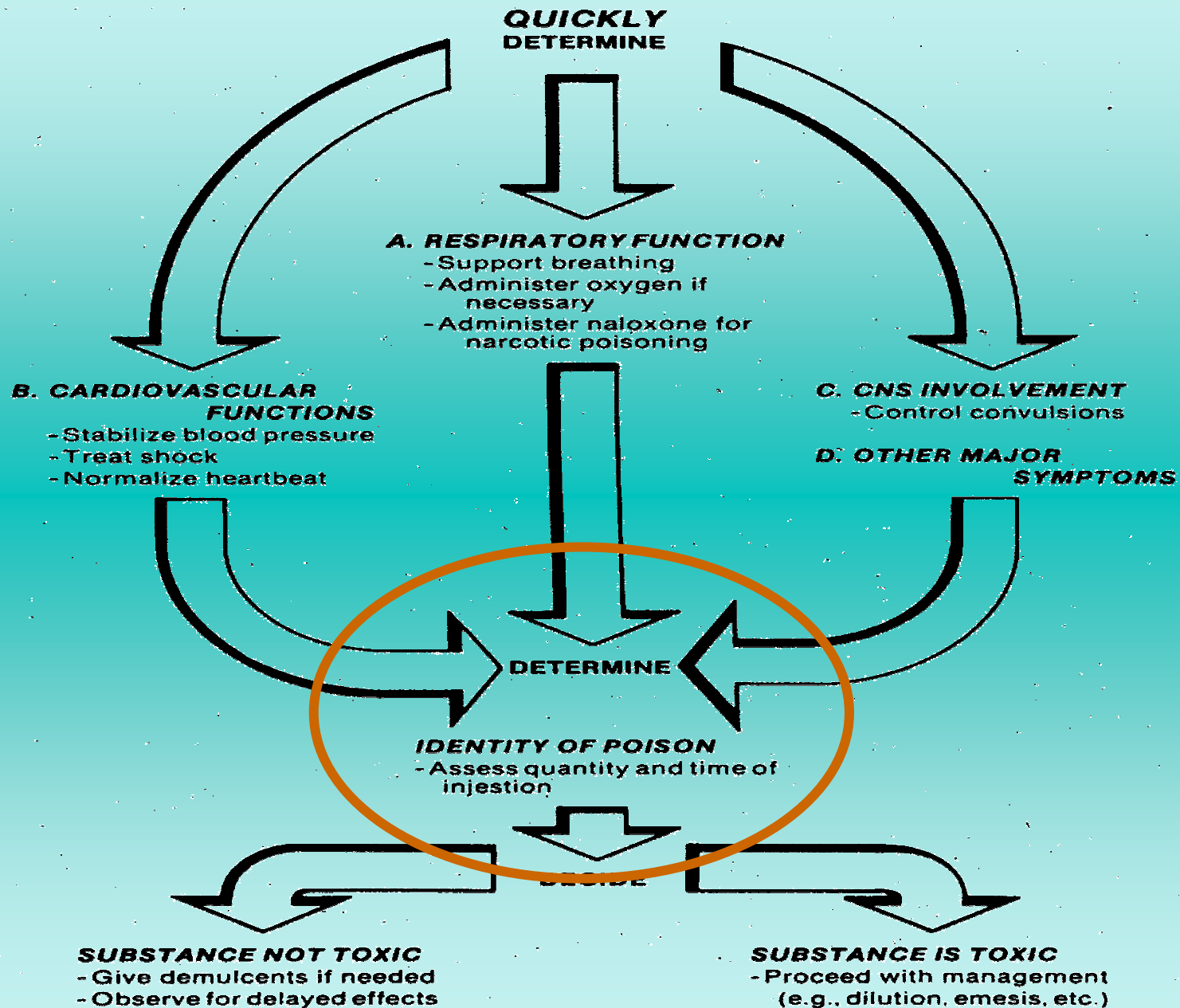


# *Toxikologische Analytik - Neue analytische Verfahren*

*Rainer Schmid*

*Sect. Biopharmaceutical & Toxicological Analysis  
Inst. for Medical & Chemical Labdiagnostics (KIMCL)  
General Hospital - Medical University of Vienna*

# ***PRINCIPLES IN MANAGEMENT OF THE POISONED PATIENT***



## ***Major causes of poisoning (in the United States)***

### **Plants**

**Soaps, detergents, cleaners**

**Household disinfectants.**

**deodorizers**

**Household bleach**

**Perfume, cologne, toilet water**

**Cosmetic lotions, creams**

**Glues, adhesives**

**Paint**

**Insecticides (excluding  
mothballs)**

**Rodenticides**

**Corrosive acids, alkalies**

**Vitamins, minerals**

### ***Miscellaneous internal and external medicines:***

**Aspirin**

**Miscellaneous analgesics**

**Antihistamines, cold medications**

**Antiseptic medications**

**Psychopharmacologic agents**

**Cough medicines**

**Hormones**

**internal antibiotics**

# *Anticipated Turn-around Times for a Full-Service Laboratory*

## *Anticipated Turnaround Time'*

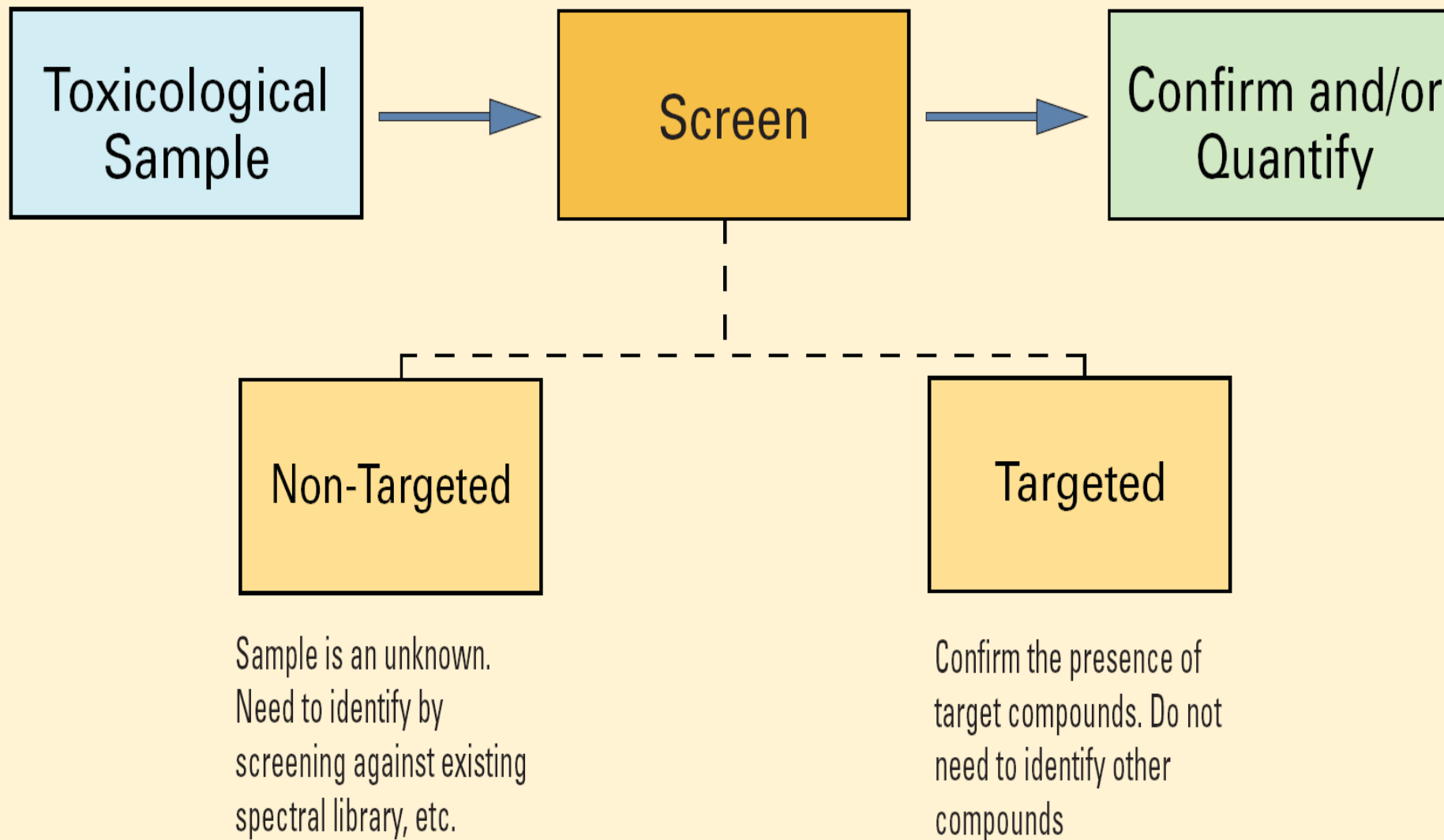
### *Quantitative Tests (Serum)*

Acetaminophen	30 min - 45 min
Carbamazepine	45 min - 1 hr
Cholinesterase	1 hr - 2 hr
Cyclosporine	1 hr - 1.5 hr
Digoxin	1 hr - 1.5 hr
Ethanol	15 min - 30 min
Ethylene Glycol	1 hr - 2 hr
Isopropanol	20 min - 45 min
Lidocaine	45 min - 1 hr
MEGX	45 min - 1 hr
Methanol	20 min - 45 min
Procainamide & NAPA	1 hr - 2 hr
Phenobarbital	45 min - 1 hr
Phenytoin	45 min - 1 hr
Quinidine	45 min - 1 hr
Salicylates	30 min - 45 min
Theophylline	45 min - 1 hr
Valproic Acid	45 min - 1 hr

### *Qualitative Tests (Urine)*

Comp Drug Screen <sup>3</sup>	2hr -4hr
Stimulant Panel	1 hr - 2.5 hr

## *Typical Screening Workflow*



*Relative Specificity, Sensitivity, and Speed of Processing of Different Laboratory Techniques*

<i>Laboratory Technique</i>	<i>Specificity</i>	<i>Sensitivity</i>	<i>Speed</i>
<b>Spectrophotometry</b>	+	++	+++
<b>Immunology</b>	+(+)	++	++++
<b>Thin-layer (TLC) Chromatography</b>	++	++	++
<b>High-performance liquid (HPLC) Chromatography</b>	+++	++	+ - +++
<b>Gas-liquid (GLC) Chromatography</b>	+++	++	+
<b>Gas chromatography-mass-spectrometry</b>	++++	++++	+
<b>Liquid chromatography-mass-spectrometry</b>	++++	++++	+++

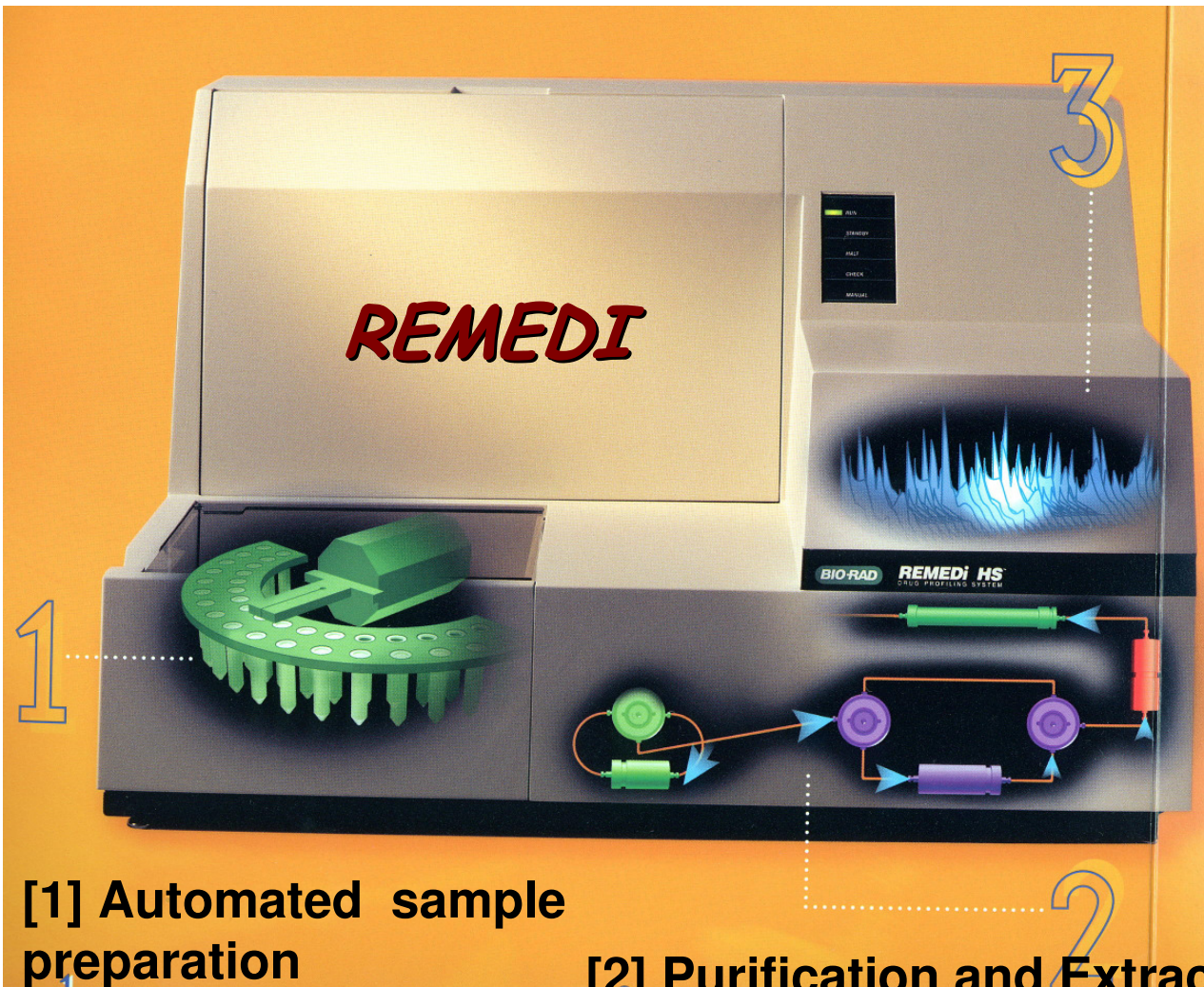
## *Causes of false immunological screening...*

- ***Screen test does not seek the drug***  
structural dissimilarity from drug class  
prototype e.g. fentanyl
- ***too broad antibody specificity***  
- group test e.g. benzodiazepines
- ***cross reactivity with other compound of different class***
- ***Toxicokinetic characteristics***  
large volume of distribution  
Short elimination half-life

*Relative Specificity, Sensitivity, and Speed of Processing of Different Laboratory Techniques*

<b>Laboratory Technique</b>	<b>Specificity</b>	<b>Sensitivity</b>	<b>Speed</b>
<b>Spectrophotometry</b>	+	++	+++
<b>Immunology</b>	+(+)	++	+++++
<b>Thin-layer (TLC) Chromatography</b>	++	++	++
<b>High-performance liquid (HPLC) Chromatography</b>	+++	++	+ - +++
<b>Gas-liquid (GLC) Chromatography</b>	+++	++	+
<b>Gas chromatography-mass-spectrometry</b>	+++++	+++++	+
<b>Liquid chromatography-mass-spectrometry</b>	+++++	+++++	+++





**[1] Automated sample preparation**

**[2] Purification and Extraction Cartridges remove impurities on-line, Separation Cartridges 1 and 2 separate drugs of diverse classes into discrete peaks.**

**[3] Scanning UV detector, resulting in a unique spectral fingerprint for each unknown peak.**

**[4] "Expert System" comparing retention time data and five distinct spectral characteristics of each unknown peak to an extensive library of drugs and metabolites.**

## ***REMEDI: Bestimmung der relativen Retentionszeit***

**RRT 1:** RT of a peak relative to the first internal standard.

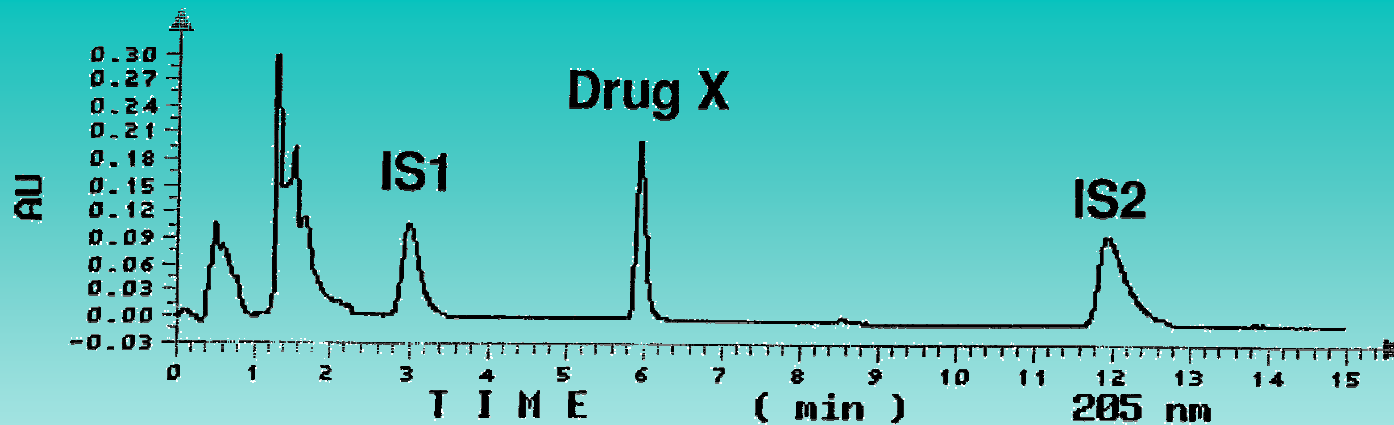
**RRT 2:** RT of a peak relative to the second internal standard.

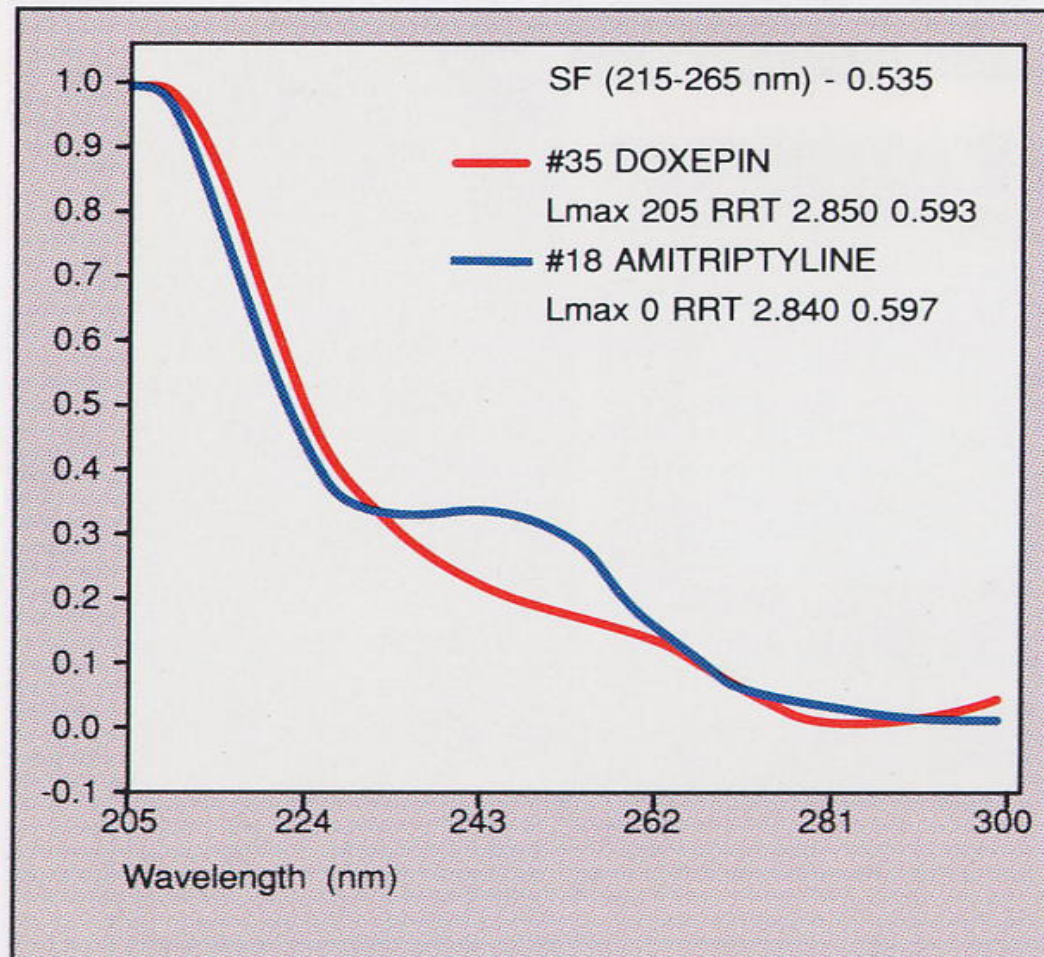
$$\frac{RT_x - dt}{RT_{I.S.} - dt}$$

$$\frac{RT_x - dt}{RT_{I.S.} - dt}$$

$$RRT1_x \approx 2.0$$

$$RRT2_x \approx 0.5$$





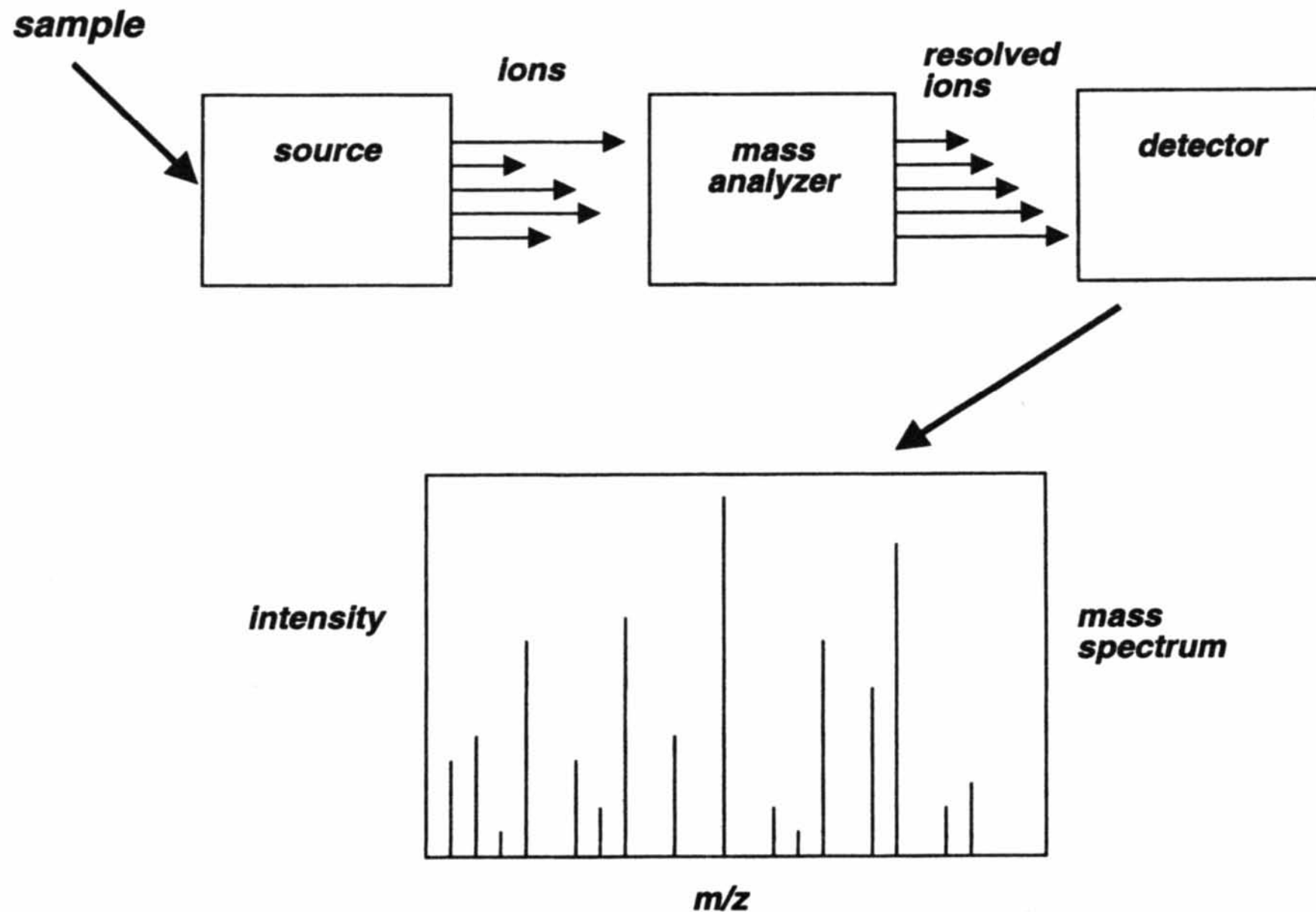
*Drug spectra are compared to a known library of candidates stored in memory.*

Derzeit ist eine  
Spektrenbibliothek von  
ca. 1000  
pharmazeutischen  
Verbindungen  
verfuegbar

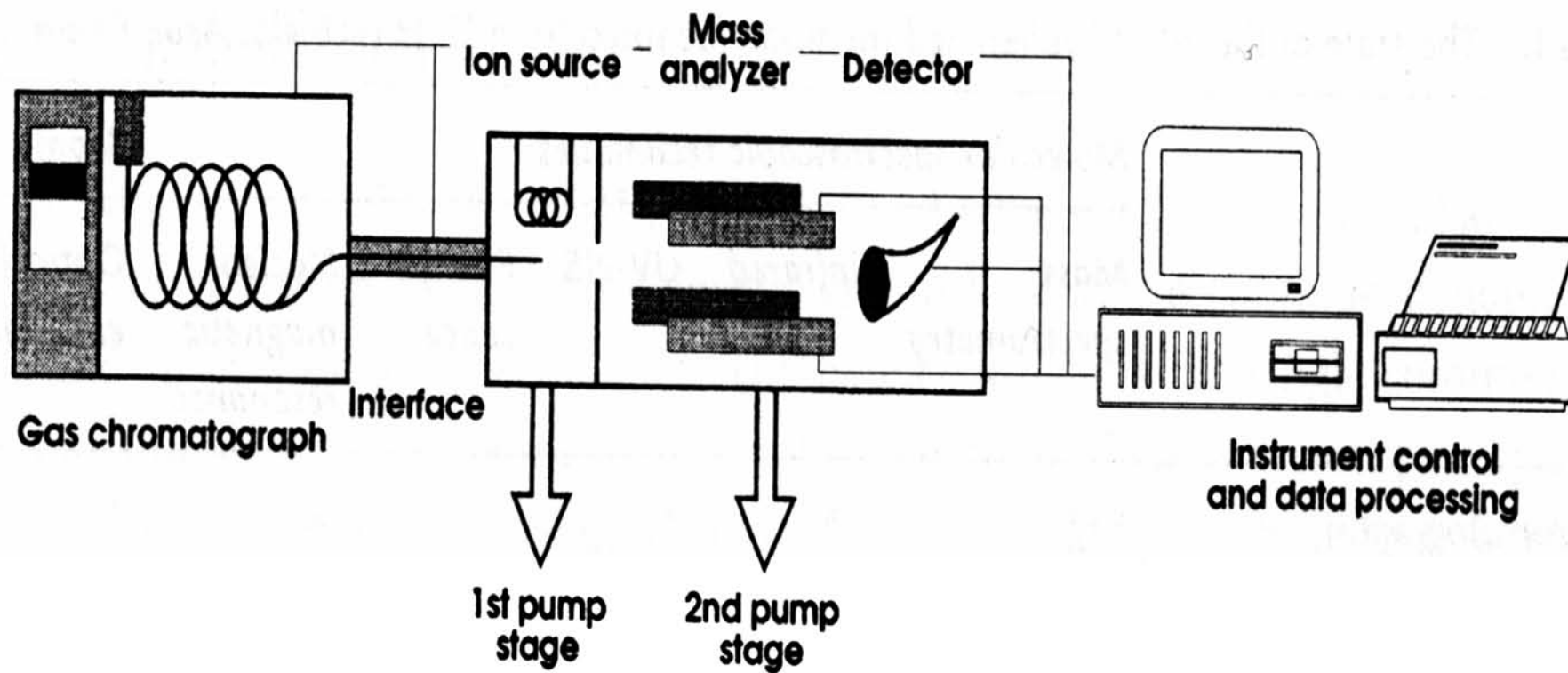


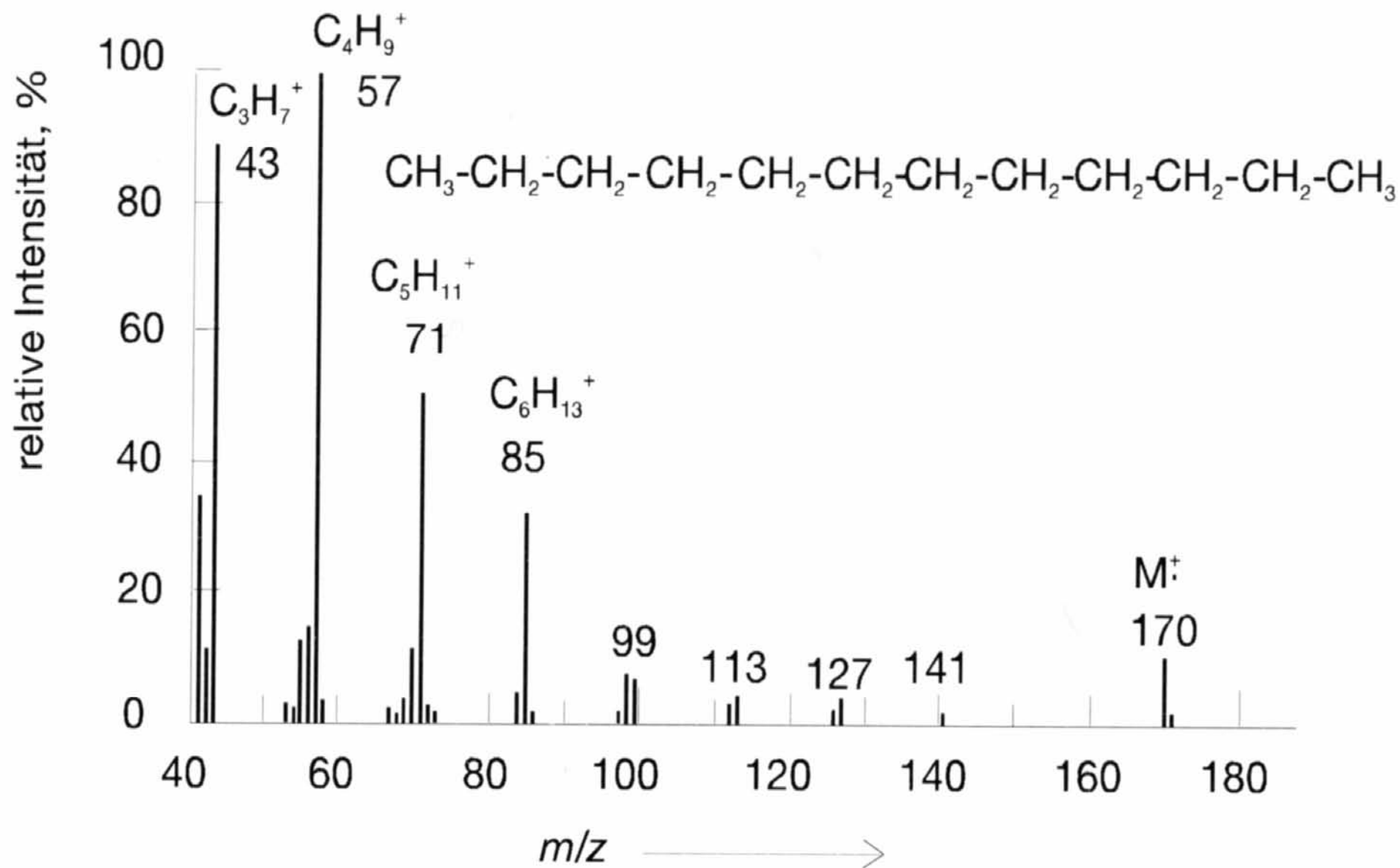
*Relative Specificity, Sensitivity, and Speed of Processing of Different Laboratory Techniques*

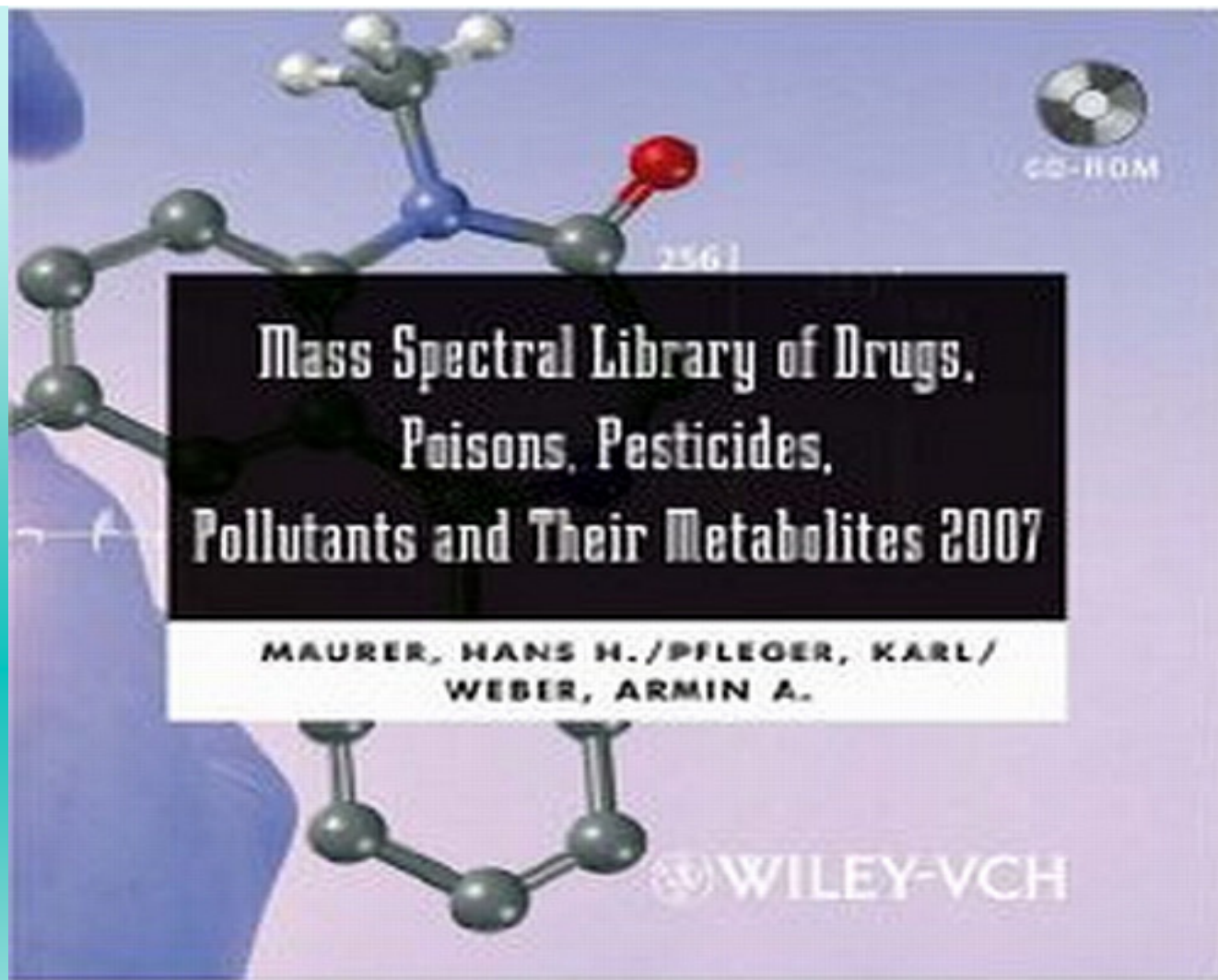
<b>Laboratory Technique</b>	<b>Specificity</b>	<b>Sensitivity</b>	<b>Speed</b>
<b>Spectrophotometry</b>	<b>+</b>	<b>++</b>	<b>+++</b>
<b>Immunology</b>	<b>+(+)</b>	<b>++</b>	<b>+++++</b>
<b>Thin-layer (TLC) Chromatography</b>	<b>++</b>	<b>++</b>	<b>++</b>
<b>High-performance liquid (HPLC) Chromatography</b>	<b>+++</b>	<b>++</b>	<b>+ - +++</b>
<b>Gas-liquid (GLC) Chromatography</b>	<b>+++</b>	<b>++</b>	<b>+</b>
<b>Gas chromatography-mass-spectrometry</b>	<b>+++++</b>	<b>+++++</b>	<b>+</b>
<b>Liquid chromatography-mass-spectrometry</b>	<b>+++++</b>	<b>+++++</b>	<b>+++</b>



***The principle of mass spectrometric analysis***







Contains 7840 data sets. Over 3300 data sets are from *metabolites*, over 2300 from *acetylated*, over 1000 from *methylyated*, over 700 from *trimethylsilylated*, over 400 from *trifluoroacetylated*, over 200 each from *pentafluoropropionylated* or *heptafluorobutyrylated compounds*.



# Systematic evaluation of 1-chlorobutane for liquid-liquid extraction of drugs

U. Demme<sup>1\*</sup>, J. Becker<sup>2</sup>, H. Bussemas<sup>3</sup>, T. Daldrup<sup>4</sup>, F. Erdmann<sup>5</sup>, M. Erkens<sup>6</sup>, P.X. Iten<sup>7</sup>, H. Käferstein<sup>8</sup>, K.J. Lusthof<sup>9</sup>, H.J. Magerl<sup>10</sup>, L.v. Meyer<sup>11</sup>, A. Reiter<sup>12</sup>, G. Rochholz<sup>13</sup>, A. Schmoldt<sup>14</sup>, E. Schneider<sup>15</sup>, H.W. Schütz<sup>13</sup>, T. Stimpfl<sup>16</sup>, F. Tarbah<sup>17</sup>, J. Teske<sup>18</sup>, W. Vycudilik<sup>16</sup>, J.P. Weller<sup>18</sup>, W. Weinmann<sup>19</sup>



\*On behalf of the "Workgroup Extraction" of the GTFCh (Society of Toxicological and Forensic Chemistry, Germany)

Institutes of Forensic Medicine of <sup>1</sup>Jena, <sup>2</sup>Mainz, <sup>4</sup>Duesseldorf, <sup>5</sup>Giessen, <sup>7</sup>Zuerich (CH), <sup>8</sup>Cologne, <sup>10</sup>Wuerzburg, <sup>11</sup>Munich, <sup>12</sup>Luebeck, <sup>13</sup>Kiel, <sup>14</sup>Hamburg, <sup>16</sup>Vienna (A), <sup>18</sup>Hannover, <sup>19</sup>Freiburg and <sup>3</sup>Praxis Labormedizin Dortmund, <sup>6</sup>Clin.-Chem. Central Laboratory, Aachen, <sup>9</sup>Nat. For. Inst., Den Haag (NL), <sup>15</sup>LKA Baden-Wuerttemberg, Stuttgart <sup>17</sup>Dubai Police Dept.

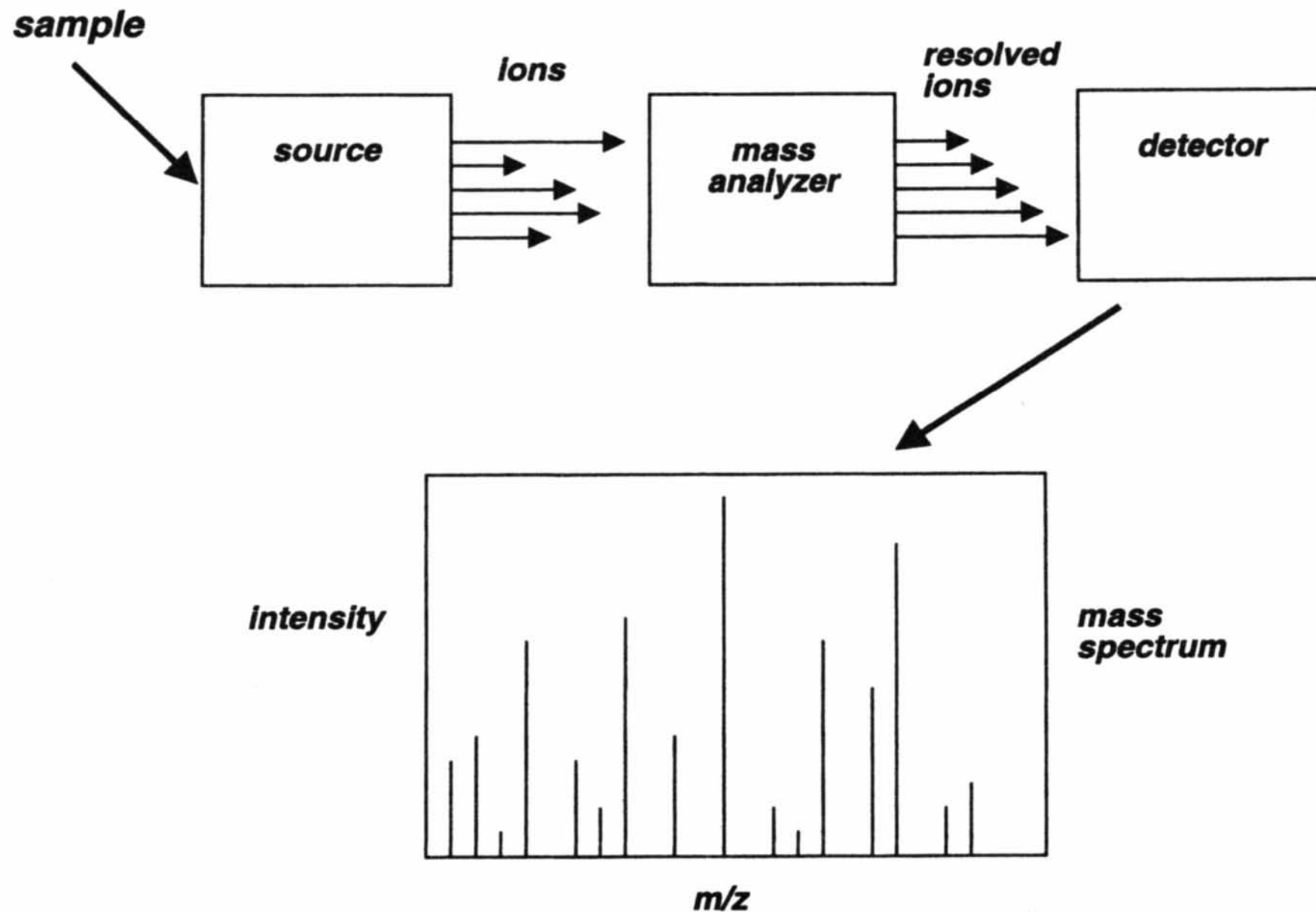
N°	drug	Y°	Ref.
24	Acebutolol	0.05	
25	Adenosin	0	
26	Ajmaline	0.5	
27	Alfentanil	1	
28	Alimemazine	1	
29	Alprazolam	0.95	[43-45]
30	Alprenolol	1	
31	Amantadine	0.5	
32	Amfebutamon	1	
33	Amfepramon	1	
34	Amfetaminil	1	
35	Amiodarone	0.95	
36	Amisulprid	0.6	
37	Amitriptyline	1	[46]
38	Amitriptyline-Oxid	0.1	
39	Amlodipine	1	
40	Amphetamine	0.5	[47-55]
41	Apomorphin	0.8	
42	Aprindin	0.95	
43	Articain	1	
44	Atenolol	0	
45	Atropine	0.6	
46	Azathioprine	0	
47	Azinphos-ethyl	1	
48	Azinphos-methyl	1	
49	Benperidol	1	
50	Benserazid	0	
51	Benzatropin	1	
52	Benzoylecgonine	0	
53	Betaxolol	1	
54	Biperidene	1	
55	Bisacodyl	0.7	
56	Bisoprolol	0.9	
57	Bromazepam	0.9	
58	Bromocriptin	1	
59	Bromophos-ethyl	1	

(cont. : see [www.gtfc.org/chlorobutaneextraction.pdf](http://www.gtfc.org/chlorobutaneextraction.pdf))

*Y<sub>o</sub>* is the extraction  
yield in the  
organic phase

*Relative Specificity, Sensitivity, and Speed of Processing of Different Laboratory Techniques*

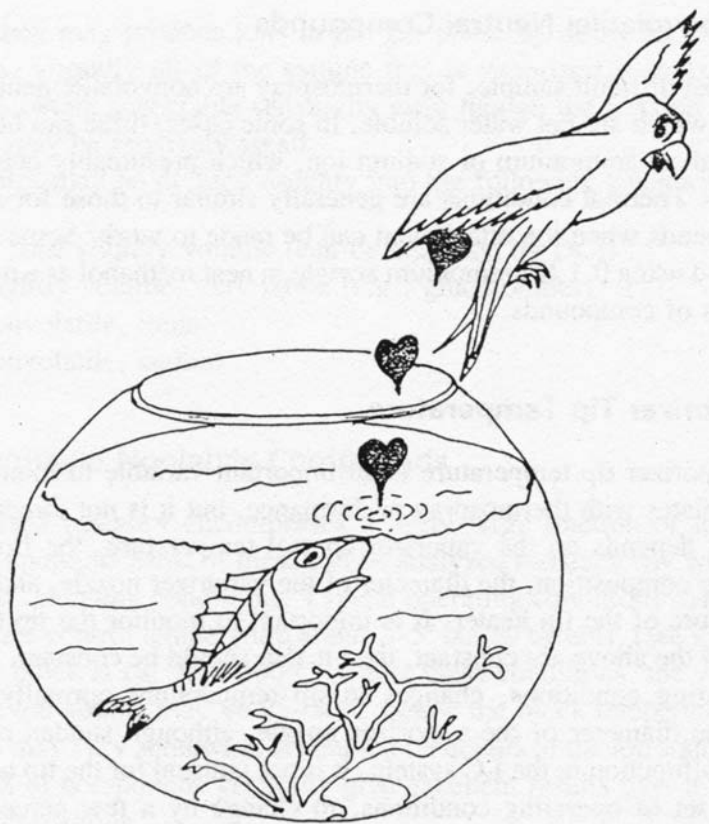
<b>Laboratory Technique</b>	<b>Specificity</b>	<b>Sensitivity</b>	<b>Speed</b>
<b>Spectrophotometry</b>	<b>+</b>	<b>++</b>	<b>+++</b>
<b>Immunology</b>	<b>+(+)</b>	<b>++</b>	<b>+++++</b>
<b>Thin-layer (TLC) Chromatography</b>	<b>++</b>	<b>++</b>	<b>++</b>
<b>High-performance liquid (HPLC) Chromatography</b>	<b>+++</b>	<b>++</b>	<b>+ - +++</b>
<b>Gas-liquid (GLC) Chromatography</b>	<b>+++</b>	<b>++</b>	<b>+</b>
<b>Gas chromatography-mass-spectrometry</b>	<b>+++++</b>	<b>+++++</b>	<b>+</b>
<b>Liquid chromatography-mass-spectrometry</b>	<b>+++++</b>	<b>+++++</b>	<b>+++</b>



***The principle of mass spectrometric analysis***

# Liquid-chromatography - Mass Spectrometry

*- a difficult marriage!!*



HPLC

=

Liquid at 1 atm



???

MS

=

Gas at  $10^{-6}$  atm



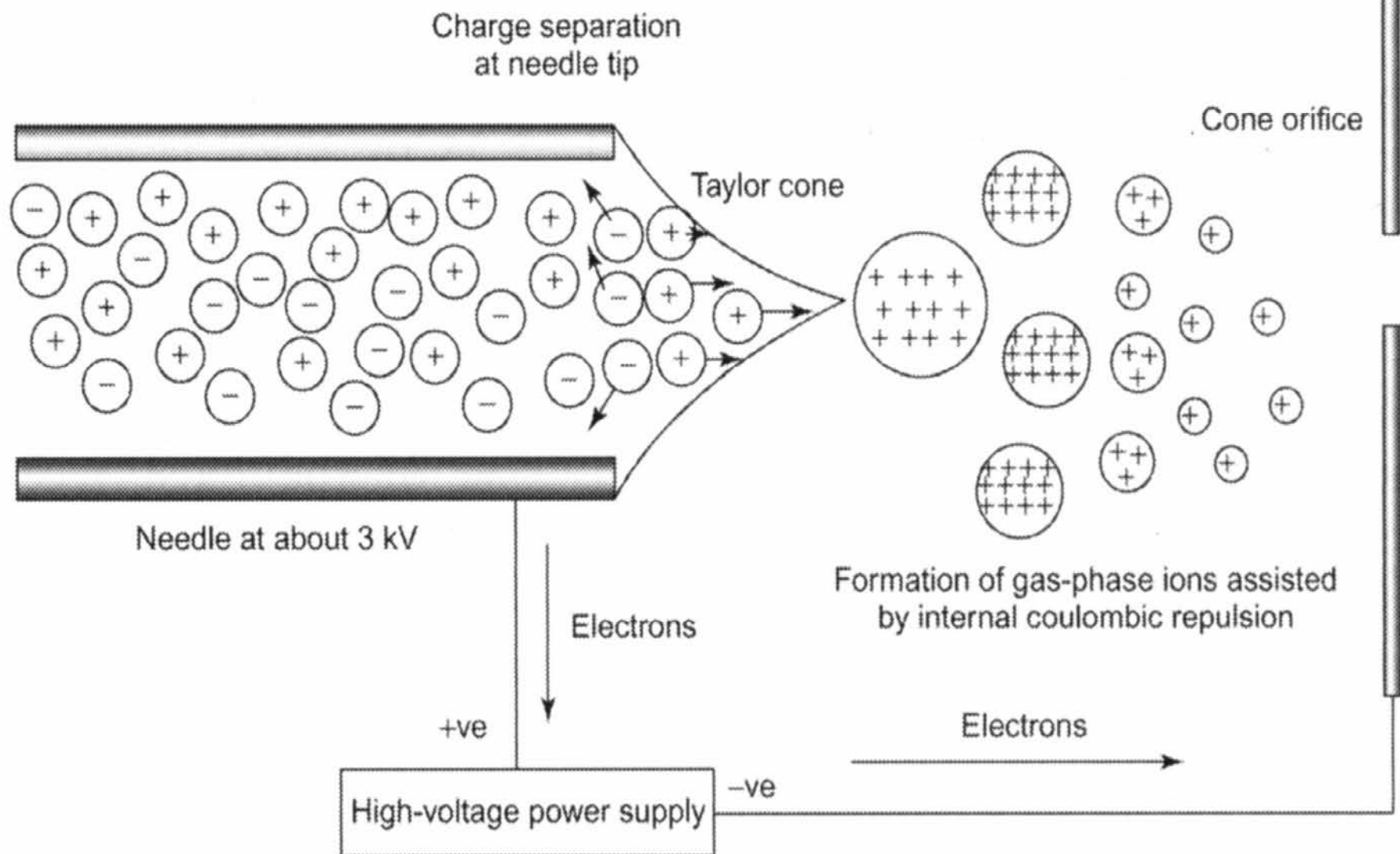
Not a new  
technique...

**First ionization with an 'ESI'  
was done more than 200  
years ago...**

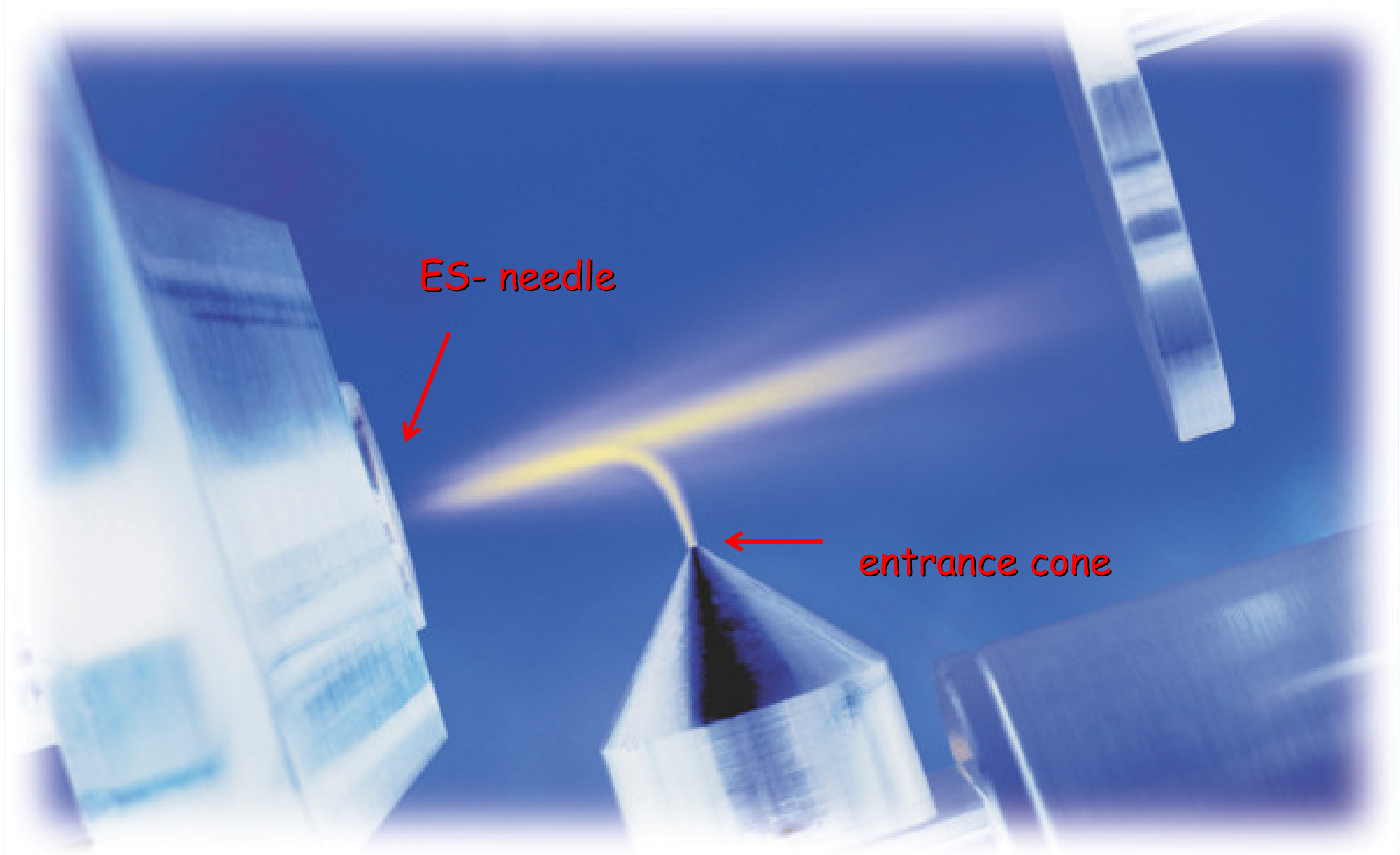


*Electrostatic spraying of water in  
the 18th century.*

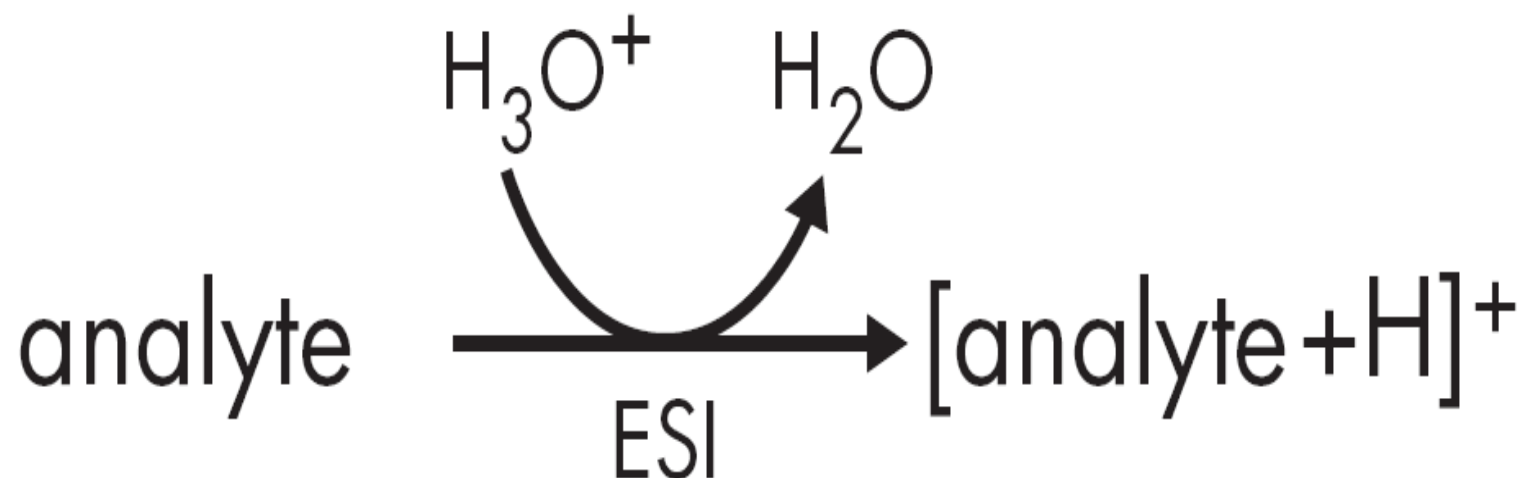
# The electrospray process (ESI)



# Electro-Spray



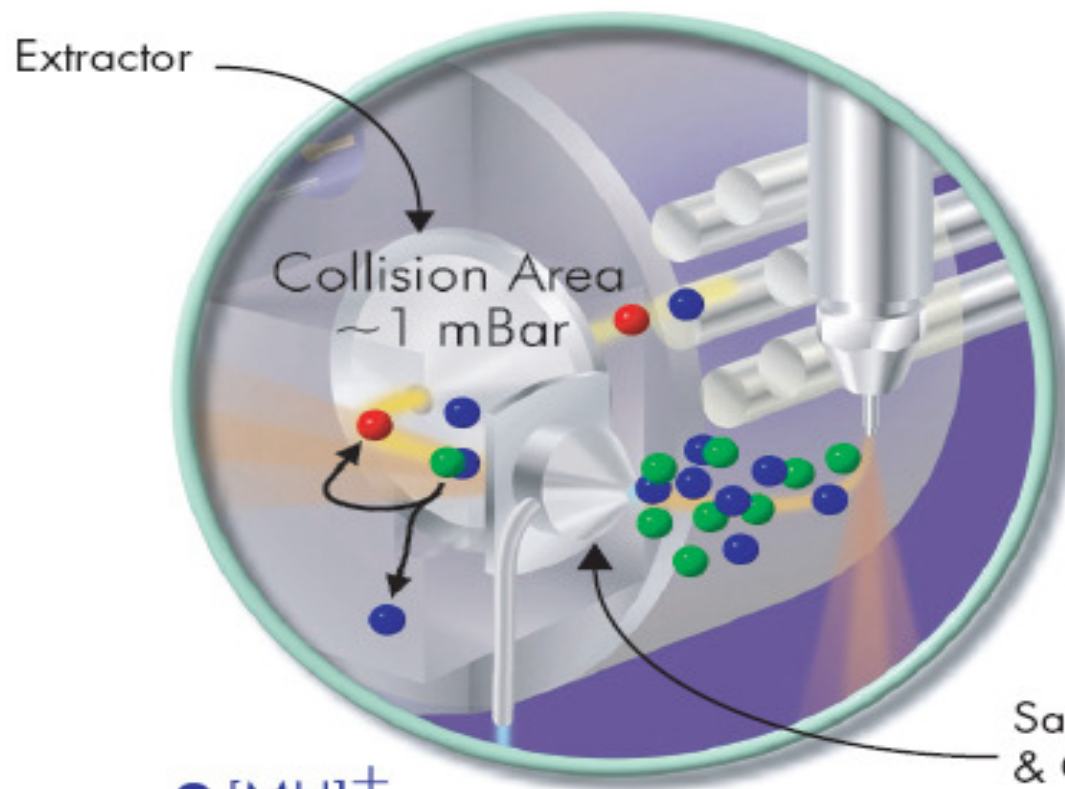




### ***Atmospheric Pressure Ionisation (API) process***

This *soft ionisation process* leads to cations in positive ion mode and anions in negative ion mode which are generally stable.

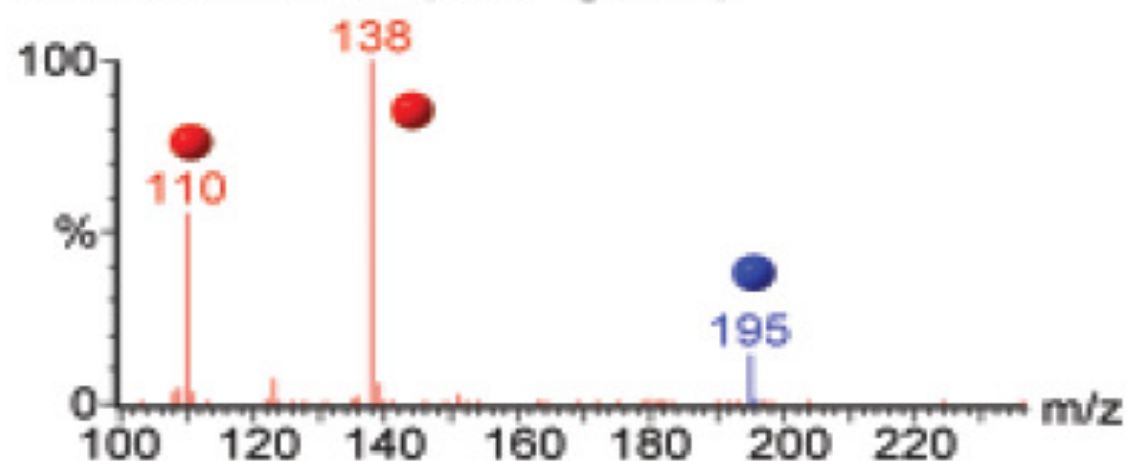
The molecular ions can be fragmented in the *source region* of LC/MS instruments

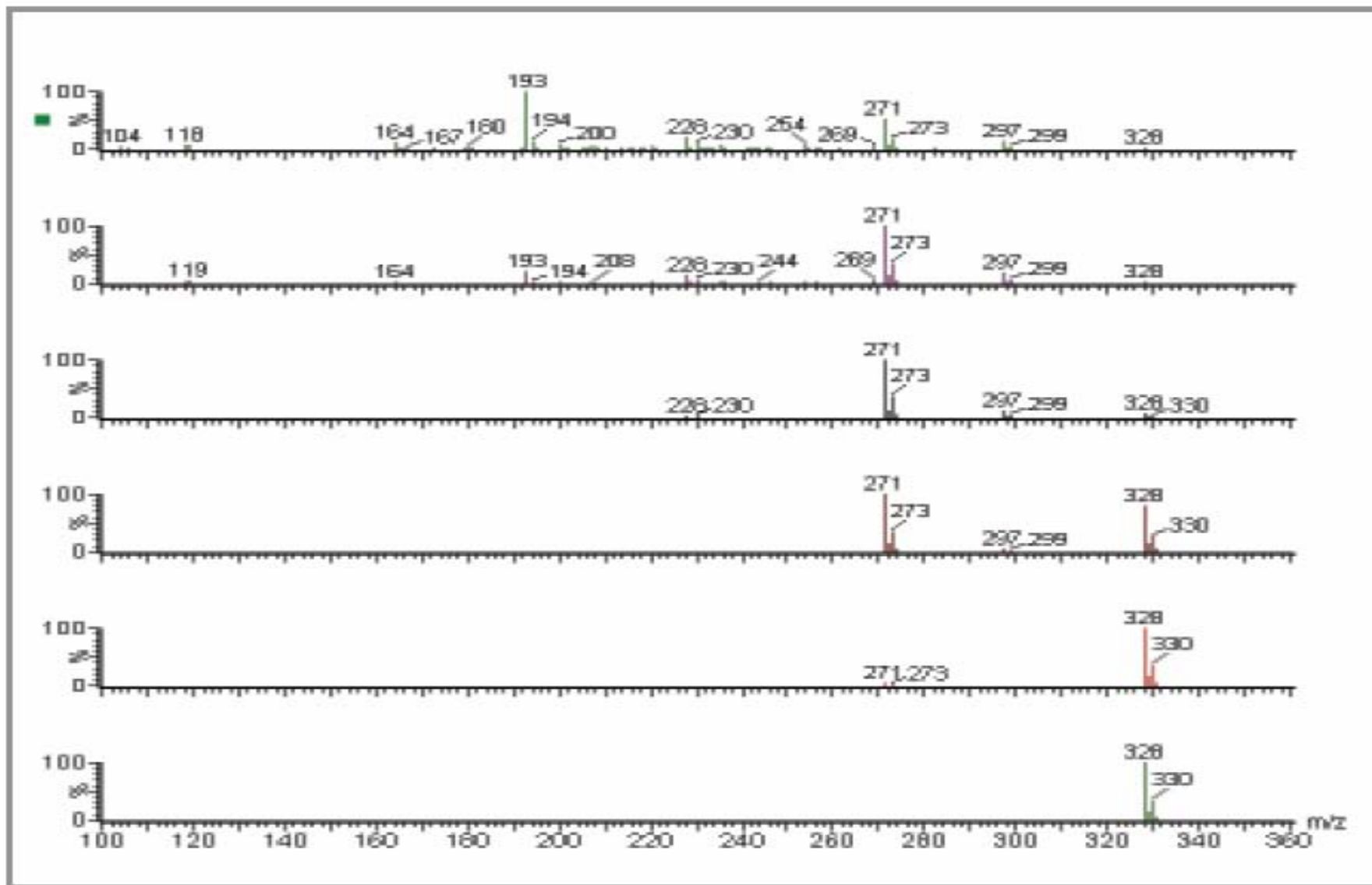


*InSource-CID fragmentation  
of the molecular ion ( $m/z$  195)  
of caffeine at 60V in the  
(Quattro micro) API ion source*

- $[MH]^+$
- $N_2$  molecules
- $[Fragment]^+$

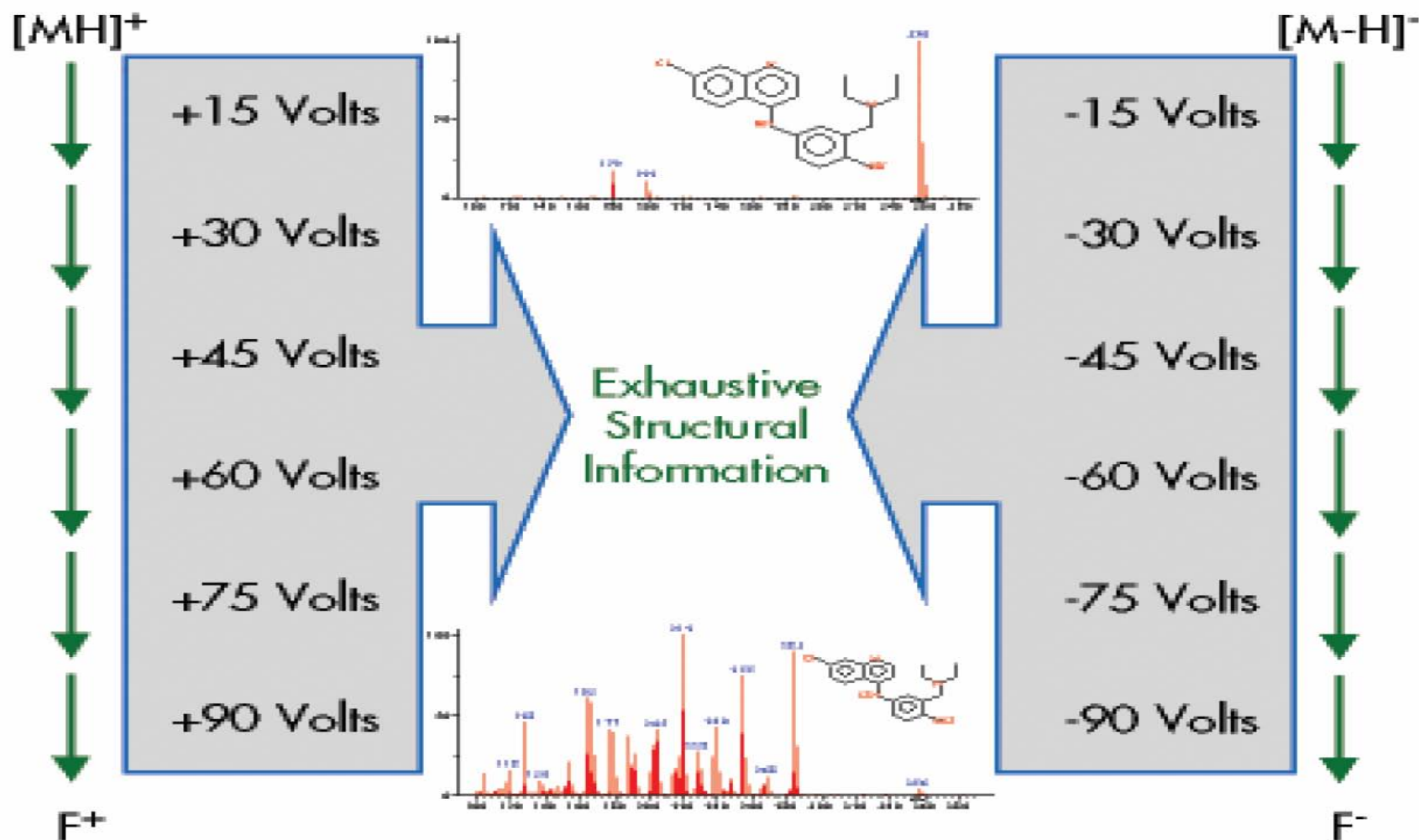
ToxML 30 Caffein [ESI+ @ 60V]

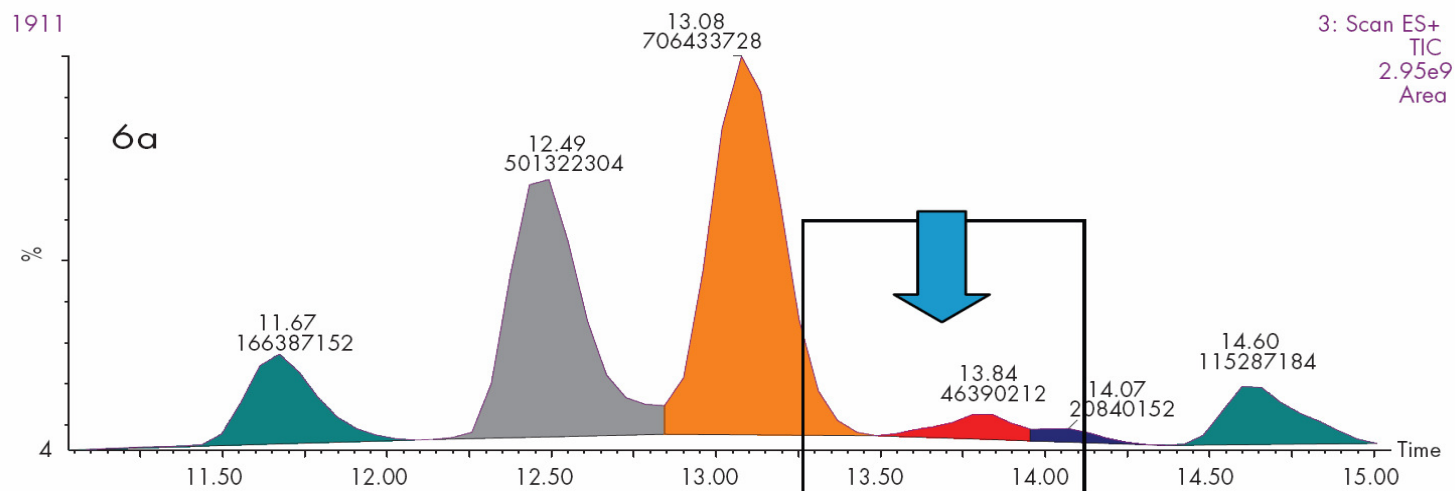




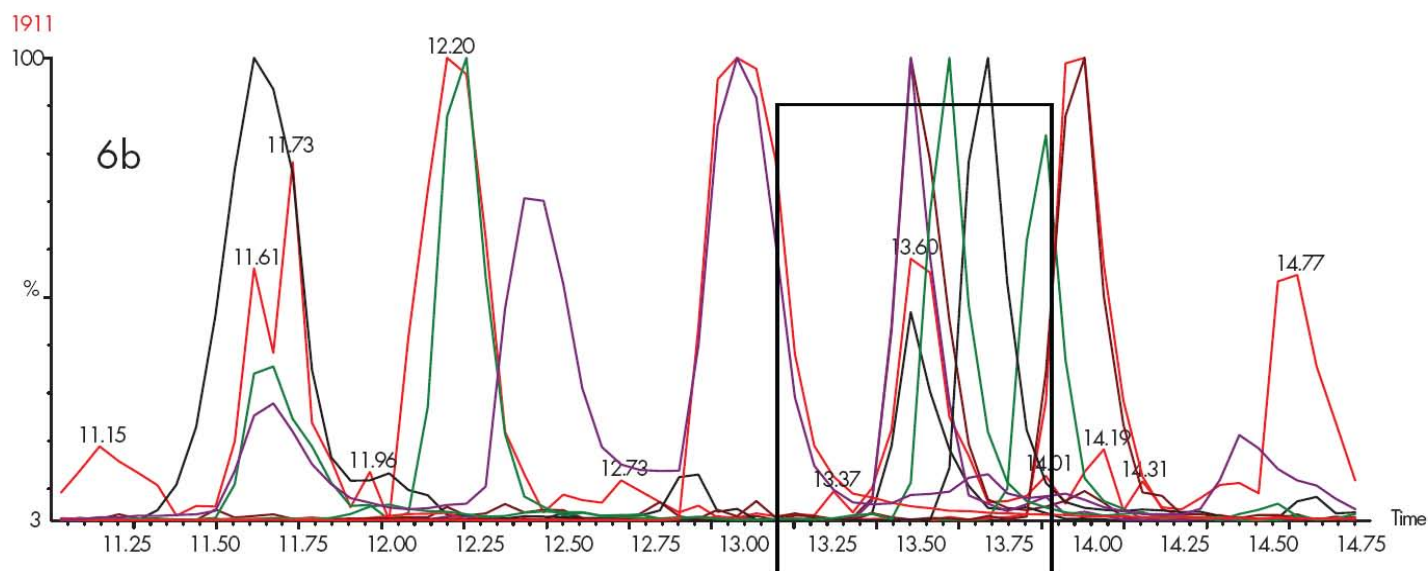
Mass spectra of Loxapine, recorded at *6 different CV values* using in-source CID.

The *degree of fragmentation* increases with the cone voltage

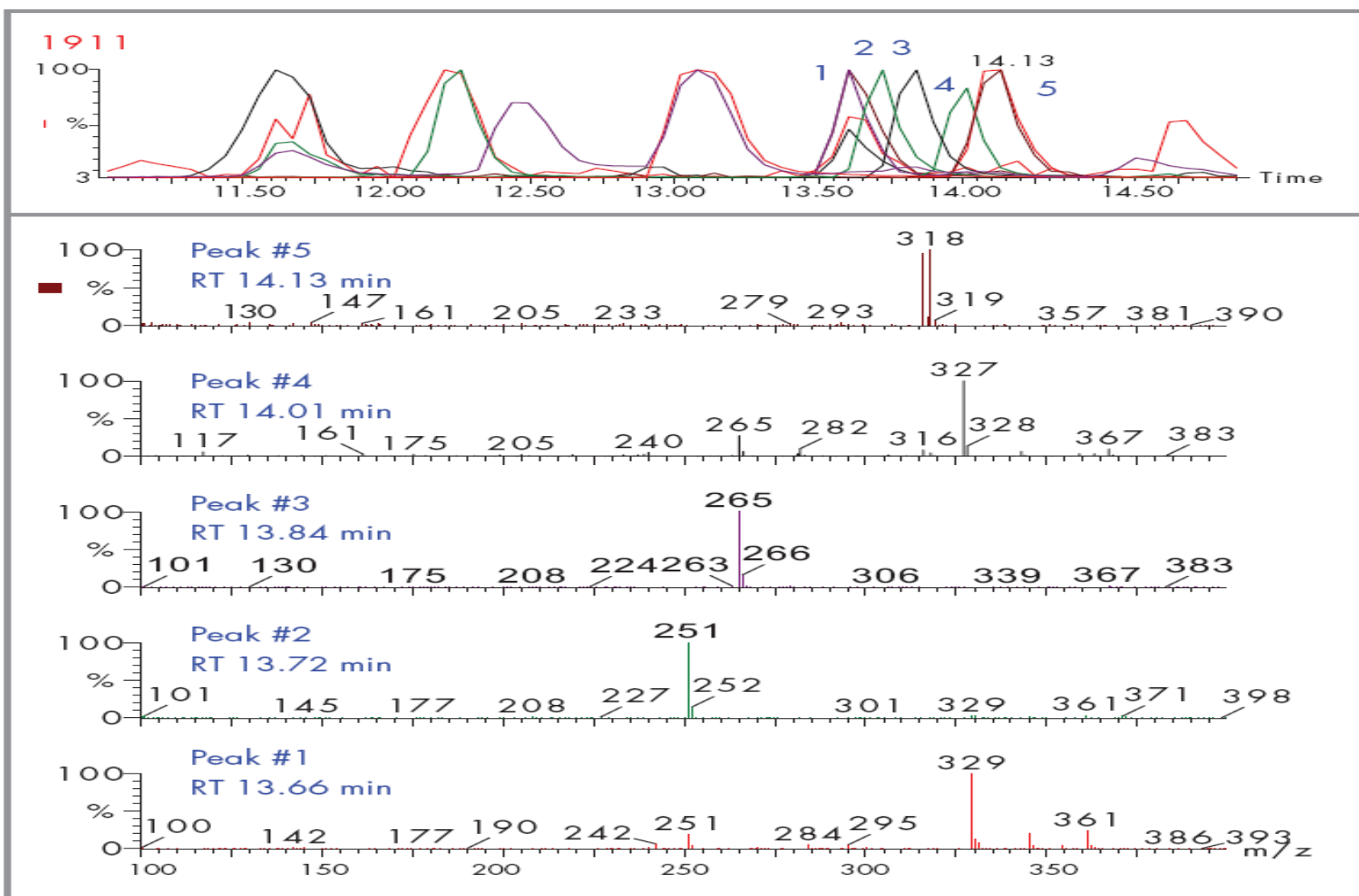




*Section of chromatogram area acquired in positive ESI @ 30 V.  
Here the total ion chromatogram (TIC) indicates that only one component elutes at 13.8 min*



*Section of chromatogram area for function 3 acquired in positive ESI @ 30 V. Using extracted ion chromatograms shows that at least three components elute between 13.5 and 13.8 minutes.*



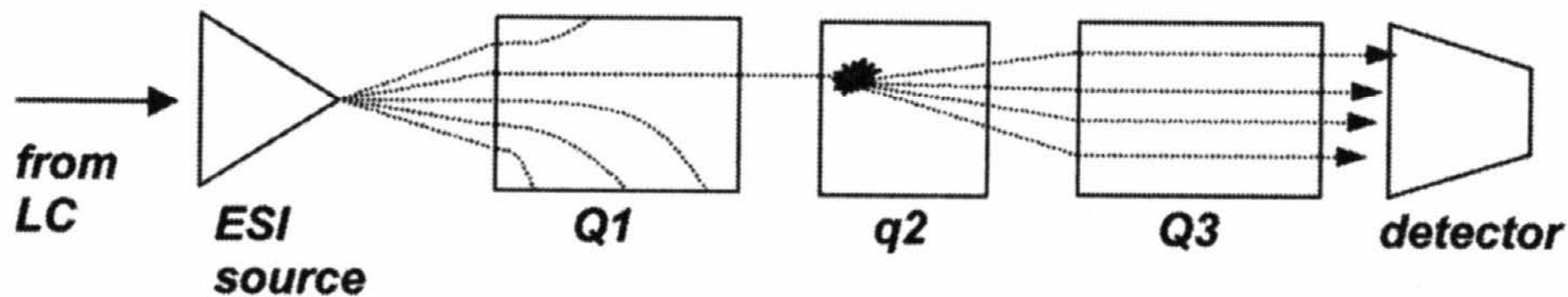
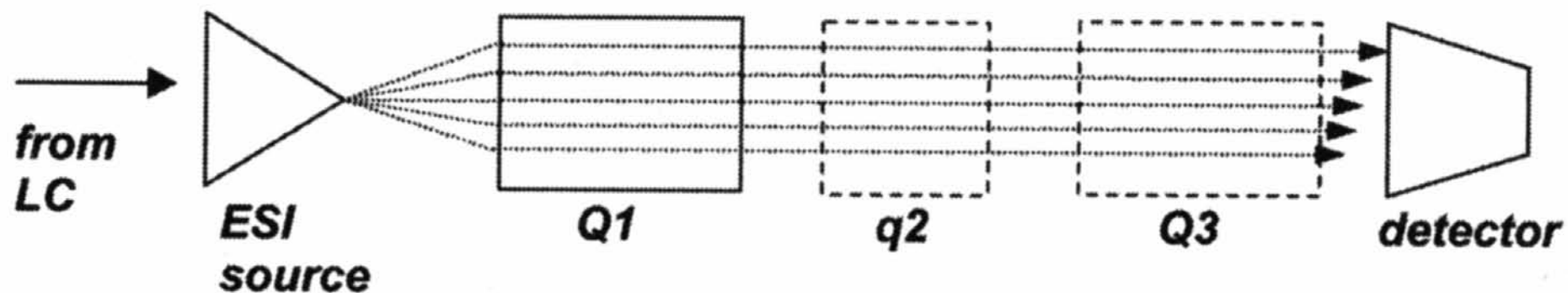
*Chromatogram acquired in positive ESI @ 30 V and corresponding mass spectra of 5 components. Automated spectral deconvolution allows extraction of clean mass spectra that can be used for library searching.*



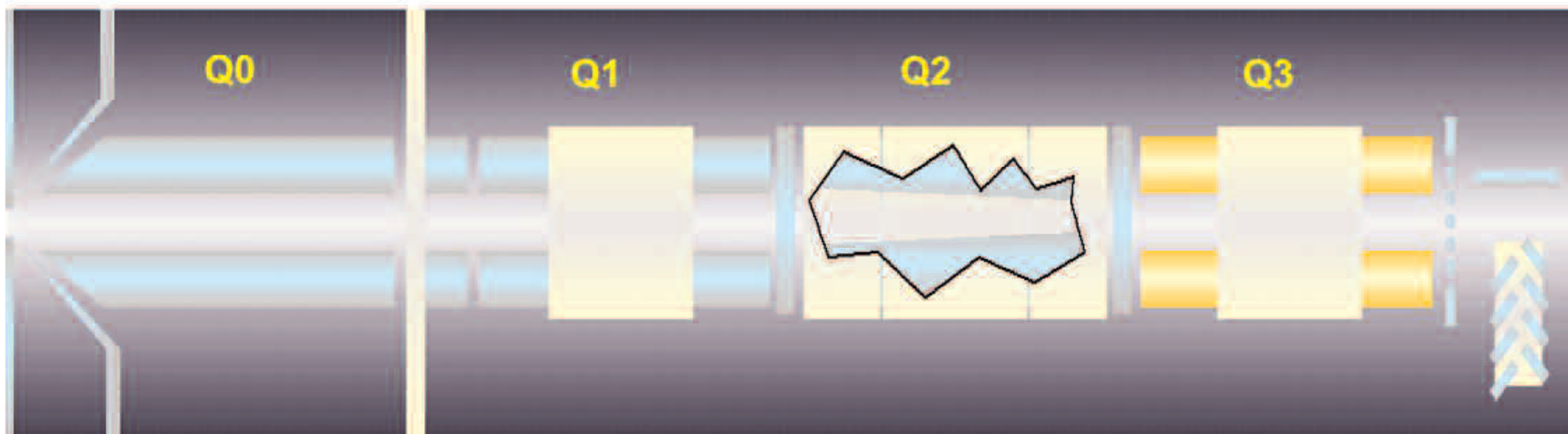
**Candidate Average Fit (%)**

#	Analyte Name	Status	Origin	6 Functions
1	Nicotine	Unexpected molecule	Smoker / Contamination	56.1
2	Trimetazidine	Expected molecule	Medication	63.3
3	Acetaminophen	Expected molecule	Medication	62.3
4	Caffeine	Expected molecule	Medication	74.0
5	Quinine	Expected molecule	Medication	70.3
6	Zolpidem	Expected molecule	Medication	94.7
7	Meprobamate	Unexpected molecule	Unknown	55.3
8	Mianserin	Expected molecule	Medication	67.1
9	Acepromazine	Unexpected molecule	Unknown	57.6
10	Bromazepam	Unexpected molecule	Unknown	53.1
11	Hydroxyzine	Expected molecule	Medication	88.2
12	Propoxyphene	Expected molecule	Medication	62.1
13	Tramadol	Expected molecule	Medication	Not found

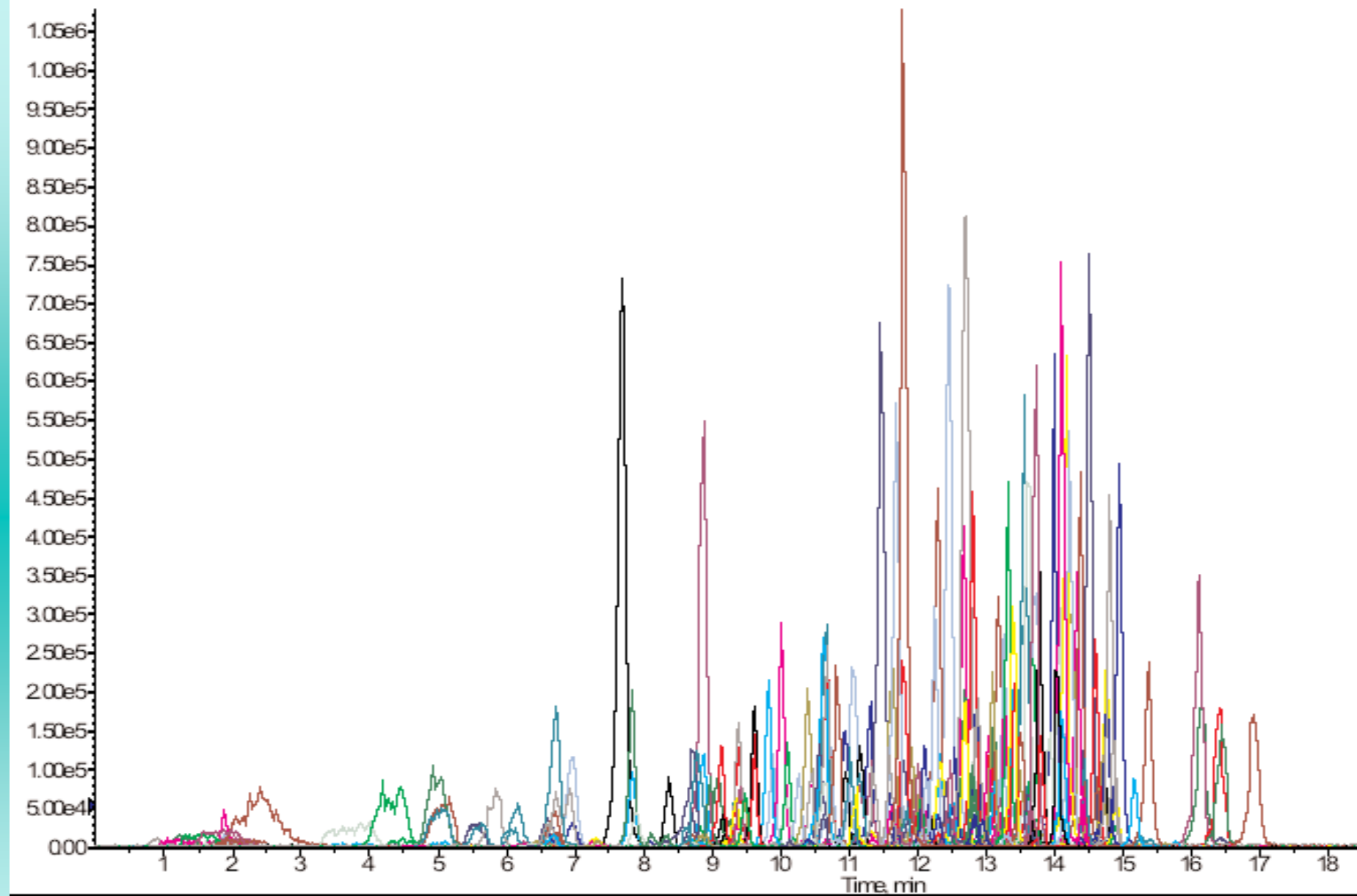
## *Funktion eines Triple-Quad Mass Spectrometer*





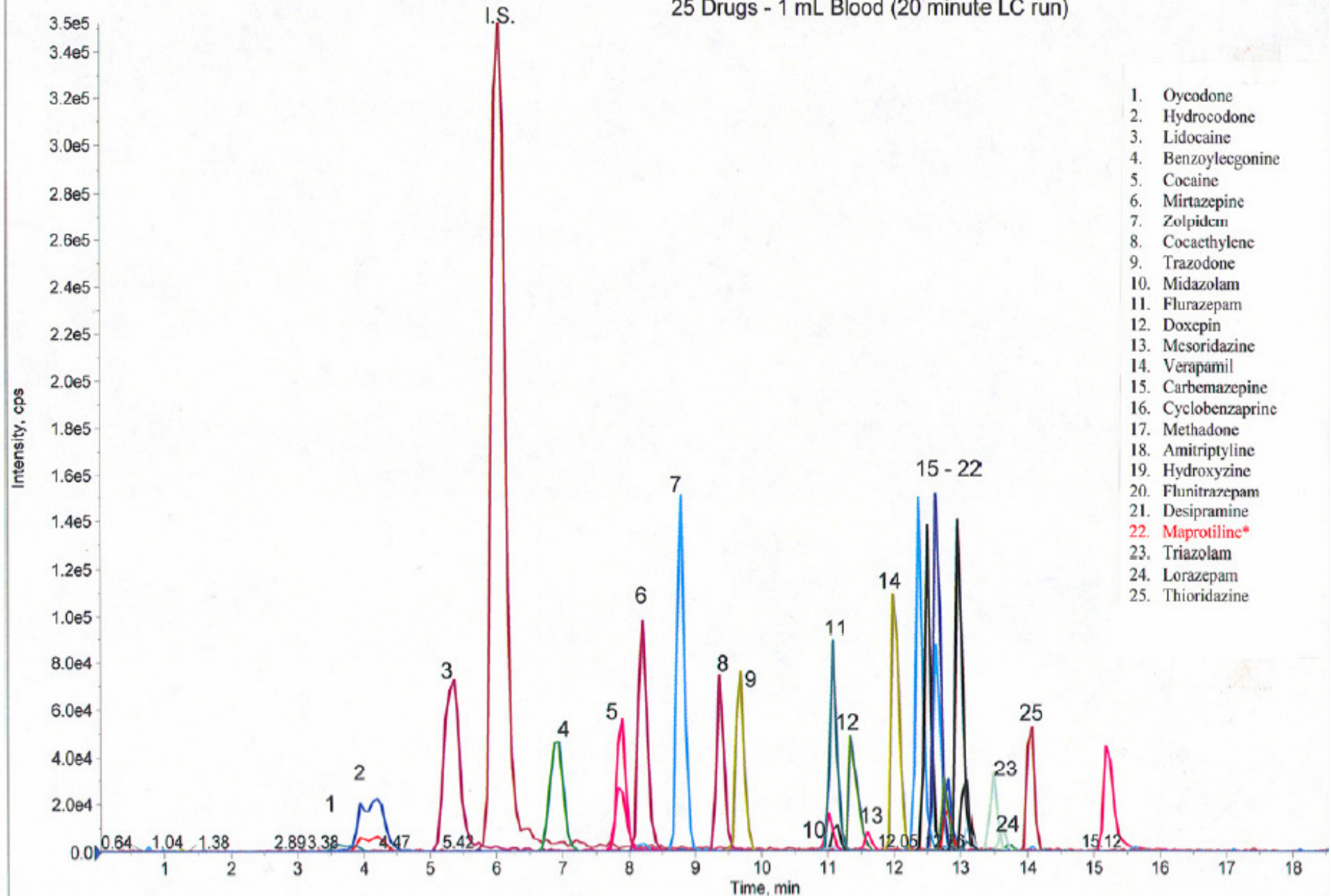


Schematic diagram of the ion path of a hybrid triple quadrupole/linear ion trap (QQQ/LIT) instrument.  
The ion path is virtually identical to a standard triple quadrupole, except Q3 has been modified to operate not only as a quadrupole, but also as an ion trap.



**Analysis and detection of 300 compounds in a single assay. All compounds were analyzed in under 20 minutes.**

# 25 Drugs - 1 mL Blood (20 minute LC run)

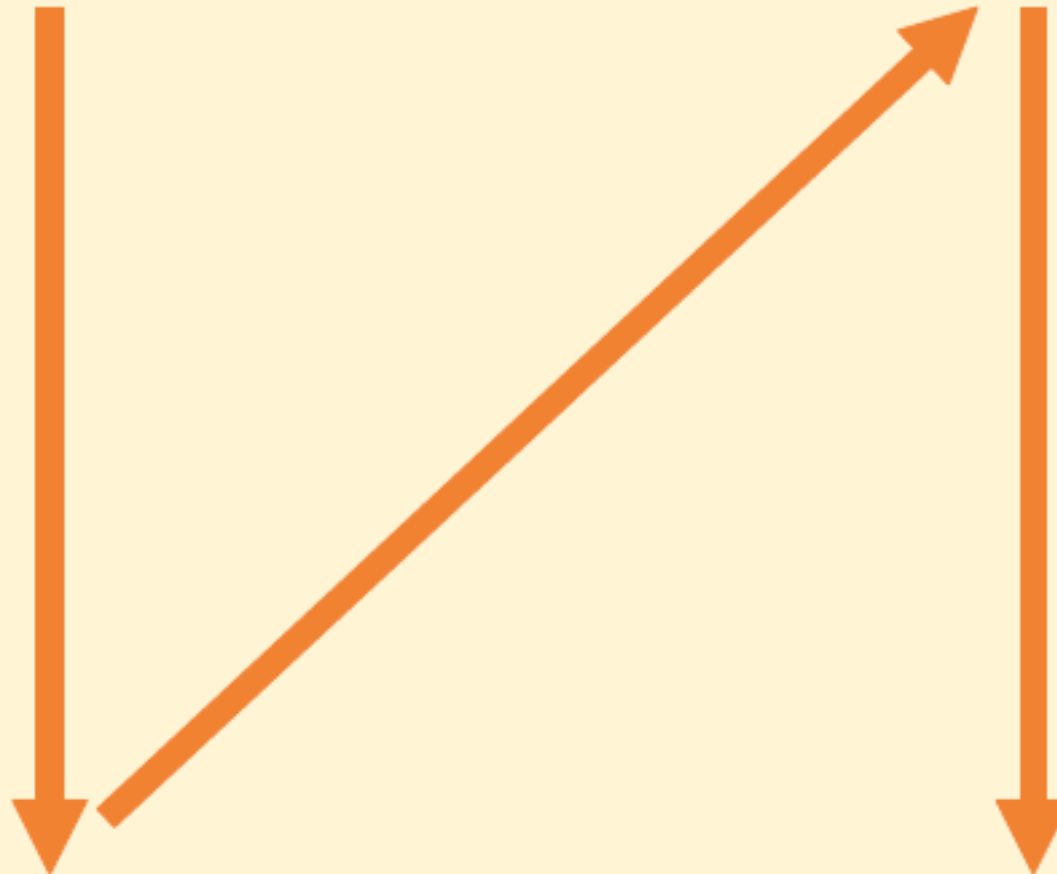


**Scan Event 1**  
+ Full Scan MS

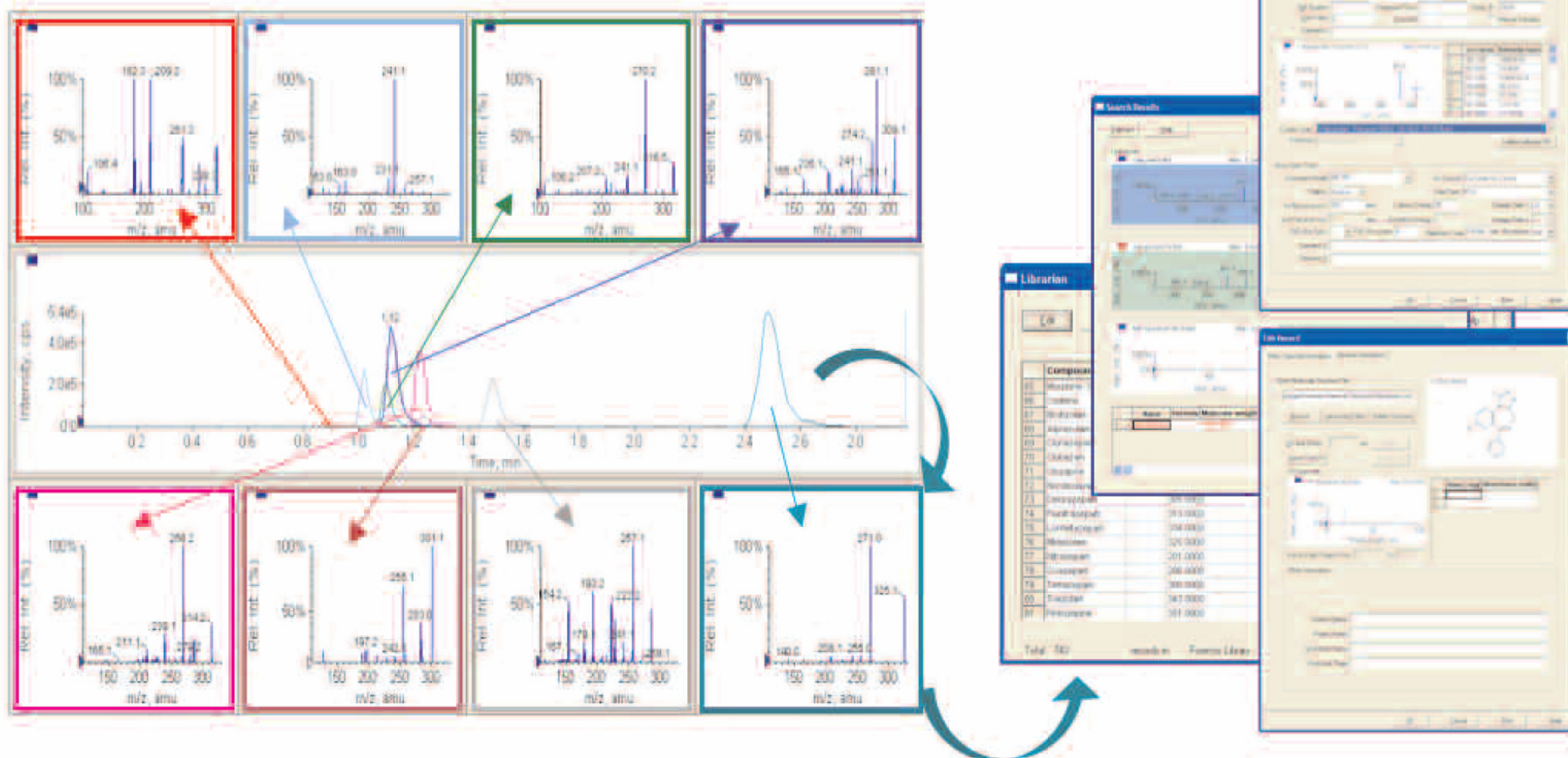
**Scan Event 7**  
– Full Scan MS

**Scan Events 2–6**  
+ MS/MS on parent list

**Scan Event 8**  
– MS/MS on parent list



## Library Search Results...



Analysis and detection on a QQQ/LIT instrument using MRM-IDA. The chromatograms for each MRM transition is shown, along with the LIT MS/MS spectra that are acquired from each signal. Quality spectra were obtained, even from compounds obtained at low levels. Also, unique spectrawere acquired even in cases where two compounds co-eluted. The MS/MS spectra were used to search against a spectral library for analyte confirmation.

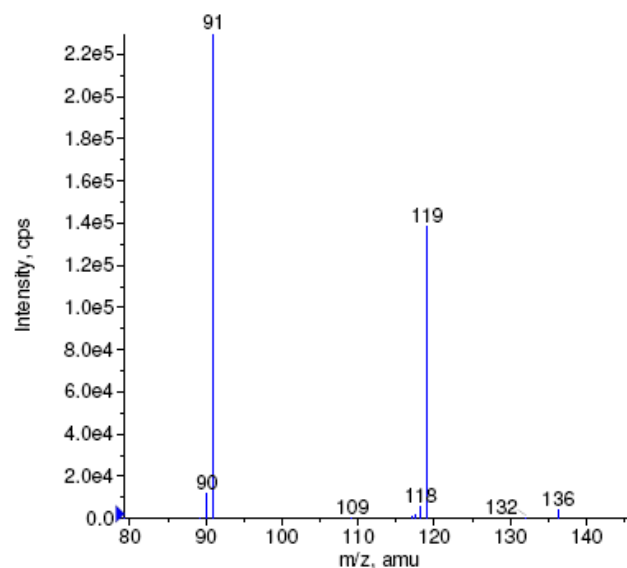


Compound Name: Amphetamine  
 Synonyms:  
 Formula: C<sub>9</sub>H<sub>13</sub>N  
 CAS Number: 300-62-9  
 Molecular weight(amu): 135.0000  
 Compound Class: illegal drug  
 Compound ID: A022  
 User Value: 0.0000  
 Keyword:  
 Internal Standard: No  
 Comment1:

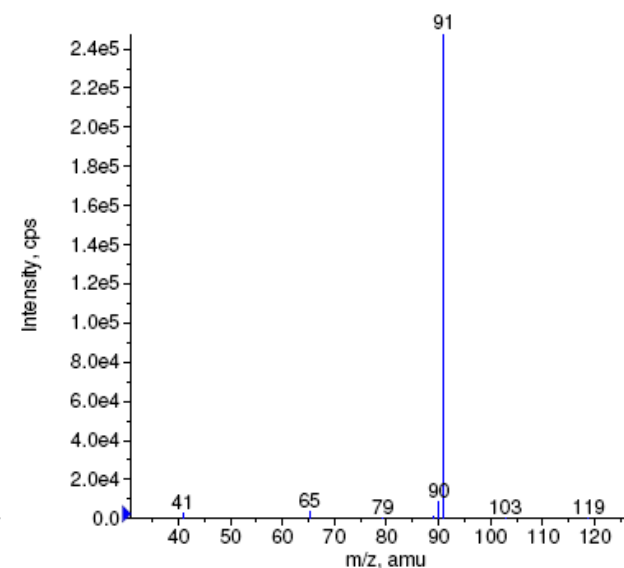
© Copyright 2004 by W. Weinmann

Instrument Model: API 365  
 Ion Source: Turbo Spray  
 Polarity: Positive  
 Scan Type: MS2  
 1st Precursor m/z: 136.0000  
 Collision Energy1(V): 20.0000  
 Charge State1: 1  
 CAD Gas Type: N<sub>2</sub>  
 CAD Gas Value: 4.0000  
 Retention Time(min): 0.20  
 Resolution: unit  
 Comment2:  
 Comment3:

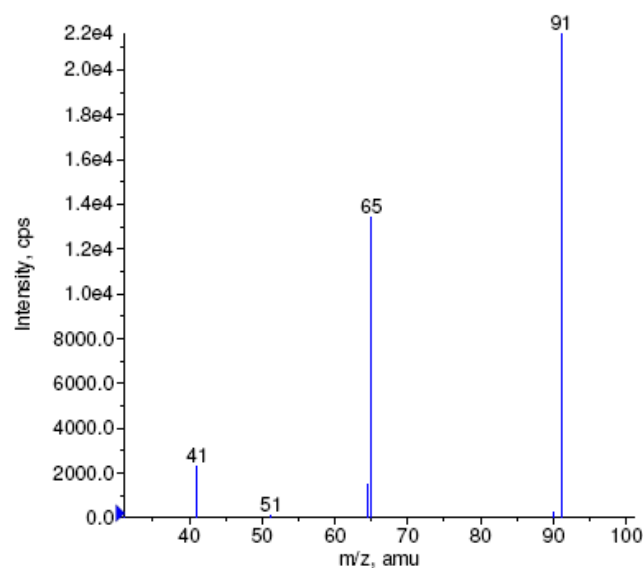
+ Amphetamine Prec(136.0) CE(20.0) RT(0.2)



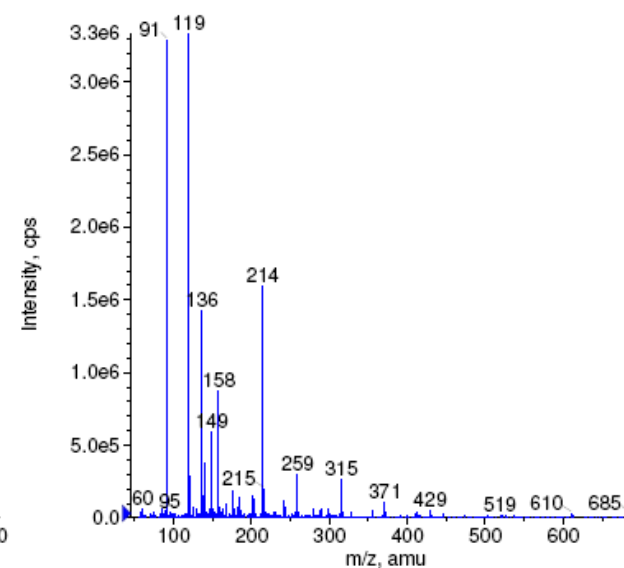
+ Amphetamine Prec(136.0) CE(35.0) RT(0.2)



+ Amphetamine Prec(136.0) CE(50.0) RT(0.3)



+ Amphetamine Prec(0.0) CE(0.0) RT(0.3)



Weinmann, W., PDF-Files of ESI-MS/MS-Product Ion Spectra (800 compounds)

Instrumentation: AB/Sciex API 365

<http://www.chemicalsoft.de/msmslib2005/MS2-2005-index.html>

## Efficiency of the different toxicological test measures...

LXQ 30 Minute	LXQ 13 Minute	LC-UV	Immunoassay
Nortriptyline	Nortriptyline	Nortriptyline	Barbiturates
Amitriptyline	Amitriptyline	Amitriptyline	Benzodiazepines
Benzoylecgonine	Benzoylecgonine	Benzoylecgonine	Cocaine
Cocaine	Cocaine	Cocaine	Opiates
Cocaethylene	—	Cocaethylene	THC
Cyclobenzaprine	—	—	—
Norbenzoylecgonine	Norbenzoylecgonine	—	—
Morphine	Morphine	—	—
Norcocaine	Norcocaine	—	—
Codeine	—	—	—
Norcocaethylene	Norcocaethylene	—	—
Methadone	—	—	—
Quinidine/Quinine	Quinidine/Quinine	—	—
Hydroxyzine	Hydroxyzine	—	—
Noskapine	Noskapine	—	—
Diltiazem	Diltiazem	—	—
Morphine-3-beta-Glucuronide	Morphine-3-beta-Glucuronide	—	—

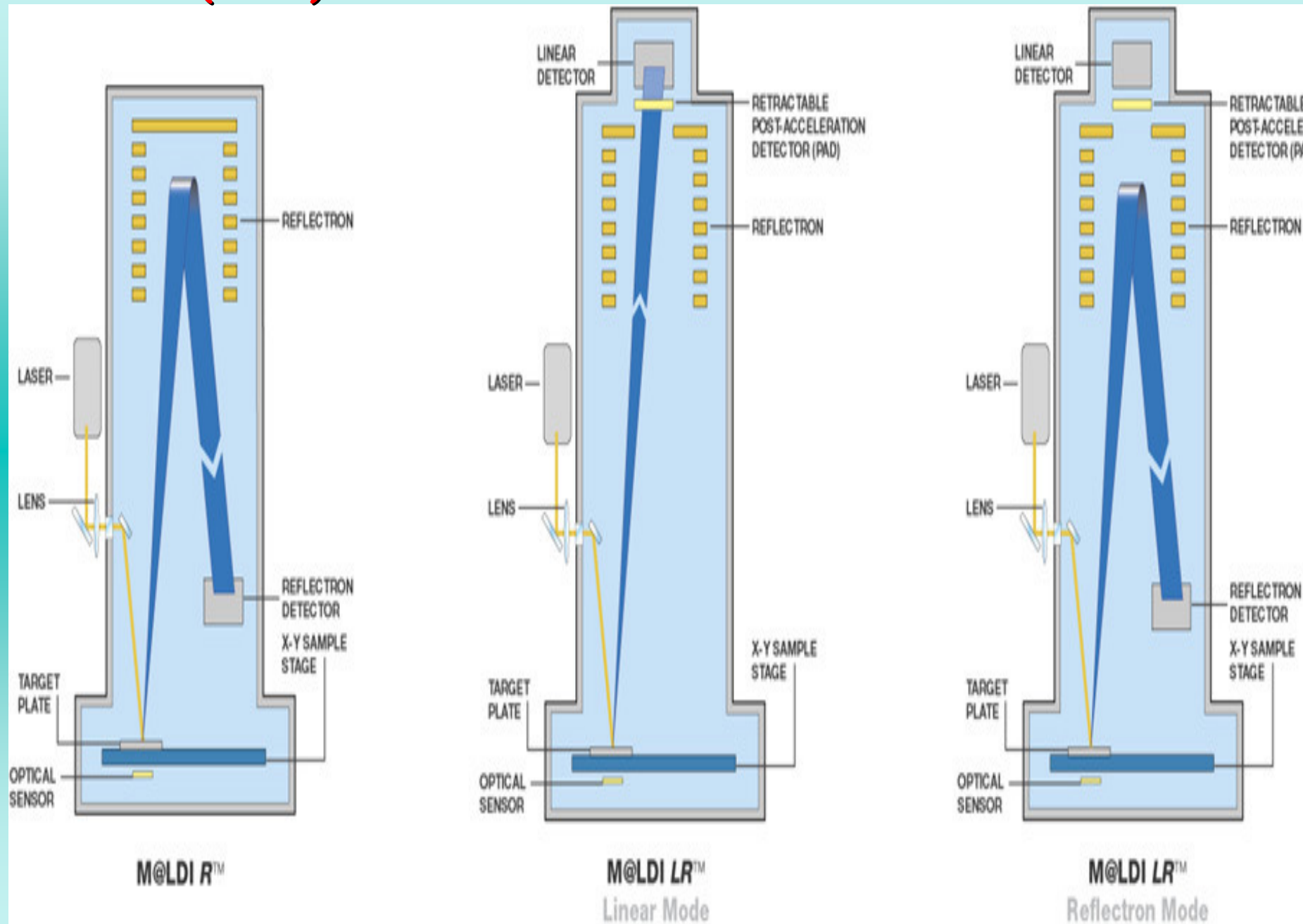
**Exact Mass Determination...**

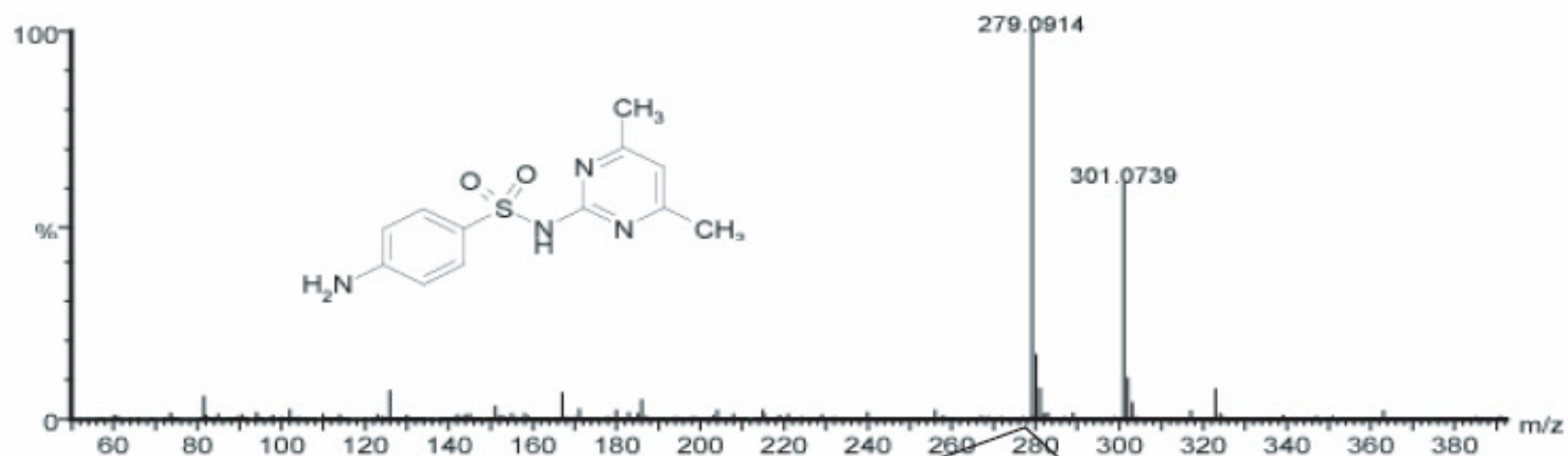


## Effect of *Mass Accuracy* and *Mass Tolerance* on (Peptide) Mass Fingerprinting Search Results

Search $m/z$	Mass tolerance (Da)	# Hits
1529	1	478
1529.7	0.1	164
1529.73	0.01	25
1529.734	0.001	4
1529.7348	0.0001	2

# MALDI - (TOF) instruments





5ppm tolerance

50ppm tolerance

Elemental Composition

File Edit View Process Help

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0

Isotope cluster parameters: Separation = 1.0, Abundance = 1.00%

Monoisotopic Mass, Odd and Even Electron Ions

327 formula(e) evaluated with 3 results within limits (up to 50 closest results for each mass)

Mass	Calc. Mass	mDa	PPM	DBE	Formula	Score	C	H	N	O	S
279.0914	279.0916	-0.2	-0.6	7.5	C12 H15 N4 O2 S	1	12	15	4	2	1
	279.0522	-0.8	-3.0	16.5	C20 H11 N2	3	20	11	2		
	279.0602	1.2	4.2	2.5	C11 H19 O6 S	2	11	19		6	1

For Help, press F1

Elemental Composition

File Edit View Process Help

Single Mass Analysis

Tolerance = 50.0 PPM / DBE: min = -1.5, max = 50.0

Isotope cluster parameters: Separation = 1.0, Abundance = 1.00%

Monoisotopic Mass, Odd and Even Electron Ions

327 formula(e) evaluated with 30 results within limits (up to 50 closest results for each mass)

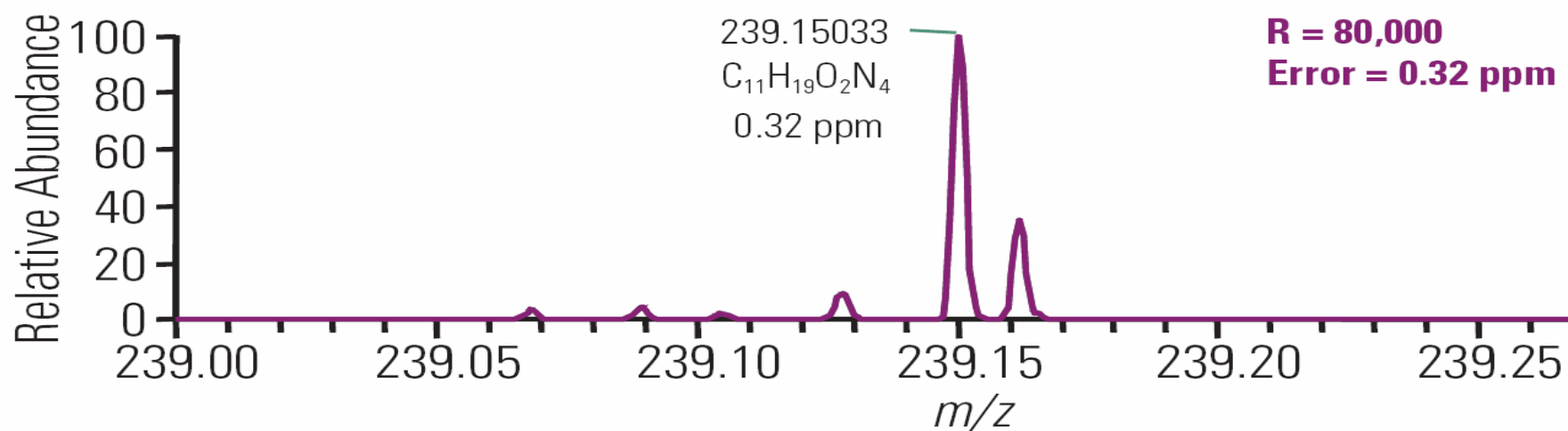
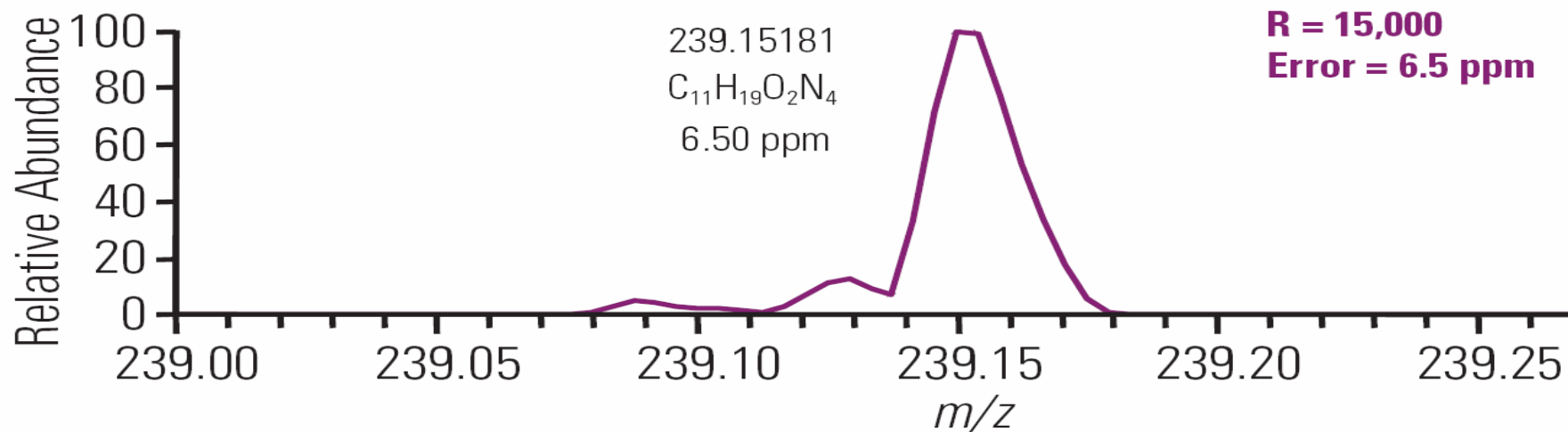
Mass	Calc. Mass	mDa	PPM	DBE	Formula	Score	C	H	N	O	S
279.0914	279.0916	-0.2	-0.6	7.5	C12 H15 N4 O2 S	4	12	15	4	2	1
	279.0932	0.8	3.0	16.5	C20 H11 N2	28	20	11	2		
	279.0902	-1.2	-4.2	2.5	C11 H19 O6 S	7	11	19		6	1
	279.0929	-1.5	-5.4	7.0	C14 H17 R O3 S	2	14	17		3	1
	279.0895	-1.9	-6.7	12.0	C7 H13 N O3	20	7	13	1	3	
	279.0869	-2.5	-9.0	3.0	C9 H17 N3 O5 S	11	9	17	3	5	1
	279.0862	-3.2	-11.5	12.5	C15 H11 N4 O2	19	15	11	4	2	
	279.0876	-3.0	-13.0	9.5	C7 H15 N6 O4 S	10	7	15	6	4	1
	279.0886	-4.2	-15.0	11.5	C17 H15 N2 S	12	17	15	2		1
	279.0889	-4.5	-16.3	7.5	C14 H15 O6	15	14	15		6	
	279.0988	5.4	19.2	8.0	C11 H13 N5 O4	22	11	13	5	4	
	279.0855	-5.9	-21.1	8.0	C12 H13 N3 O5	18	12	13	3	5	
	279.0961	-6.7	-24.0	7.5	C13 H15 N2 O5	17	13	15	2	5	
	279.0844	-7.0	-25.2	11.5	C18 H15 O S	14	18	15		1	1

For Help, press F1

Influence of exact mass resolution on identification power

Compound	Actual Mass	Measured Mass	mDa Error	ppm Error
4-acetamidophenol	152.0711	152.0707	-0.4	-2.6
sulfaguanidine	215.0602	215.0597	-0.5	-2.3
Sulfadimethoxine	311.0814	311.081	-0.4	-1.3
Val-Tyr-Val	380.2185	380.219	0.5	1.3
Terfenadine	472.3215	472.321	-0.5	-1.1
Leucine enkephalin	556.2771	556.2775	0.4	0.7
Reserpine	609.2812	609.2828	1.6	2.6
Erythromycin	734.469	734.4695	0.5	0.7
RMS ppm error				1.8

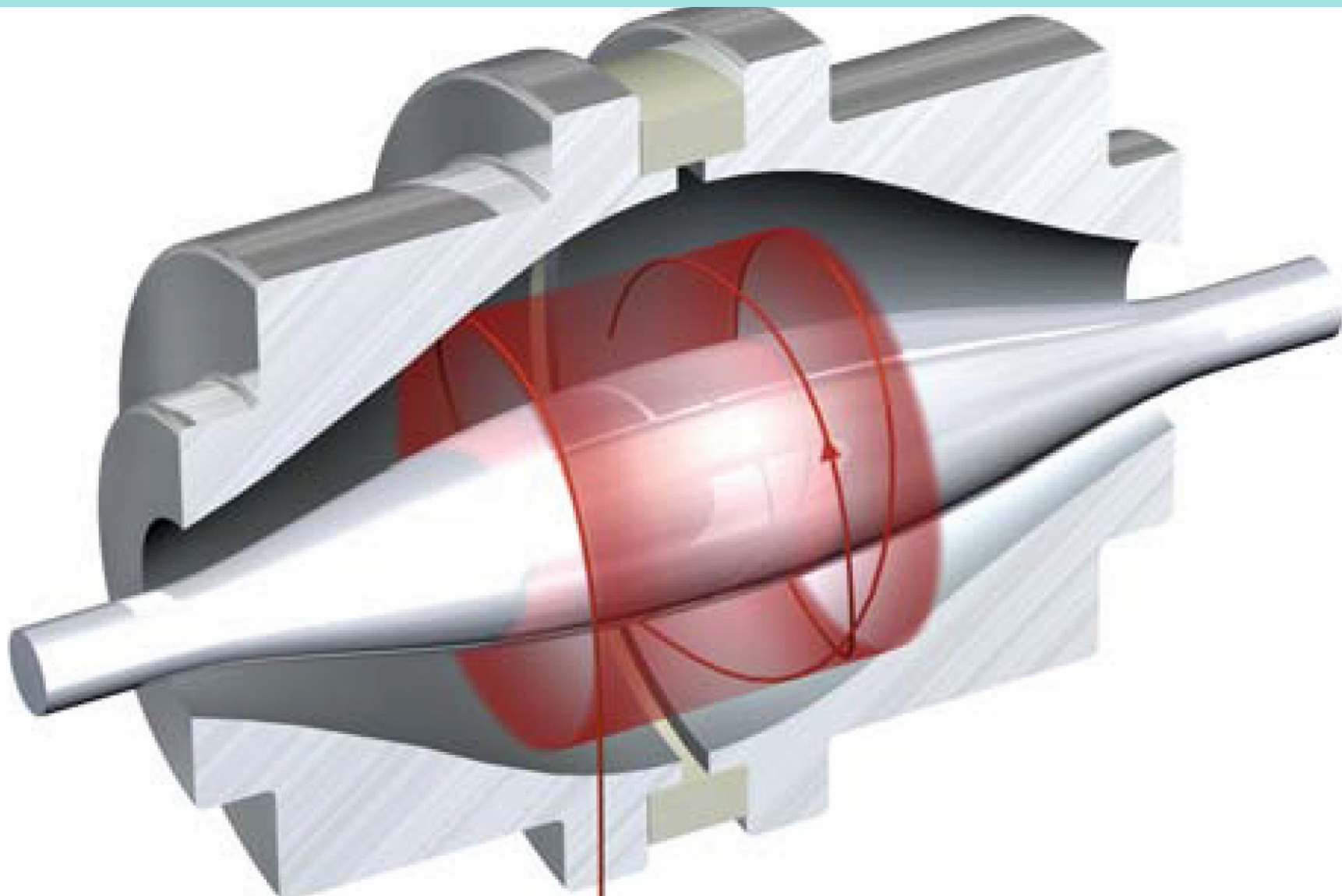
**Identification of drugs by exact mass measurements**



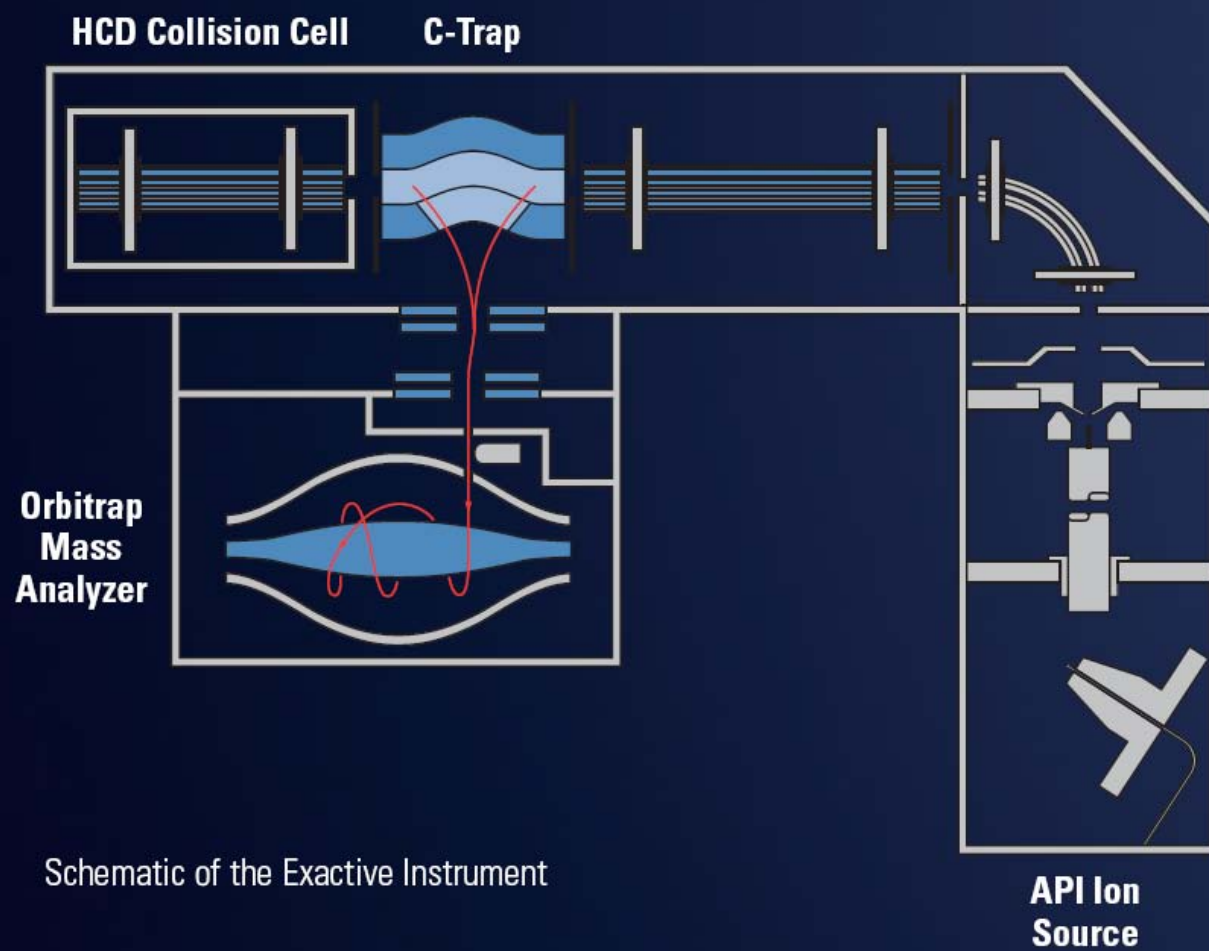
The unusually high error of 6.5 ppm (top) indicates the need for higher resolution. Re-analysis performed at 80,000 (bottom) showing baseline separation of the targeted pesticide from the matrix interference



## *The Orbitrap - Technology*







But there are softwaretools needed...

## ToxID Summary Report

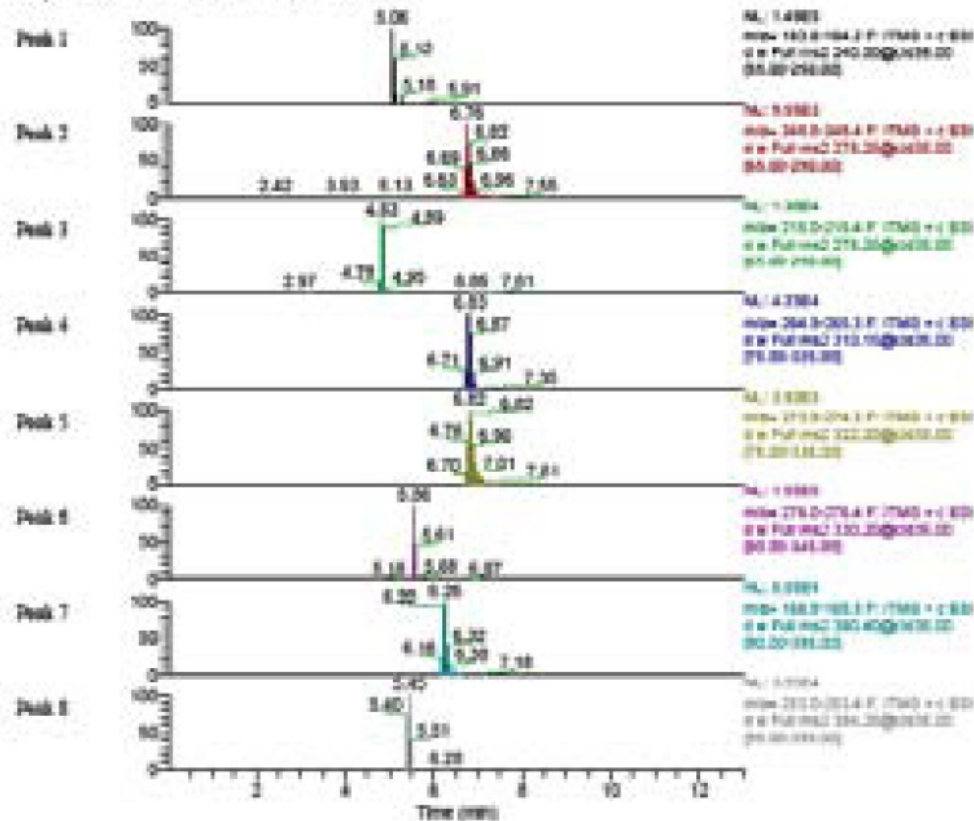
Raw File Name: C:\Documents and Settings\user\Local Desktop\Application\_Names\ToxID\10 RAW

Config File Name: C:\Oxlab\examples\ToxID\ToxID\_config\_10min.cnv

Sample Name:

Laboratory: ChemLab

Acquisition Start Time: 3/13/2007 1:04:34 AM



Identified compounds  
MS/MS scans related  
chromatograms

Peak Number	Compound Name	Code	SI	RSI	m/z	Expected RT	Real RT	Intensity	Library Name
1	Bupropion	p	000	000	140.0	5.30	5.06	148731	Tox_Library
2	EDDP	p	017	073	278.2	6.80	6.76	9540	Tox_Library
3	Veratrine	p	016	037	278.2	6.00	4.83	13044	Tox_Library
4	Methadone	p	032	032	100.2	6.70	6.83	42362	Tox_Library
5	Chlorpromazine-D3	i	030	030	323.2	6.80	6.83	2004	Tox_Library
6	Propoxy-D3	i	069	074	300.2	5.80	5.56	154827	Tox_Library
7	Haloperidol-D4	i	030	037	380.4	6.30	6.30	81580	Tox_Library
8	Quetiapine	p	070	071	384.2	5.40	5.40	38512	Tox_Library

Identified compounds  
with search indexes

## ToxID Long Report

Raw File Name: C:\Documents and Settings\marta\Local Desktop\Desktop\Application\_Video\_Test20111118.RAW

Config File Name: C:\Weather\examples\TwoIP\TwoIP.config\_1.htm.rcv

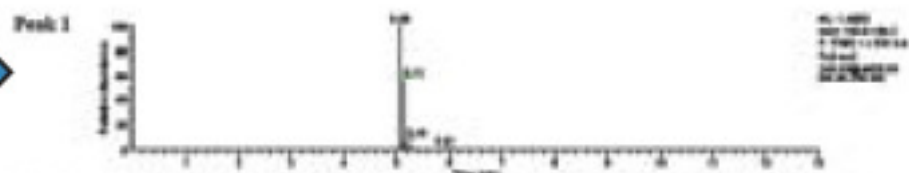
### Sample Number

Laboratory Chemicals

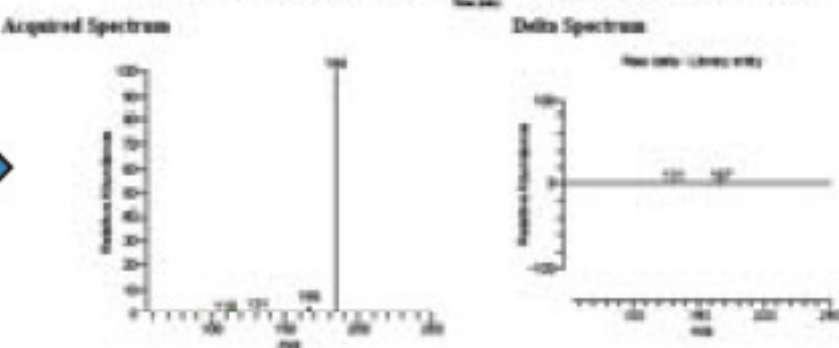
Acquisition Start Time: 1/13/2007 1:04:14 AM

Book Number	Component Name	Code	SI	BIT	ms	Expected RT	Real RT	Intensity	Library Name
1	Supragas	p	008	008	140.0	1.20	1.00	148721	Gas Library

**MS/MS scans  
chromatogram**

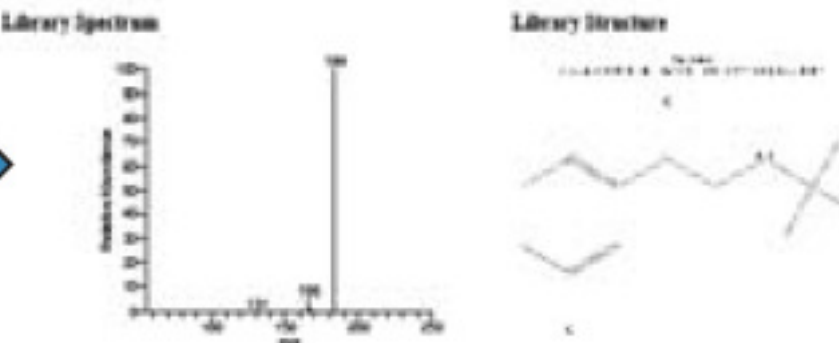


**Acquired spectrum  
for visual inspection**



**Delta spectrum shows the difference between the experimental spectrum and the library spectrum**

### Library spectrum for verification



### Compound structure

## Practical strategies in toxicological analysis....

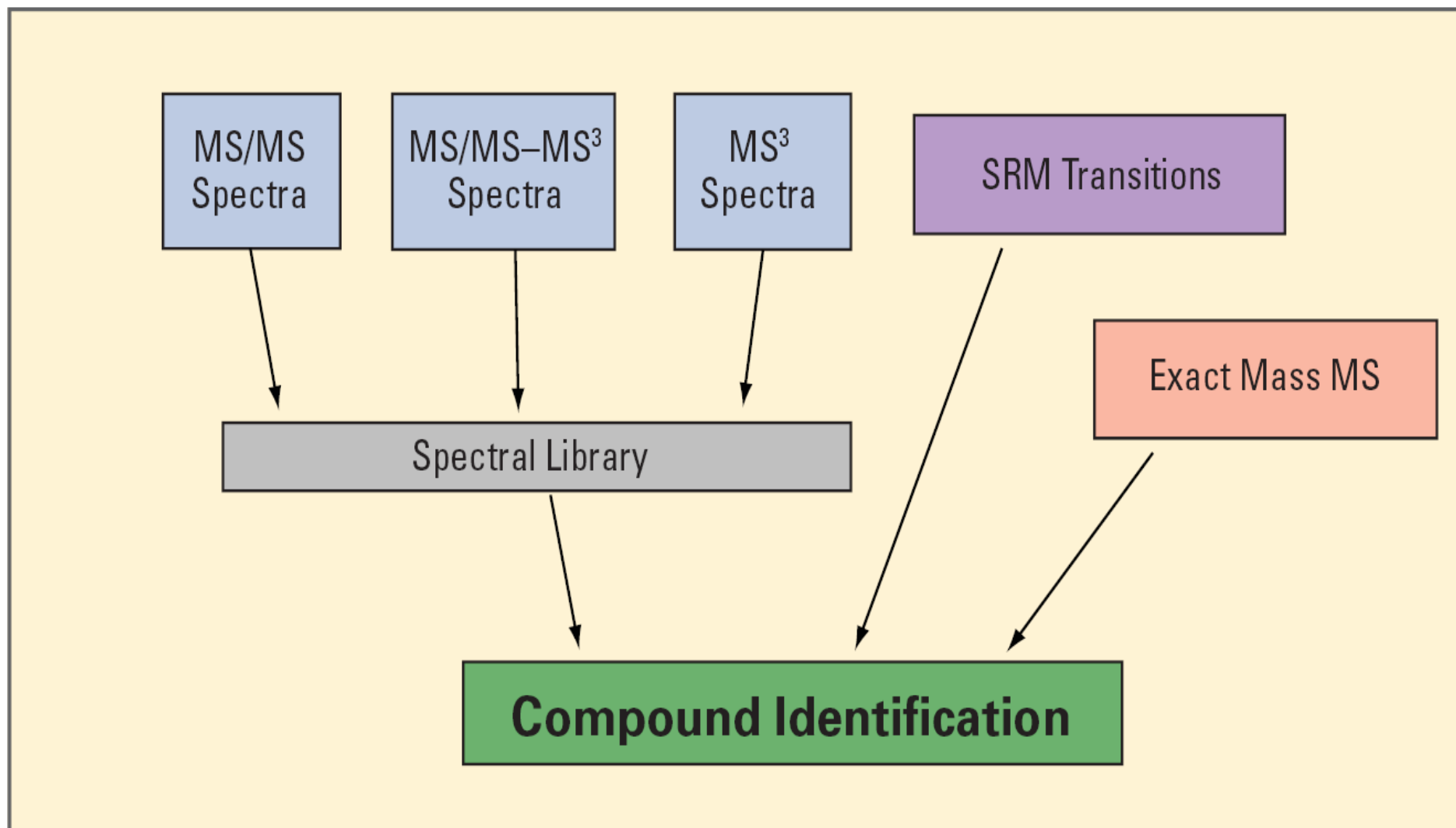
Step 1: Extract analytes from urine with SPE procedure

Step 2: Analyze the samples with LC-MS/MS method

Step 3: Automated library search and Reporting with ToxID



## Practical strategies in toxicological analysis.... (2)





Edited by W. R. Külpmann

WILEY-VCH

# Clinical Toxicological Analysis

Procedures, Results, Interpretation

Volume 1



**Danke  
für  
Ihre  
Aufmerksamkeit!**



