

Spectra Interpretation: Small Molecules

Josef Cvačka

General interpretation procedure for mass spectra

- 1/ Disclosing ions that are related to the analyte and that are not
- 2/ Determination of the molecular weight – looking for molecular ions M^+ , adducts $[M+H]^+$, $[M+Na]^+$, $[M+Cl]^-$, deprotonated molecules $[MH]^-$, dimers and multiply charged ions
- 3/ Estimating elements which are present: inspecting isotope clusters, application of nitrogen rule
- 4/ Determination of the elemental formula from exact mass measurement
- 5/ Comparison the spectrum with a library, searching for at least a similar spectrum
- 6/ Solving fragmentation spectra (requires knowledge of fragmentation mechanisms and empirical rules)

1.

Ions, which are not related to the analyte

Background ions, contaminants

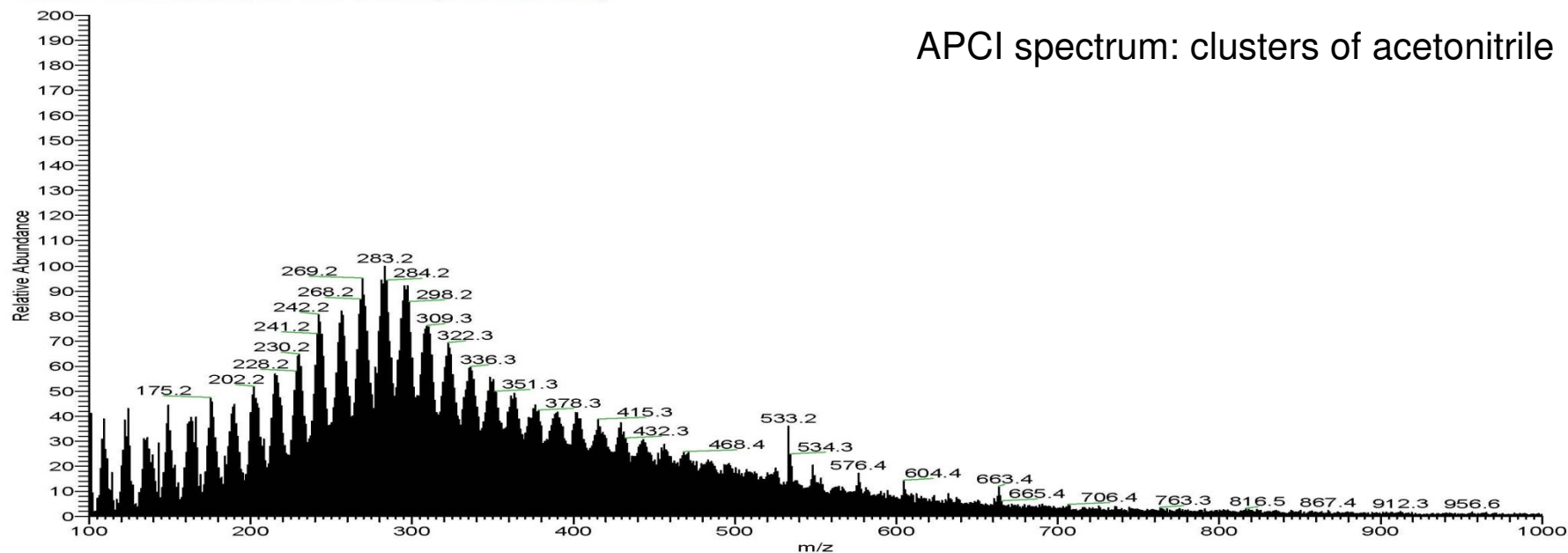
- solvent cluster ions
- impurities from solvents
- impurities from plastic ware and glassware
- impurities in the mass spectrometer from previous injections
- column bleeding
- impurities from sample

c:\Documents and Settings\...\0962VV01

3/10/2009 12:32:40 PM

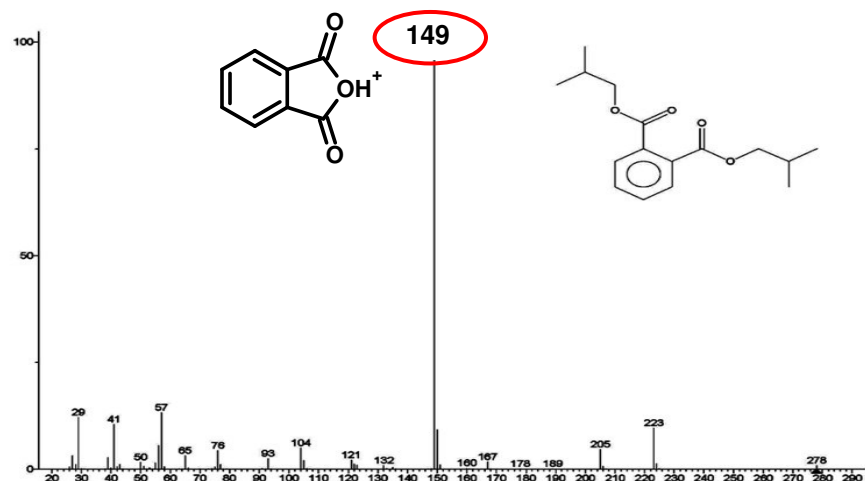
Bee Wax esters

0962VV01 #7-262 RT: 0.08-3.33 AV: 256 NL: 3.63E1
F: ITMS + c APCI corona sid=8.00 Full ms [100.00-1000.00]

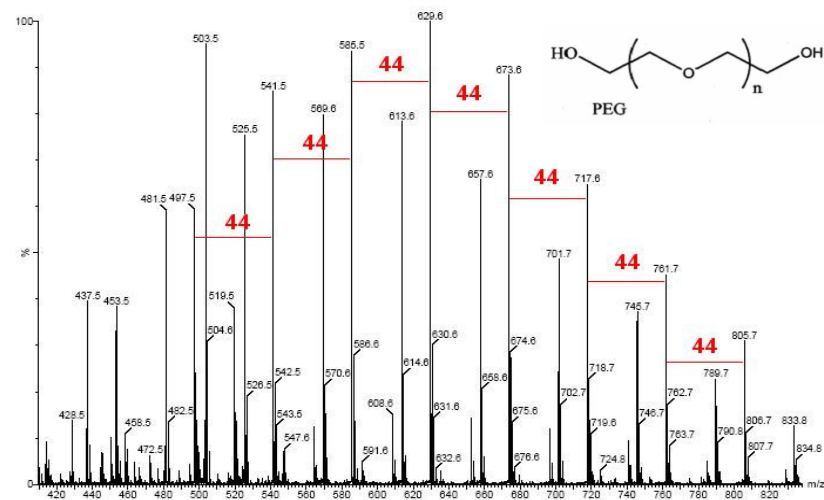


Background ions, contaminants

Phthalates: common plasticizers,
from laboratory plastic
 m/z 149, 279, 301, 391, 413 ...

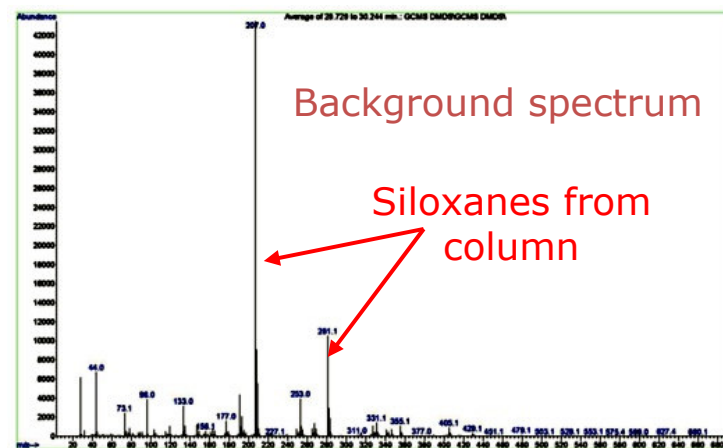
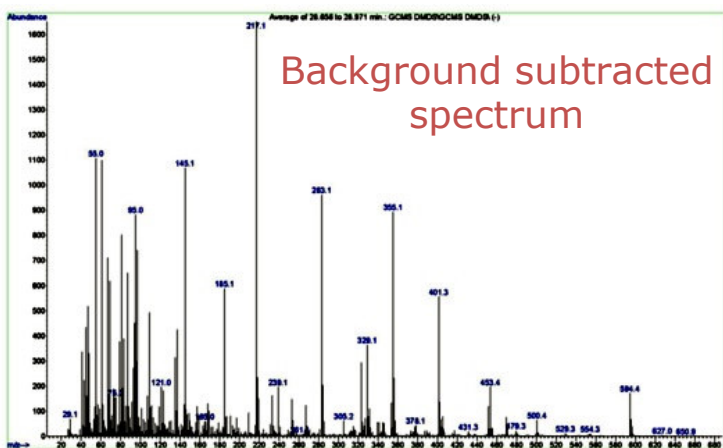
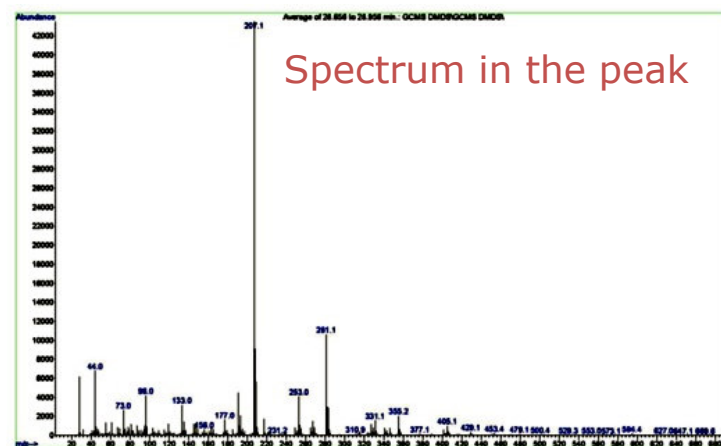
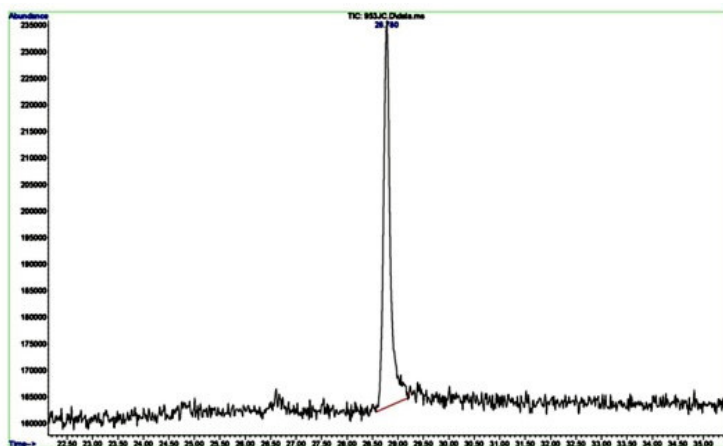


Polyethyleneglycols:
from laboratory plastic, gloves,
skin lotion
peak difference 44 u



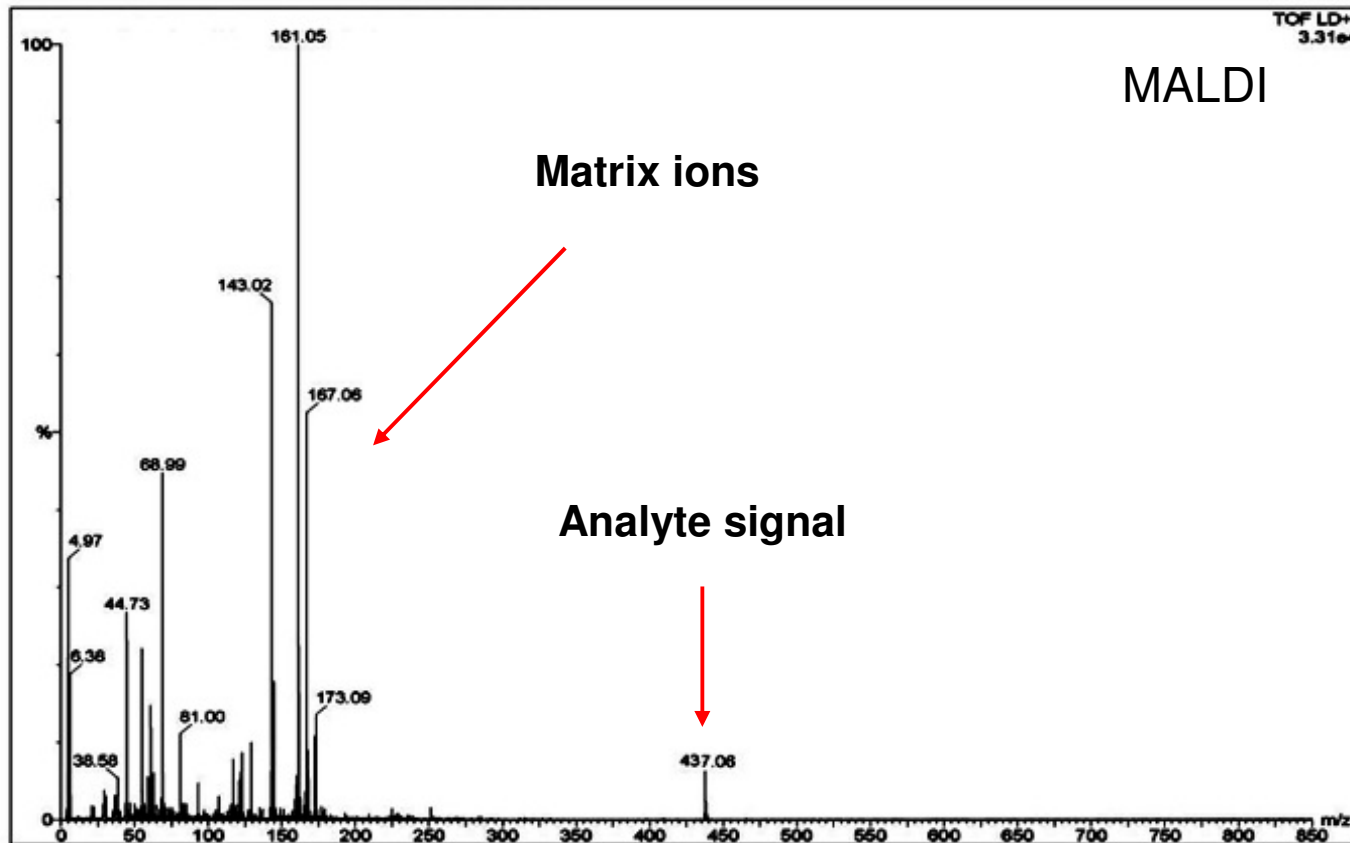
MeOH wash of laboratory gloves

Background subtraction



Matrix ions in MALDI

- Clusters, fragments, adducts of the matrix ions in the low mass range; very intense
- MALDI spectra are usually recorded starting from $\sim m/z$ 500



2.

Determination of the molecular weight

Determination of molecular weight

Identification of molecular ion or molecular adduct

I. The electron ionization



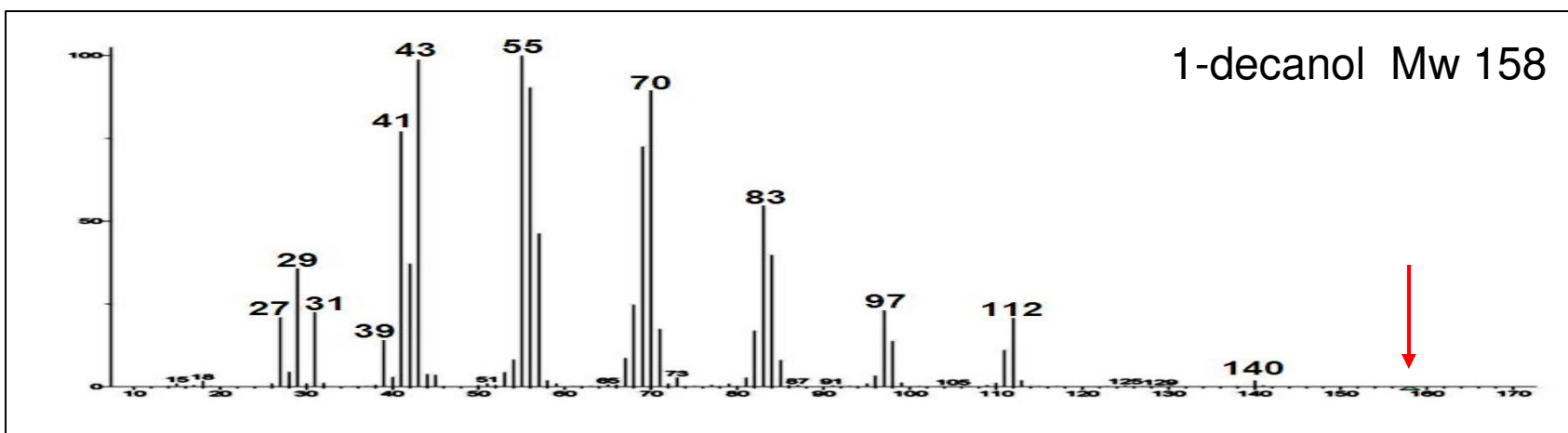
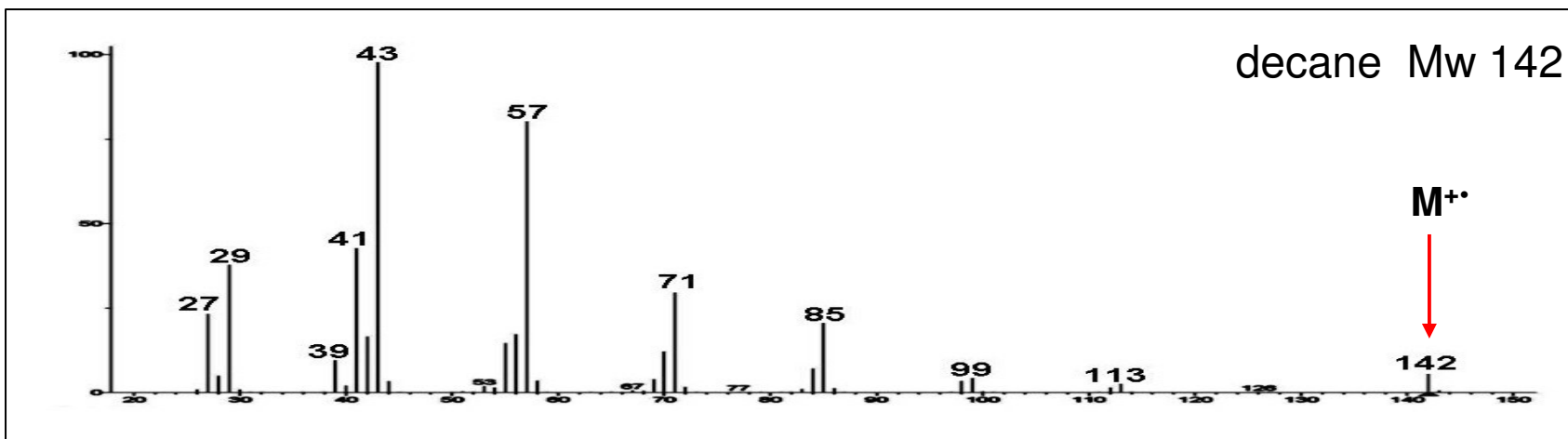
Molecular ion ($M^{+\cdot}$) is a radical cation. The m/z corresponds to the monoisotopic mass of analyte.

Identification of the molecular ion in the spectrum

- 1/ the molecular ion may not be present
 - 2/ if present, it must have the highest m/z value
 - 3/ the molecular ion provides logical neutral losses
-

Determination of molecular weight

I. The electron ionization

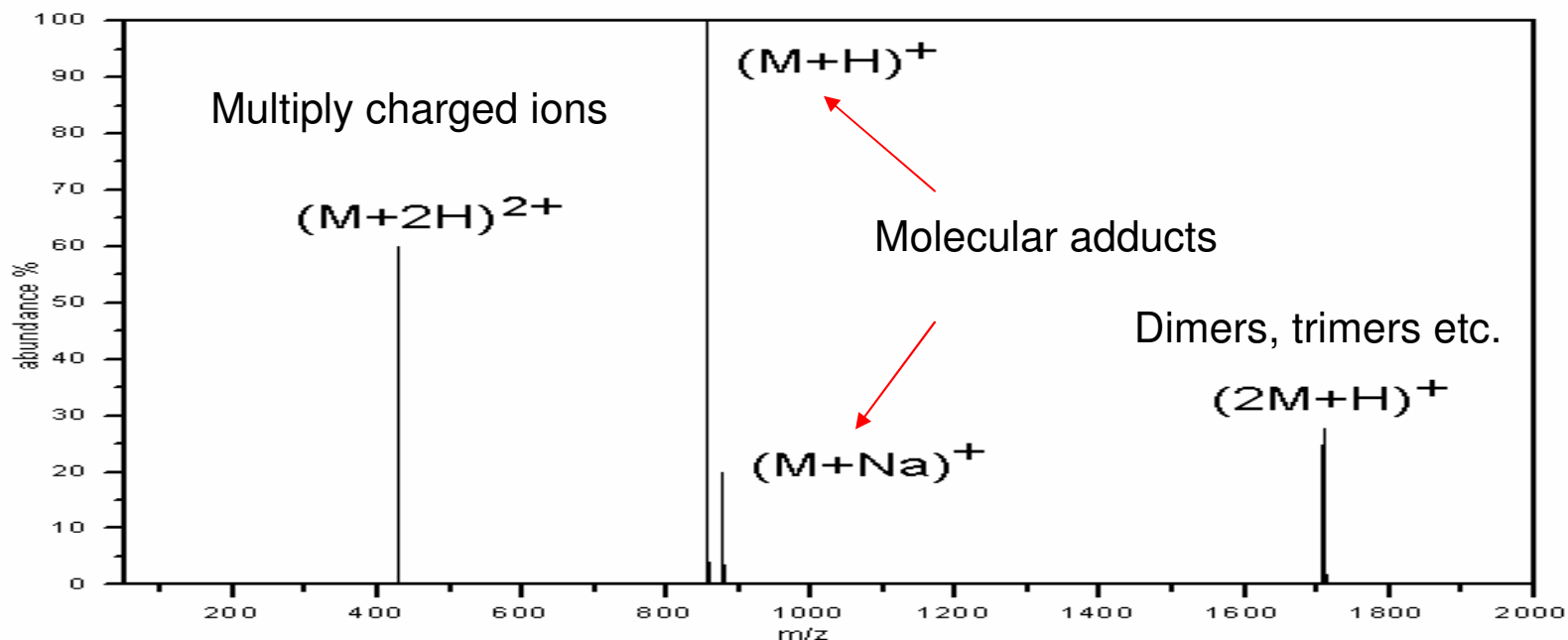


Determination of molecular weight

II. Soft ionization techniques (ESI, APCI, MALDI)

Molecular adducts ($[M+H]^+$, $[M+Na]^+$) or deprotonated molecules ($[M-H]^-$)

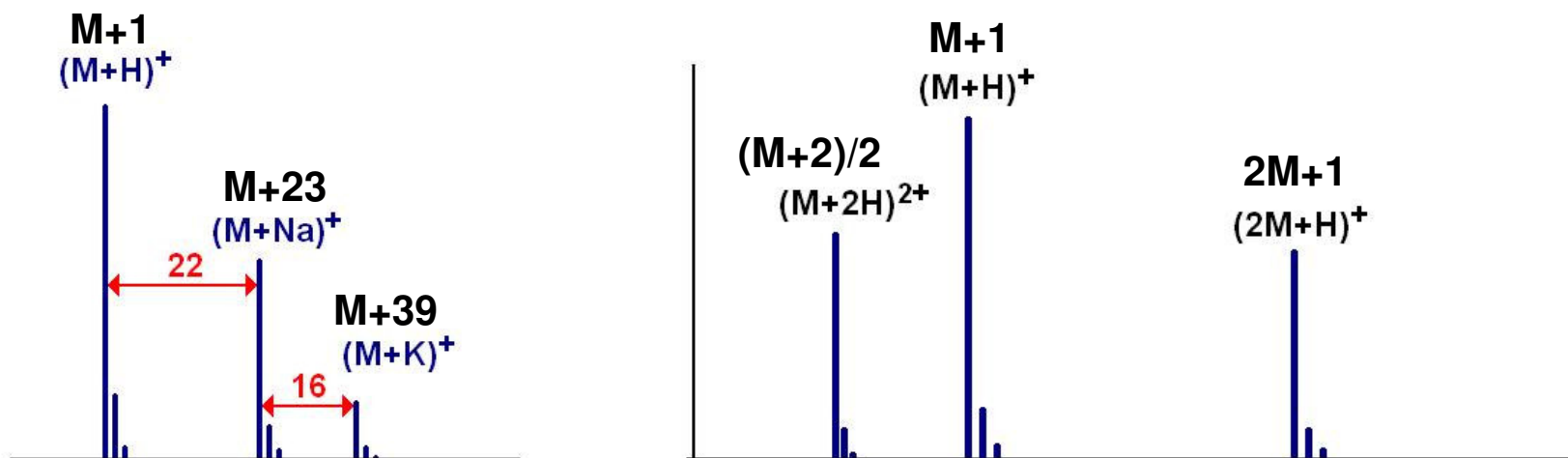
The molecular adduct is an ion with even number of electrons and may not be the highest ion in the spectrum.



Determination of molecular weight

II. Soft ionization techniques (ESI, APCI, MALDI)

The molecular weight is determined based on the presence of adducts, dimers or multiply charged ions.

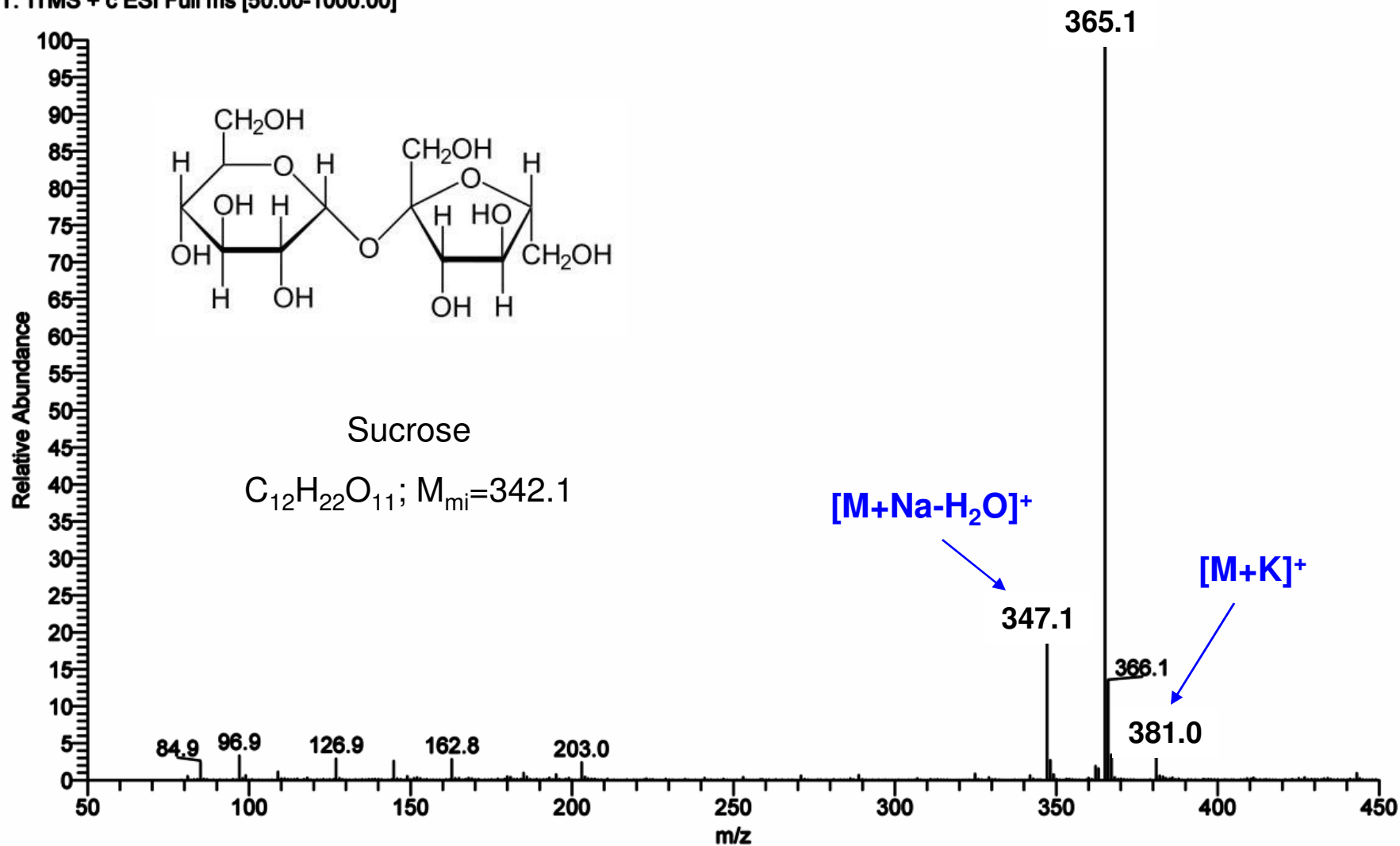


Calculation of adducts, dimers or multiply charged ions: software EIC
I:\MISC\MS\DOWNLOAD\

Determination of molecular weight



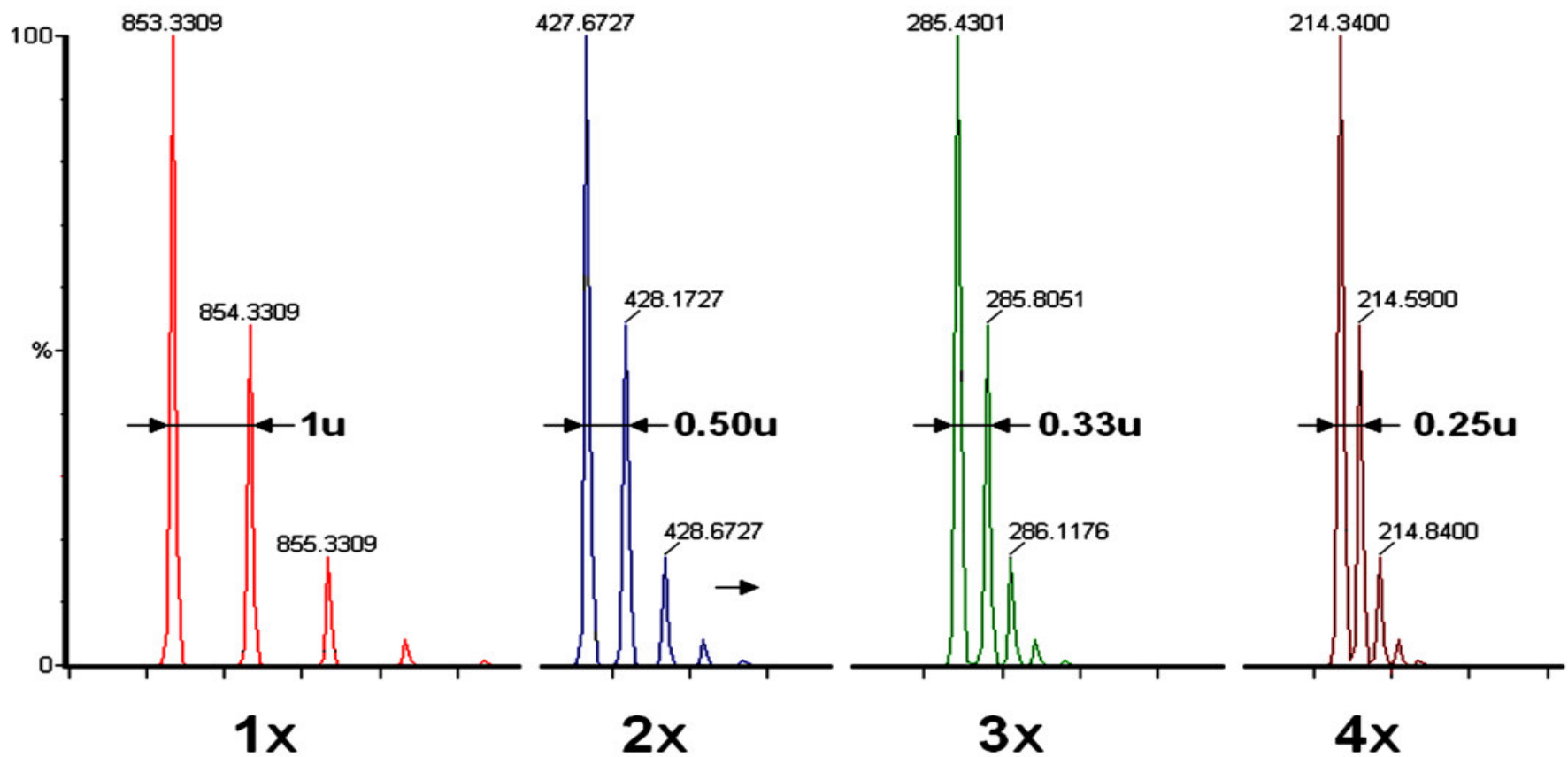
2256jc #1-51 RT: 0.00-0.75 AV: 51 NL: 1.77E4
T: ITMS + c ESI Full ms [50.00-1000.00]



Determination of molecular weight – charge state

Determining number of charges

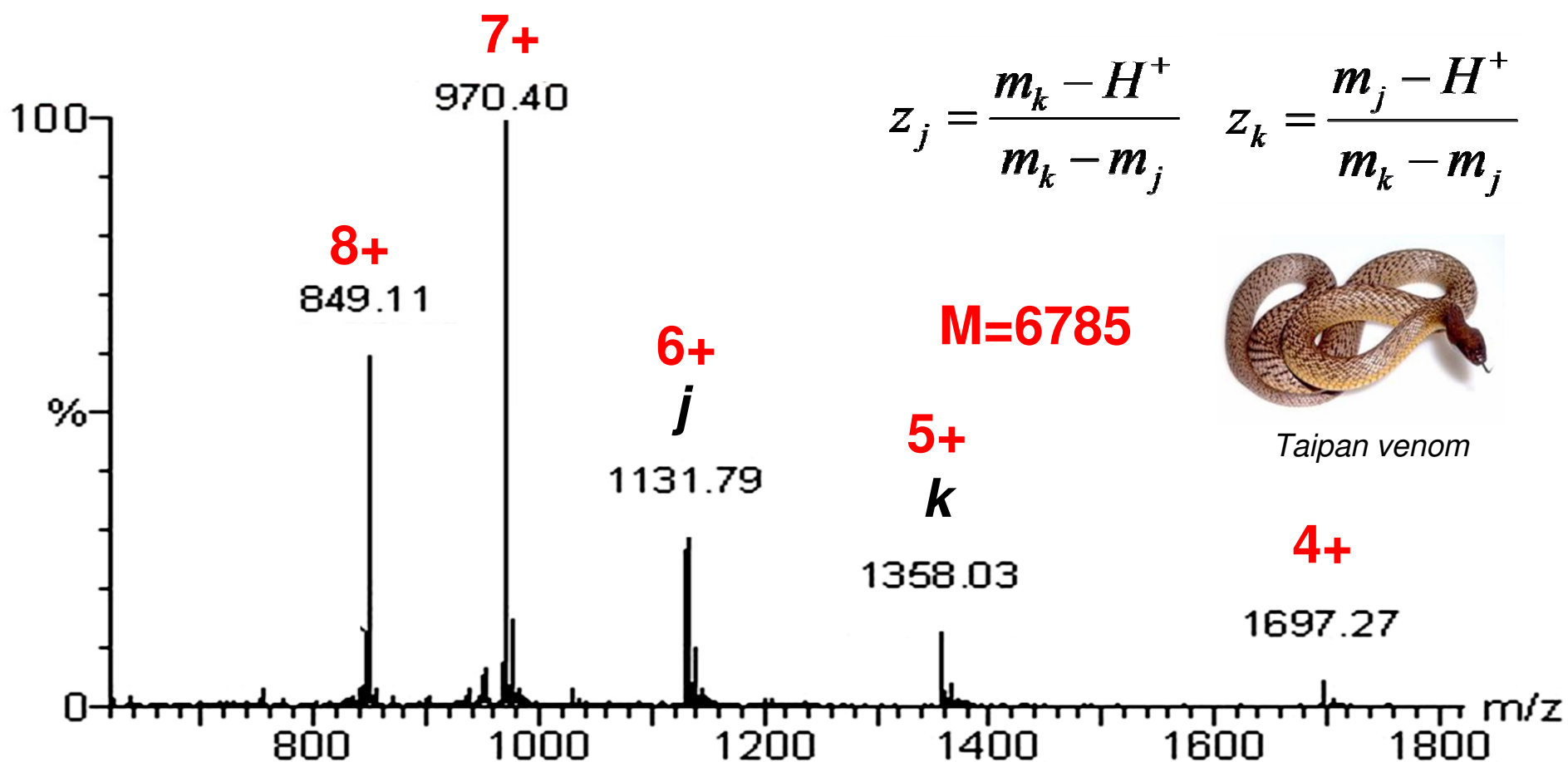
Number of charges can be determined from the distance between the peaks in the isotopic clusters.



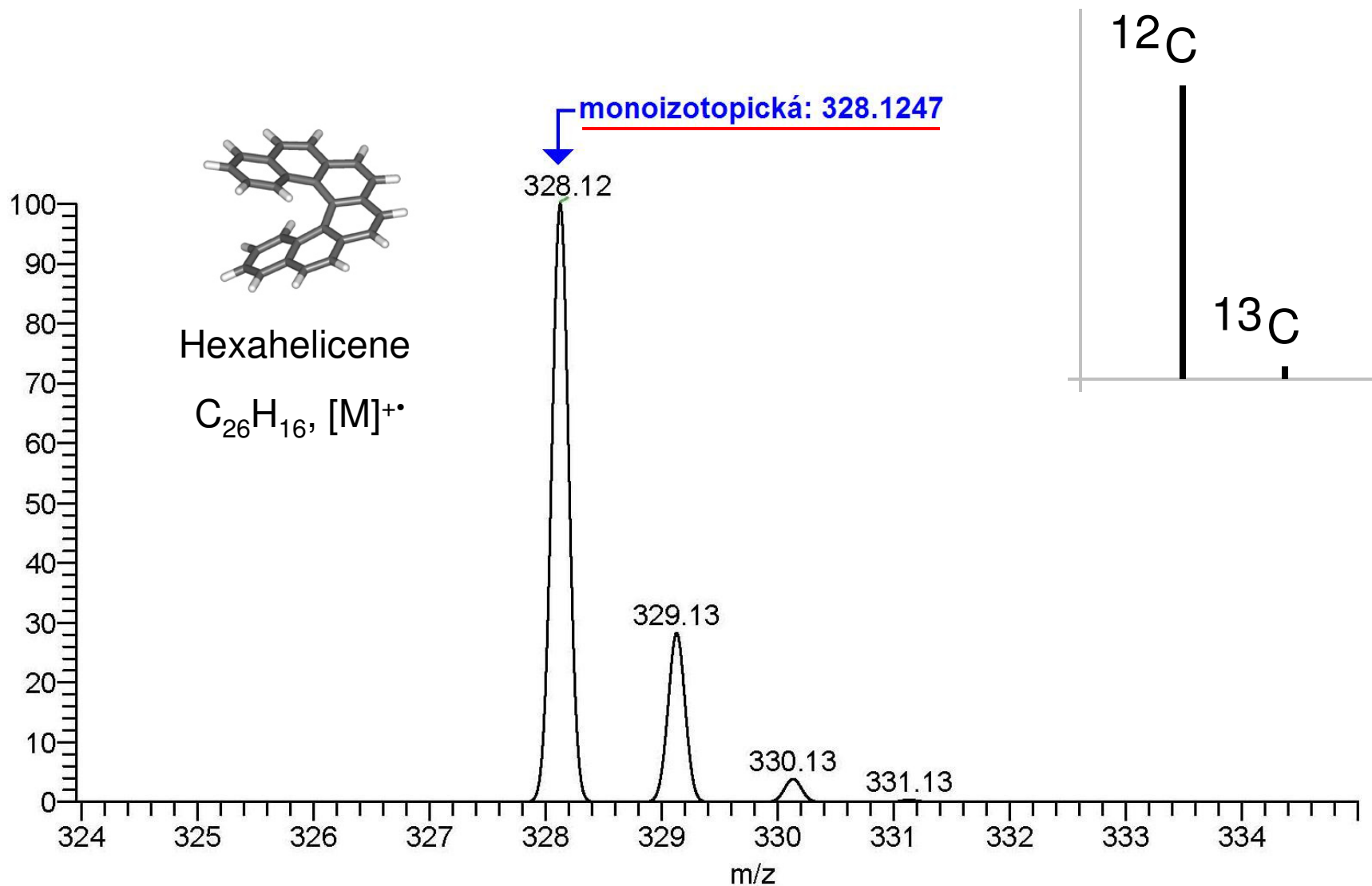
Determination of molecular weight – charge state

Determining number of charges

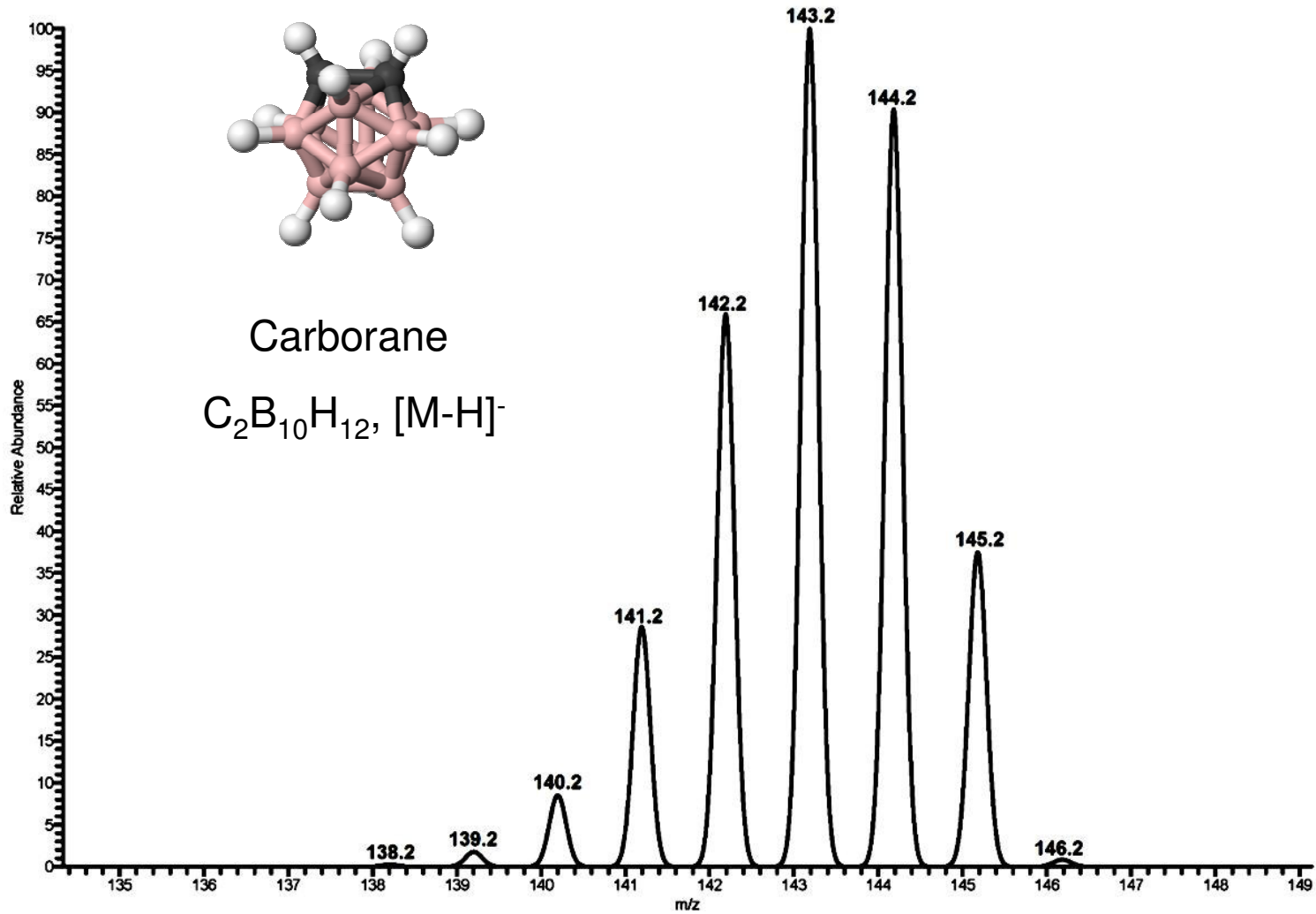
Number of charges can be determined from the distance between the neighboring peaks representing different charge states.



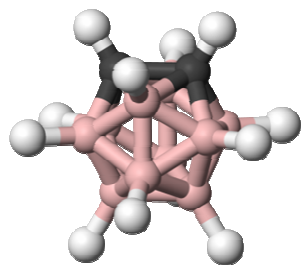
Monoisotopic mass



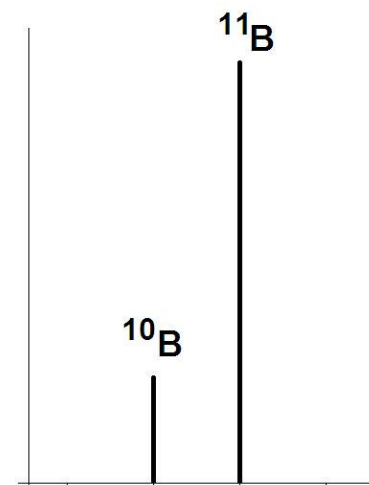
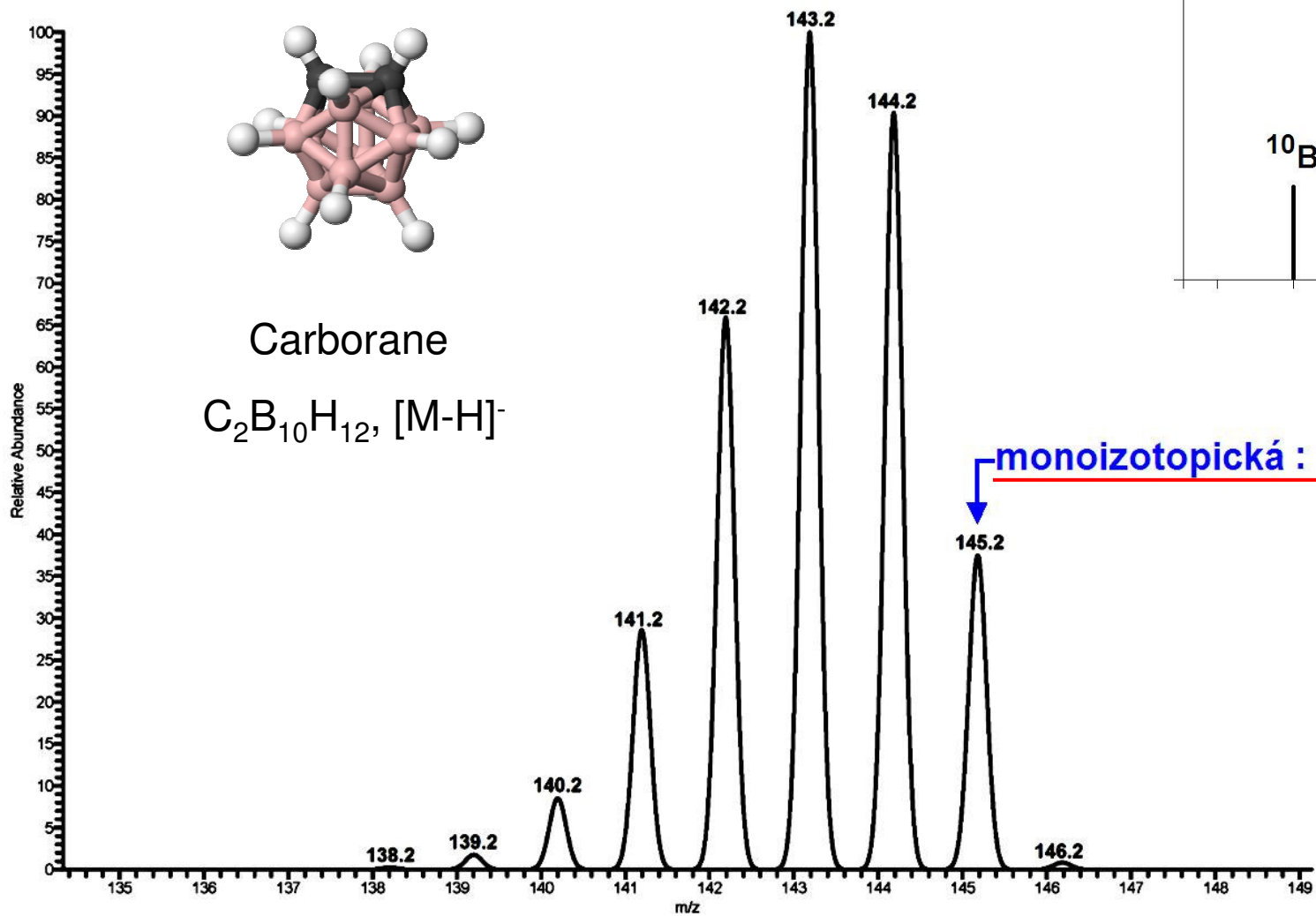
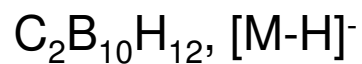
Monoisotopic mass



Monoisotopic mass



Carborane



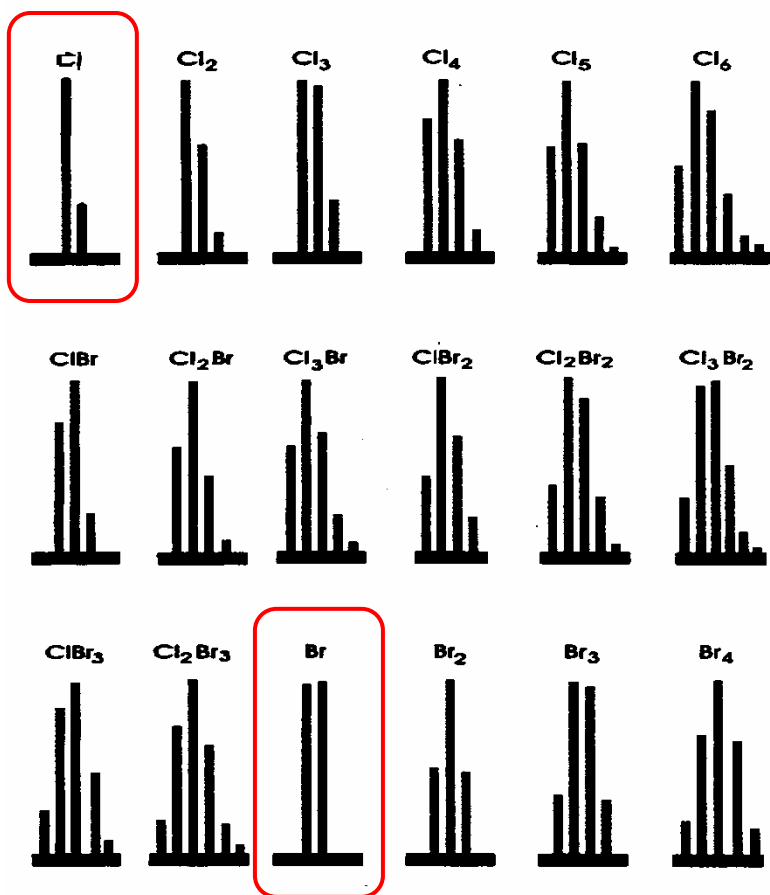
monoizotopická : 145.1791

3.

Elements present in a ion

Isotope clusters

Isotopic clusters indicate the presence of some elements in an ion (e.g., Cl, Br, metals etc.).



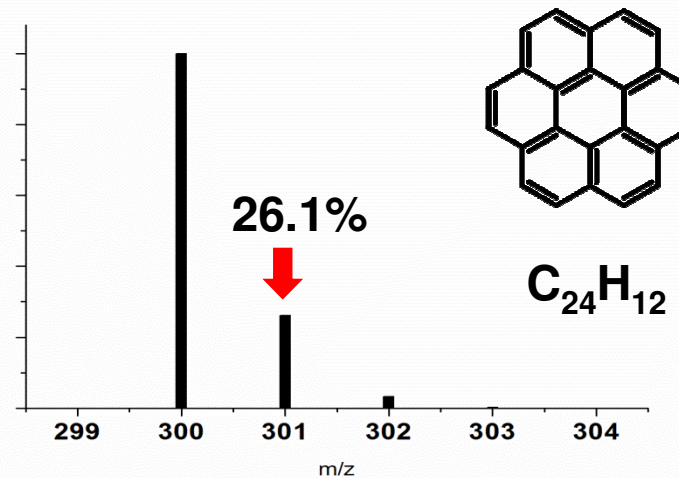
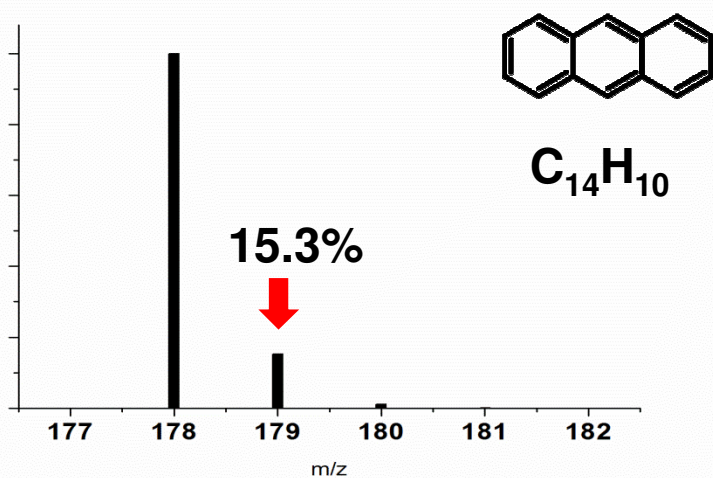
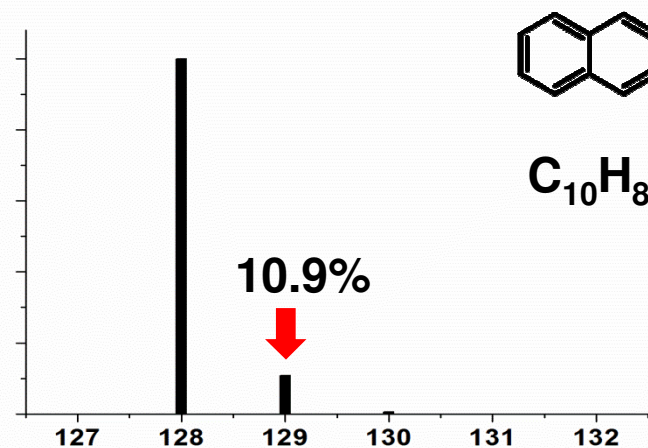
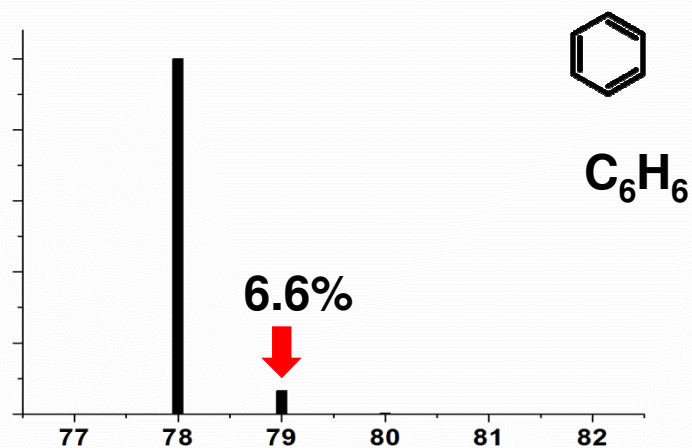
Computer programs allow you to calculate the composition of the cluster from the specified summary formula and compare it with experiment.



<http://www.colby.edu/chemistry/NMR/IsoClus.html>

Number of carbon atoms

The number of carbons in an ion can be estimated based on the intensity of ^{13}C isotope (relative ratio $^{13}\text{C}/^{12}\text{C}$ is $\sim 1.1\%$)



Nitrogen rule

- Elements with odd nominal masses form odd numbers of covalent bonds.
- Elements with even masses form even numbers of covalent bonds, with the exception of nitrogen (nominal mass of 14, valency of 3).

Nitrogen rule applies to organic compounds containing C, H, N, O, S, P, F, Cl, Br, I

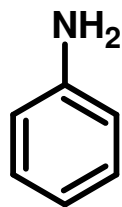
Odd value of molecular weight = odd number of nitrogens

Even value of molecular weight = even (zero) number of nitrogens

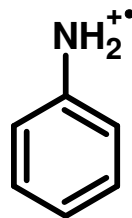
Applying the rule for ions

EI – valid for $M^{+\bullet}$ as stated above

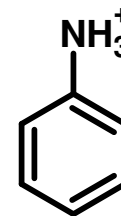
ESI, APCI, MALDI – the rule must be reversed for molecular adducts!



M=93



m/z 93



m/z 94

4.

**Determination of elemental composition from
accurate mass**

Elemental composition from accurate mass

The more accurately we determine mass of the ion, the more we restrict the number of possible structures.

Example. paclitaxel,
 $C_{47}H_{51}NO_{14}$, mon. mass
854.3388

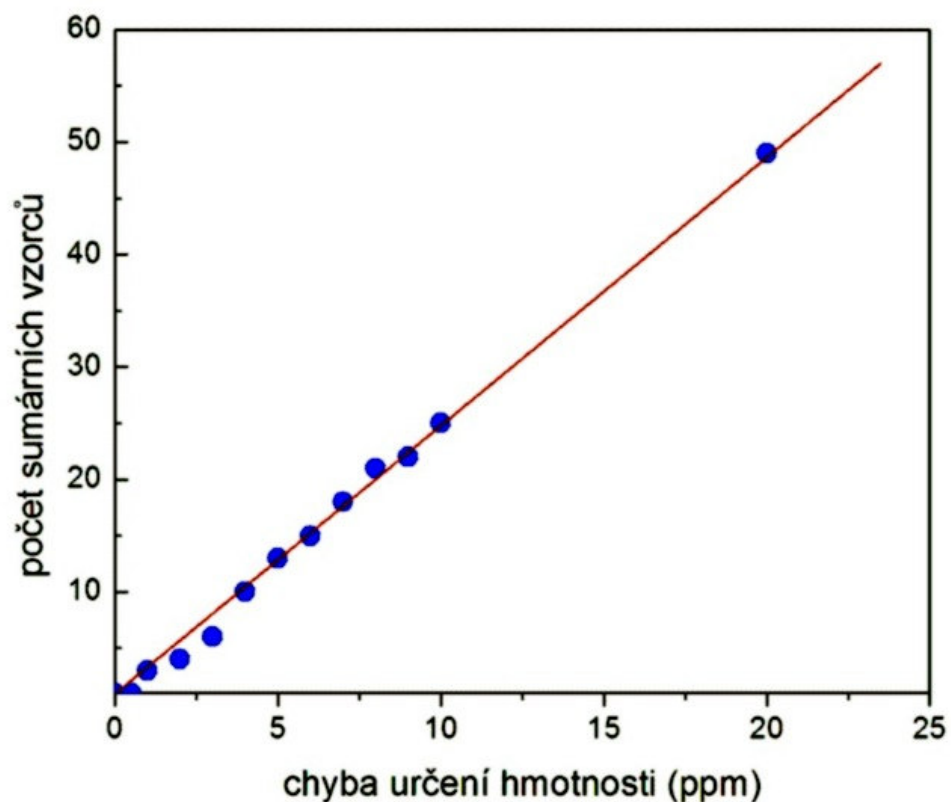
Constrains:

C: 0-100

H: 0-100

N: 0-10

O: 0-30

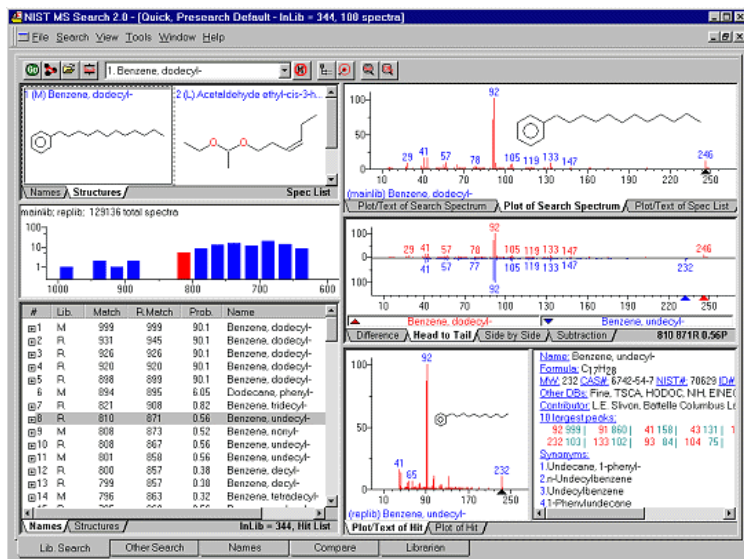


5.

Searching mass spectra libraries

Libraries of EI mass spectra

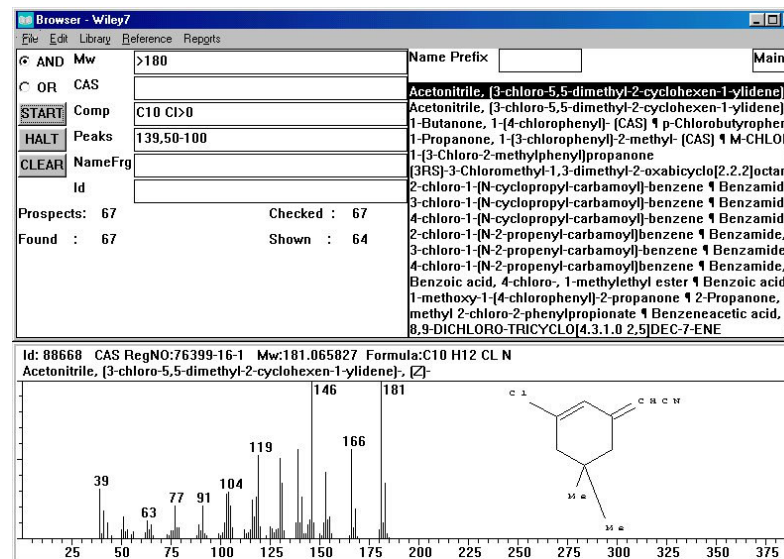
NIST/EPA/NIH Mass Spectral Library



NIST 11:

EI spectra of 212,961 compounds
 MS/MS spectra
 Kovats retention index values

Wiley Registry of Mass Spectral Data



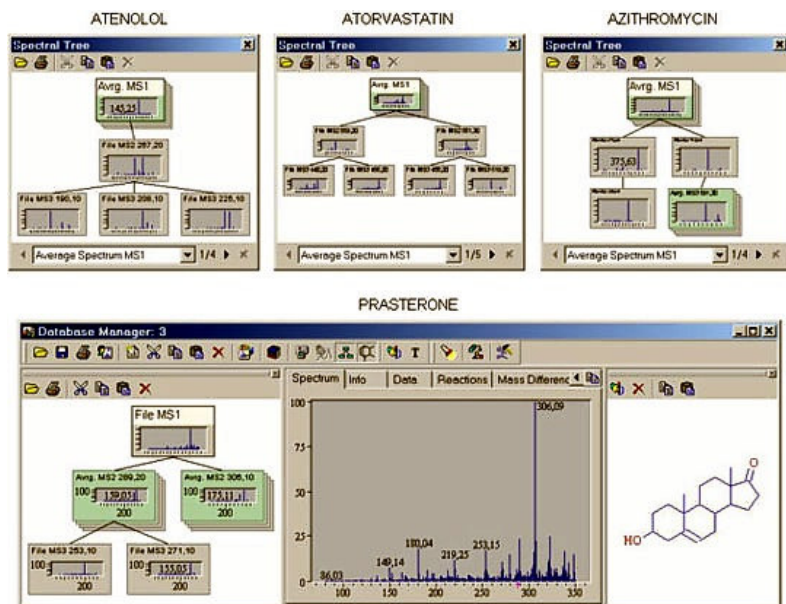
9th Edition:

663,000 EI spectra(70 eV)

NIST 05 installed on open access GC/MS

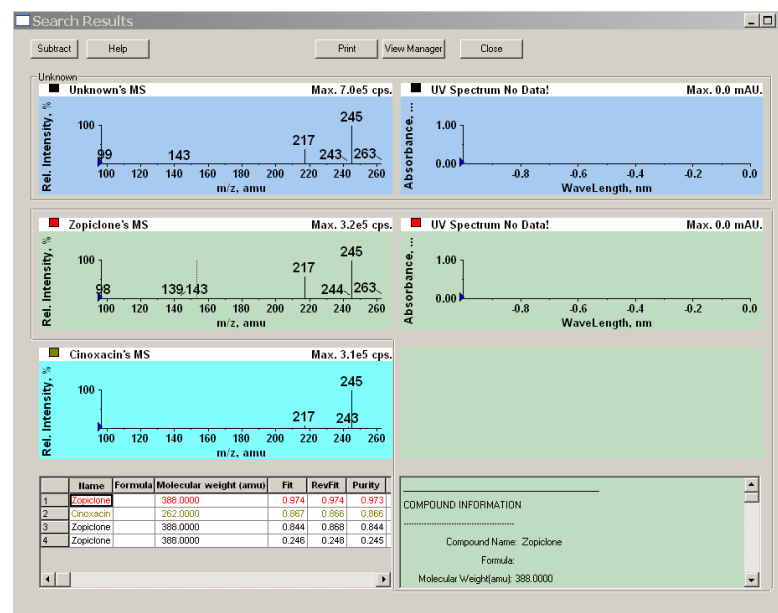
Libraries of ESI MS/MS mass spectra

HighChem's spectral library collection



MS/MS spectra > 1000 compounds
Drugs, natural products

MS Libraries for LC-MS/MS and ESI in-source CID-MS, University of Freiburg



MS/MS spectra ~ 800
compounds, drugs



<http://www.massbank.jp>

6.

Solving fragmentation spectra

Fragmentation of ions with even number of electrons (EE⁺)

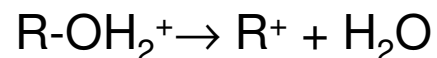
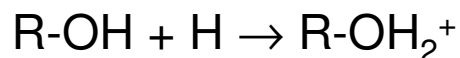
ESI, APCI (APPI, MALDI, DESI ...)

Fragmentation of EE⁺

1/ CID (MS/MS) of EE⁺ ([M+H]⁺, [M+Na]⁺, [M-H]⁻) formed by ESI

2/ fragmentation of ([M+H]⁺, [M+Na]⁺, [M-H]⁻) during APCI, APPI

Cleavage of neighboring bond to the charge site, charge migration



FRAGMENTATION of EE⁺:

The fragments are EE⁺ and a neutral fragment (not seen in the spectra)



EE⁺ ions are more stable than OE⁺

The spectra are simpler than EI spectra, thus provide less information.
They are sensitive to small changes in the structure.

Fragmentation of EE^+

Elimination of a neutral molecule depends on basicity and stability of the forming ion

Typical logical neutral losses:

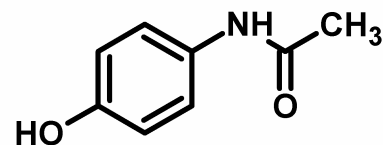
- 17: NH_3 – amines aliphatic, aromatic (+)
- 18: H_2O – oxygen-containing compounds (+/-)
- 27: HCN – amines aliphatic, aromatic, nitriles aromatic (+/-)
- 28: CO – aldehydes, ketones, nitroaromates (+/-)
- 32: CH_3OH – methyl esters (+)
- 42: $CH_2C=O$ – N-acetyl derivatives (+/-)
- 44: CO_2 – carboxylic acids, carbamates (+/-)
- 80: SO_3 – sulfonic acids(+/-)
- 162: anhydroglucose – glucosides (+/-)

Impossible “forbidden” neutral losses: 3-14, 21-25, 37-40

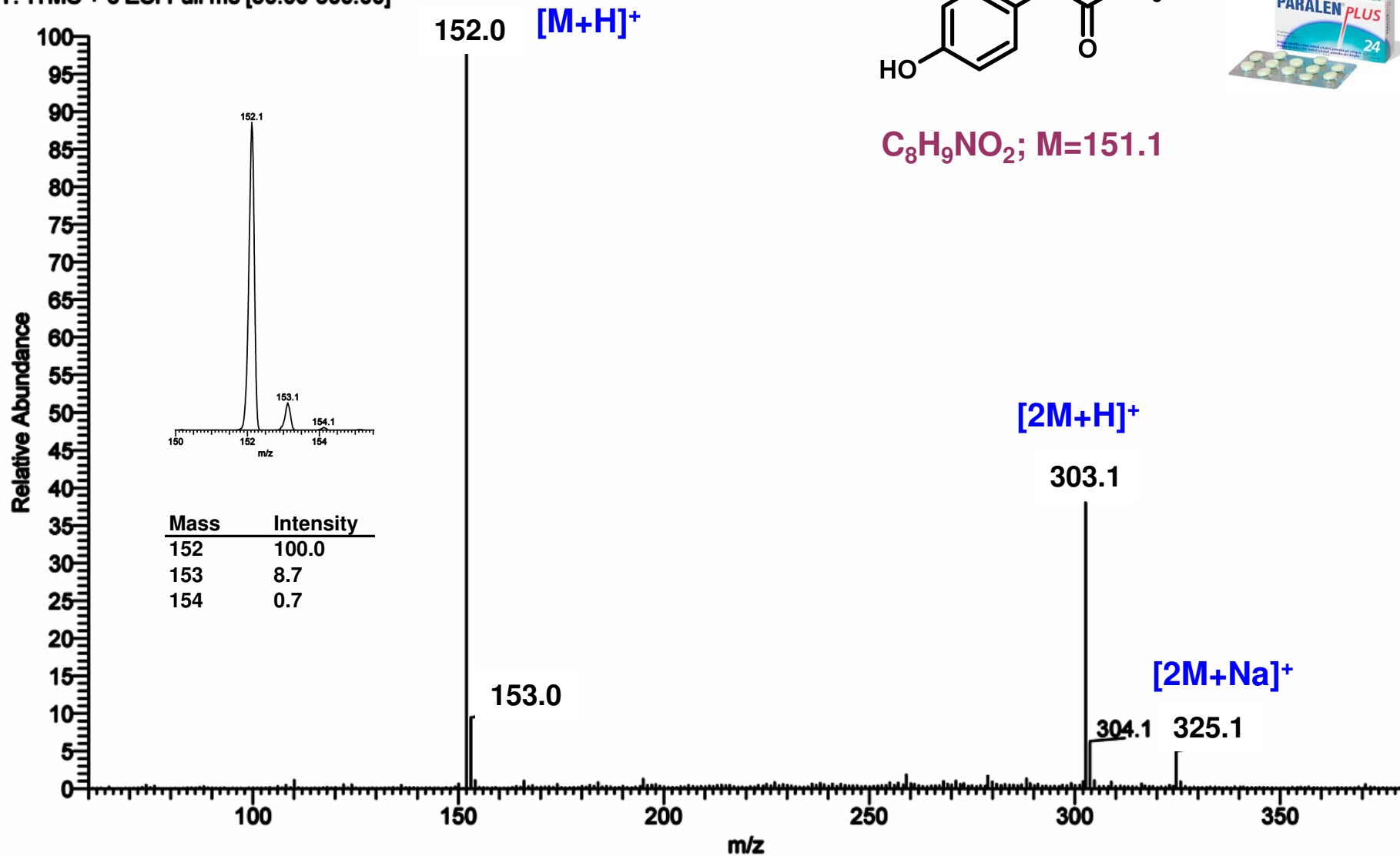
Paracetamol N-(4-hydroxyphenyl)acetamide

ESI+, MS

JC_LEKY_Pan_04_080720211508 #1-22 RT: 0.00-0.25 AV: 22 NL: 2.40E4
T: ITMS + c ESI Full ms [50.00-500.00]



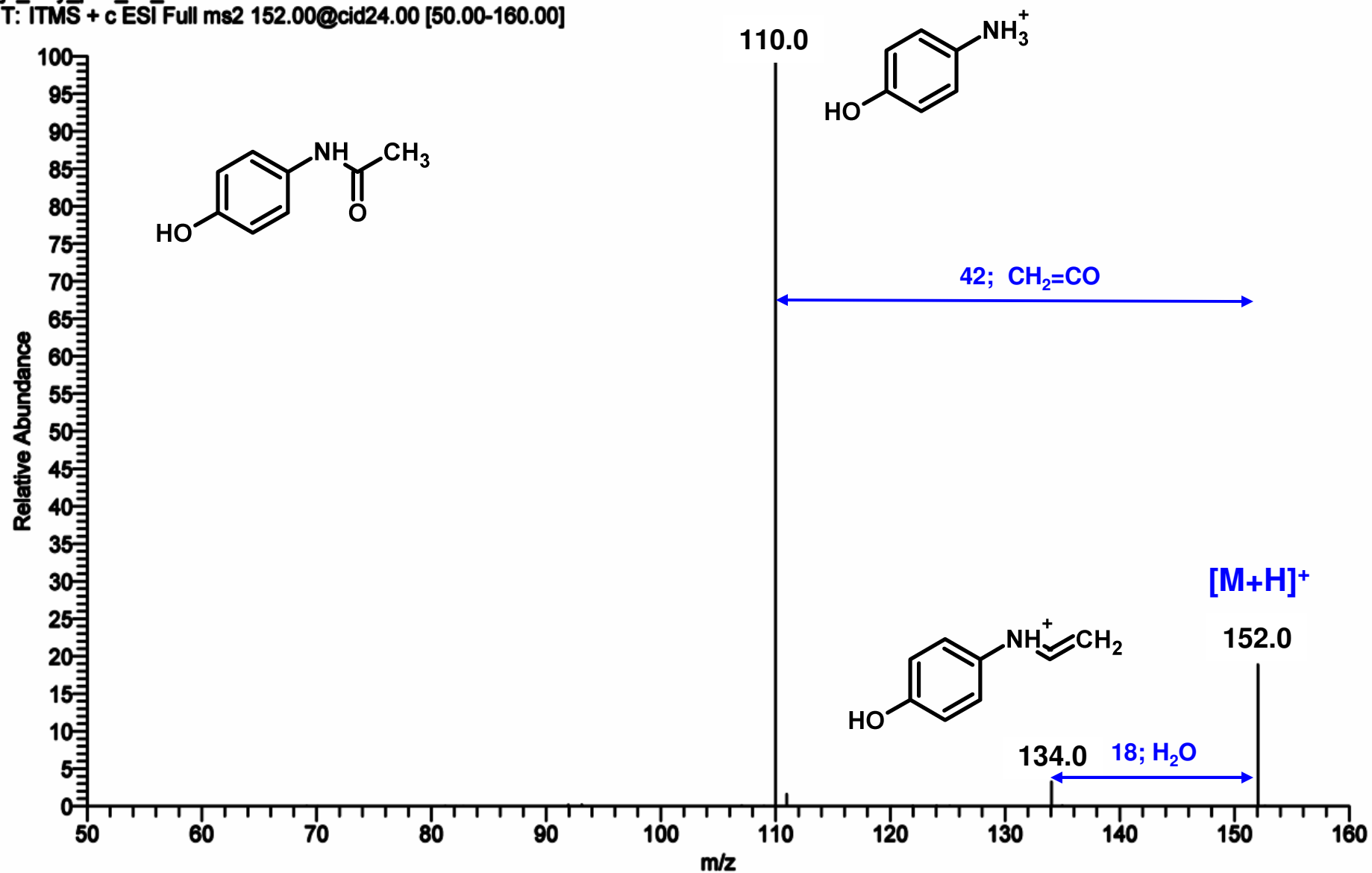
$C_8H_9NO_2$; $M=151.1$



Paracetamol N-(4-hydroxyphenyl)acetamide

ESI+, MS/MS

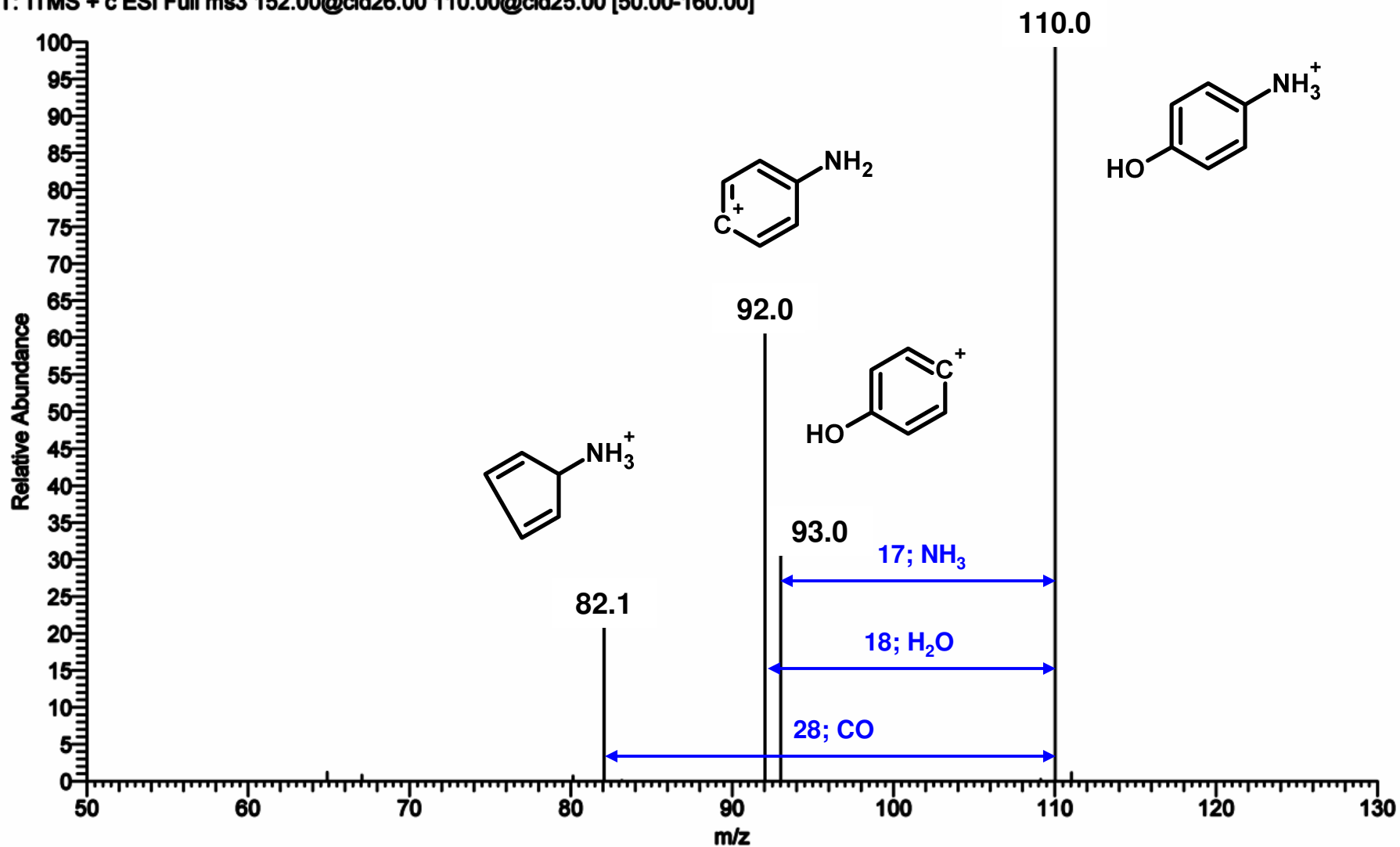
jc_leky_pan_03_080720211358#1-36 RT: 0.00-0.24 AV: 35 NL: 1.07E3
T: ITMS + c ESI Full ms2 152.00@cid24.00 [50.00-160.00]



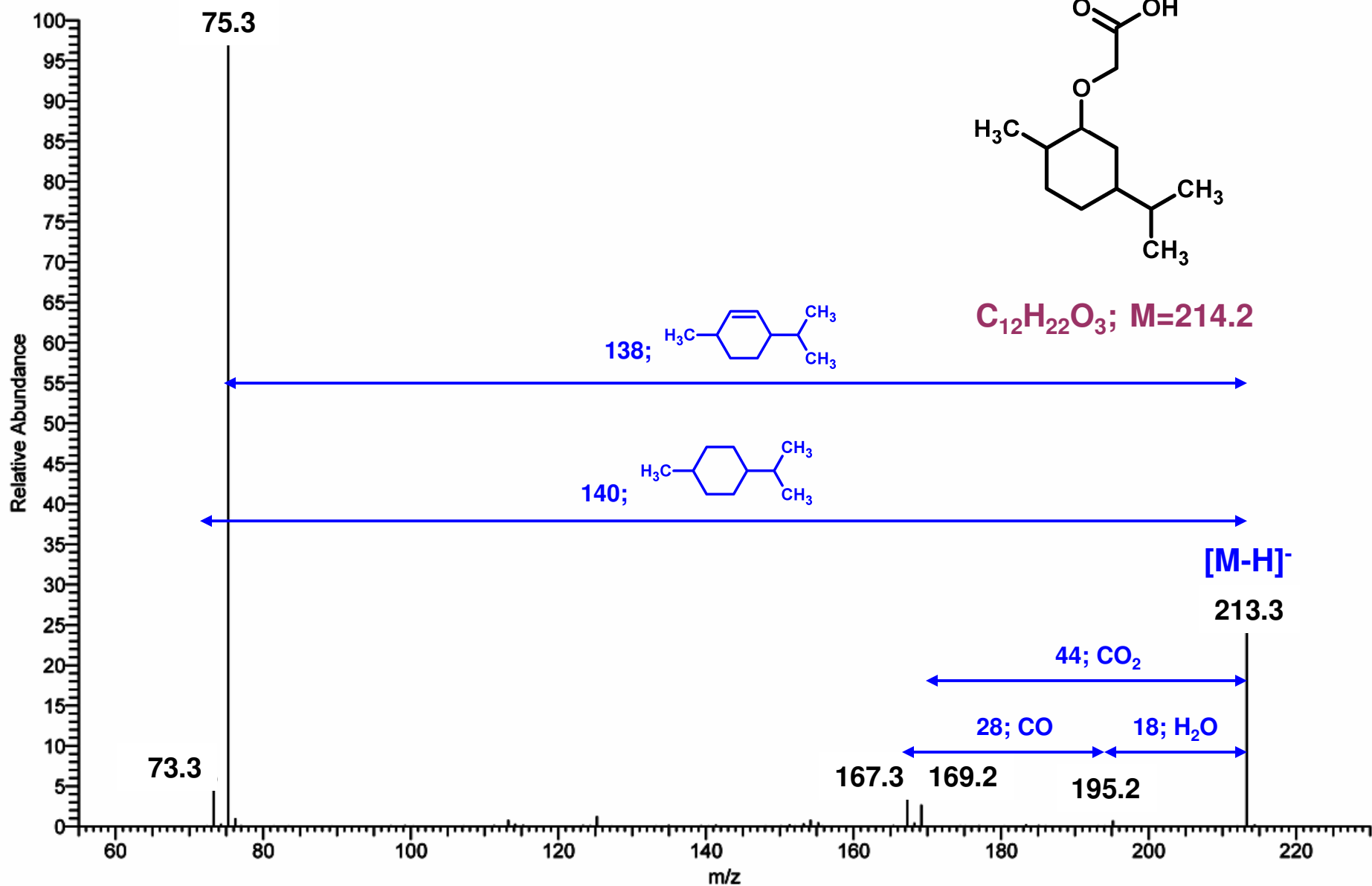
Paracetamol N-(4-hydroxyphenyl)acetamide

ESI+, MS³

JC_LEKY_Pan_18 #1-464 RT: 0.00-2.00 AV: 464 NL: 1.12E1
T: ITMS + c ESI Full ms3 152.00@cid26.00 110.00@cid25.00 [50.00-160.00]



Menthyloxyacetic acid

ESI-, MS/MS2789jc_090716170932 #1-149 RT: 0.00-2.00 AV: 149 NL: 9.82E1
T: ITMS - c ESI Full ms2 213.40@cid29.00 [55.00-250.00]

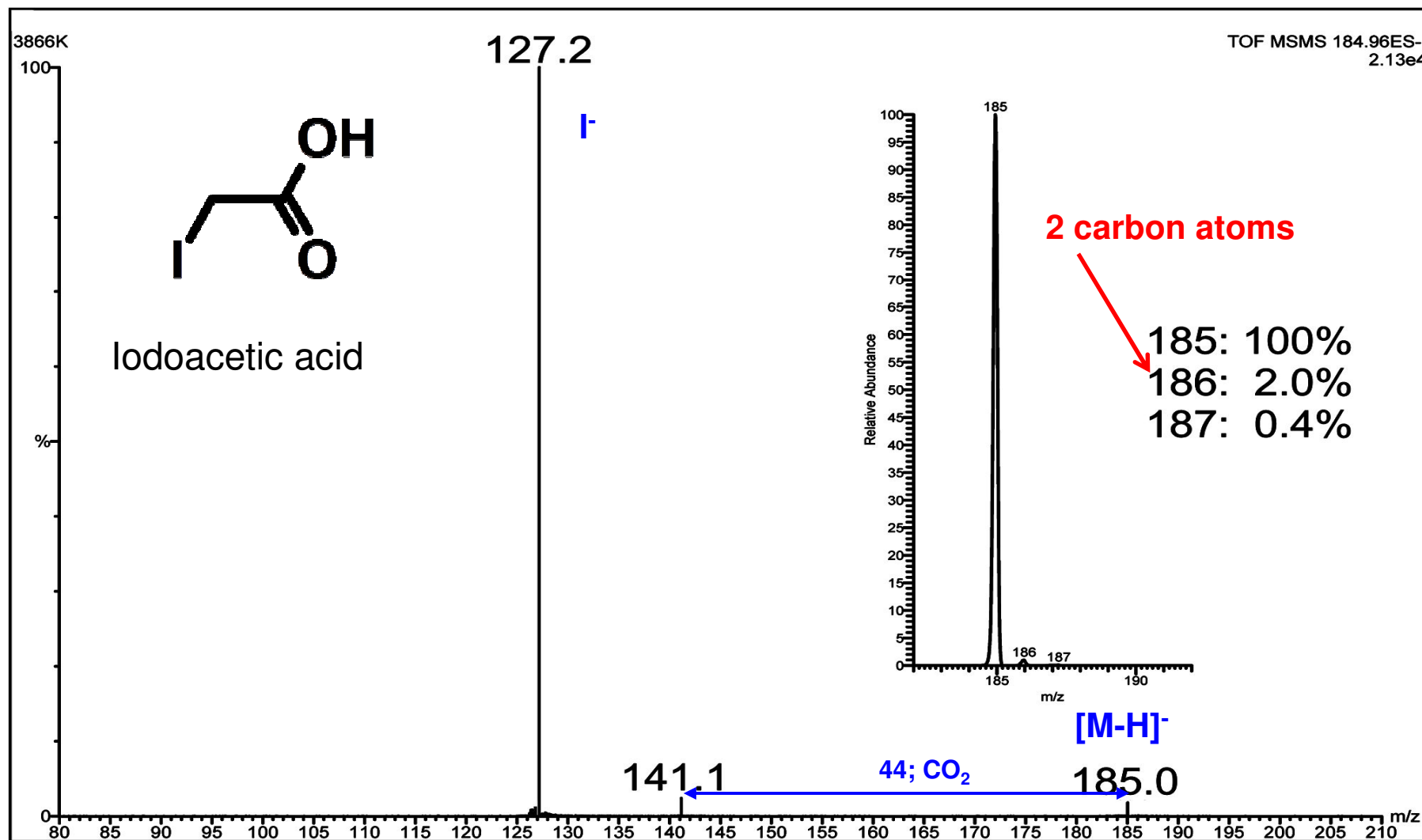
Unknown

ESI-, MS/MS



What is the structure?

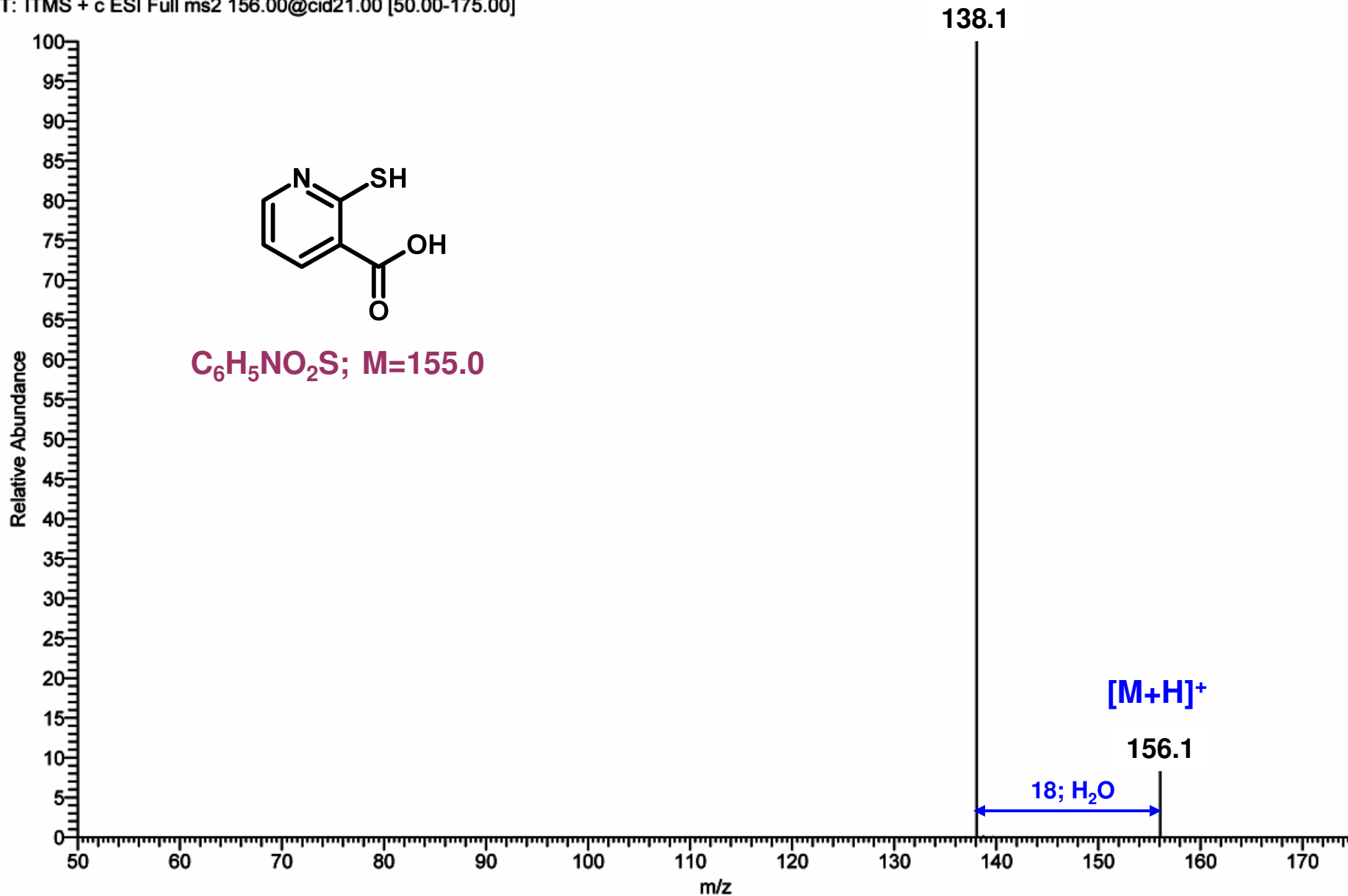
MS/MS m/z 185



2-Mercaptopyridine-3-carboxylic acid

ESI+, MS/MS

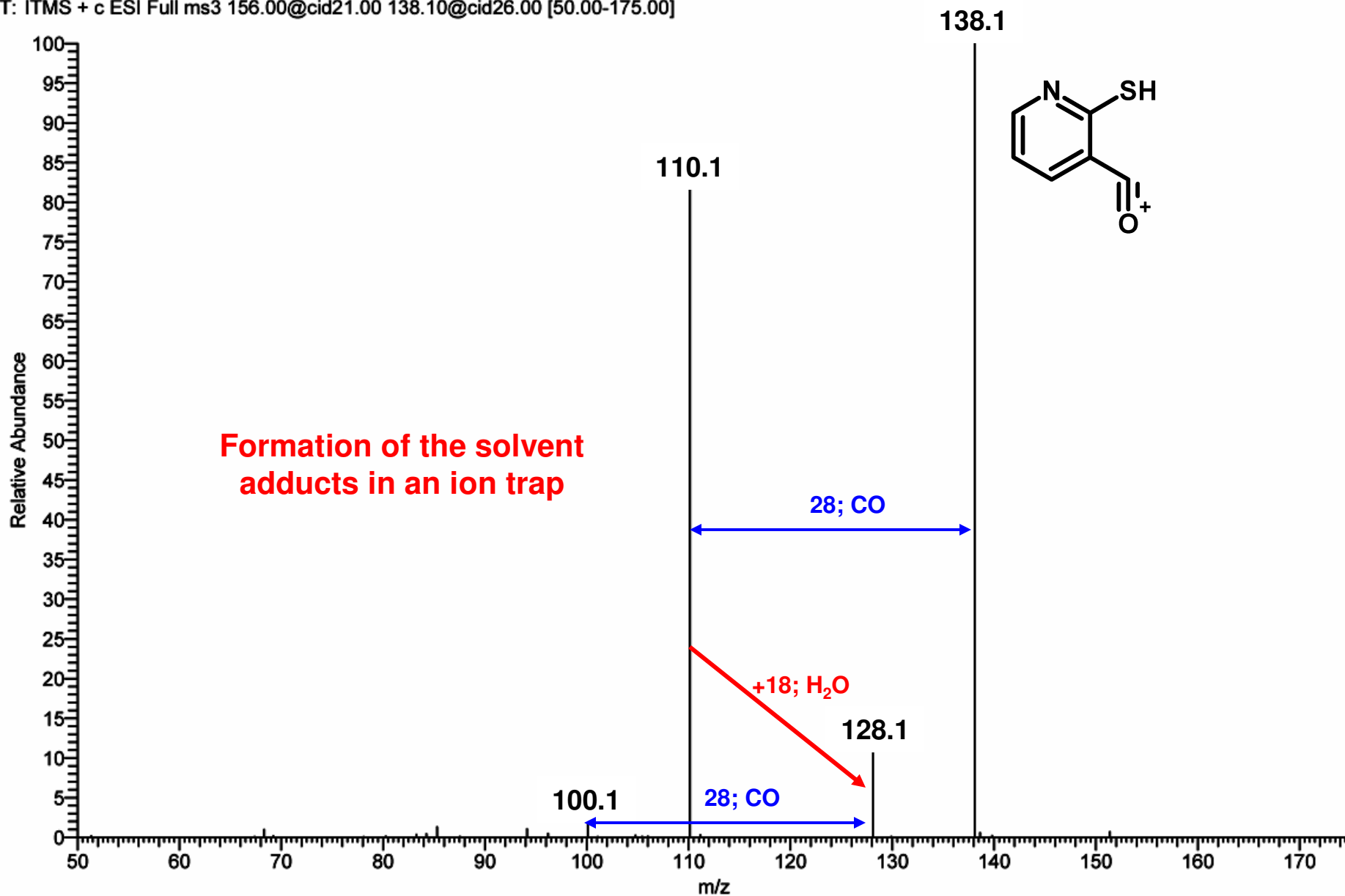
2786jc #1-147 RT: 0.00-1.02 AV: 147 NL: 2.76E3
T: ITMS + c ESI Full ms2 156.00@cid21.00 [50.00-175.00]



2-Mercaptopyridine-3-carboxylic acid

ESI+, *MS*³

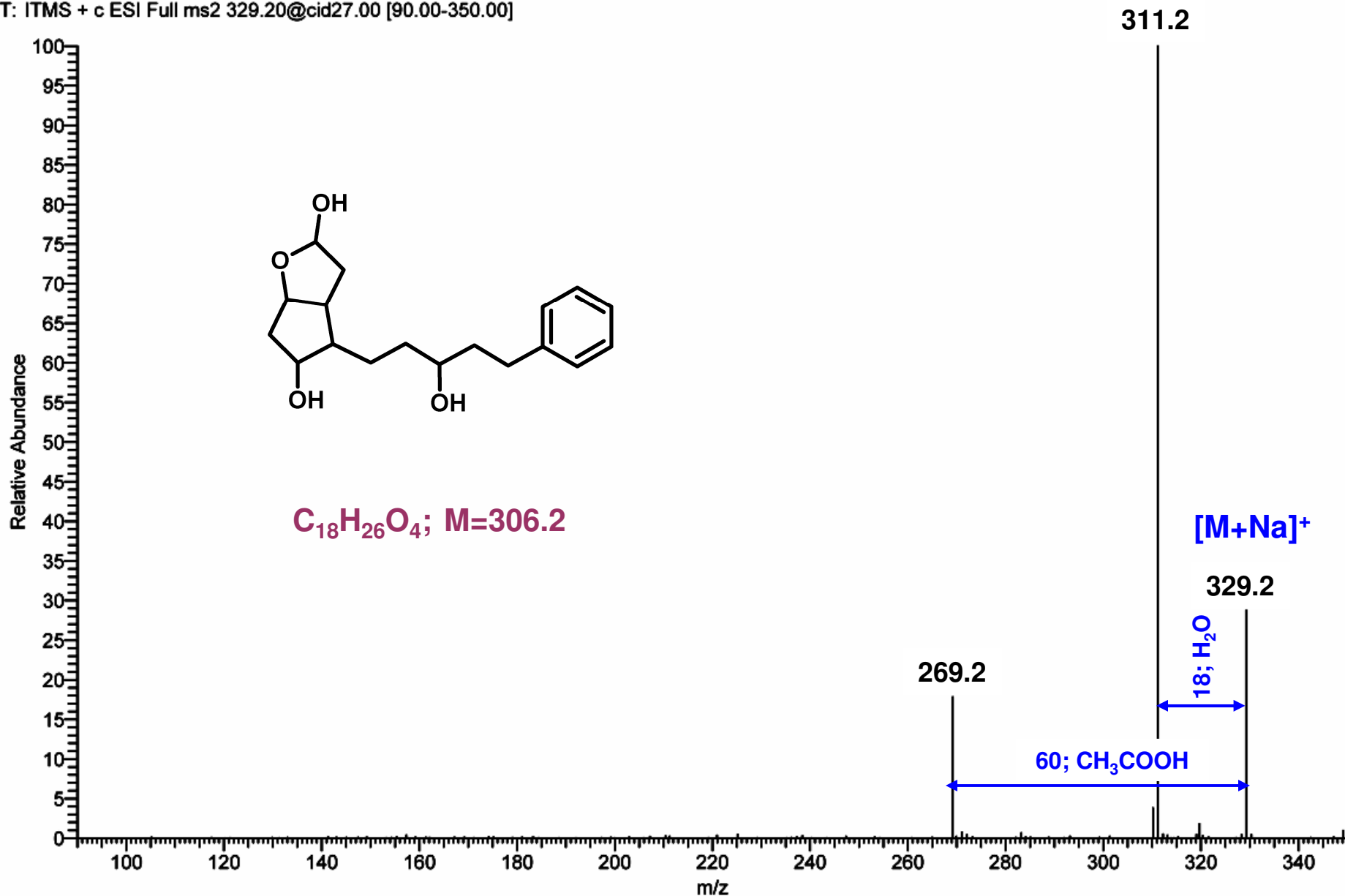
2787jc #1-230 RT: 0.00-2.01 AV: 230 NL: 1.73E1
T: ITMS + c ESI Full ms3 156.00@cid21.00 138.10@cid26.00 [50.00-175.00]



Cyclopentafuranol-derivative

ESI+, MS/MS

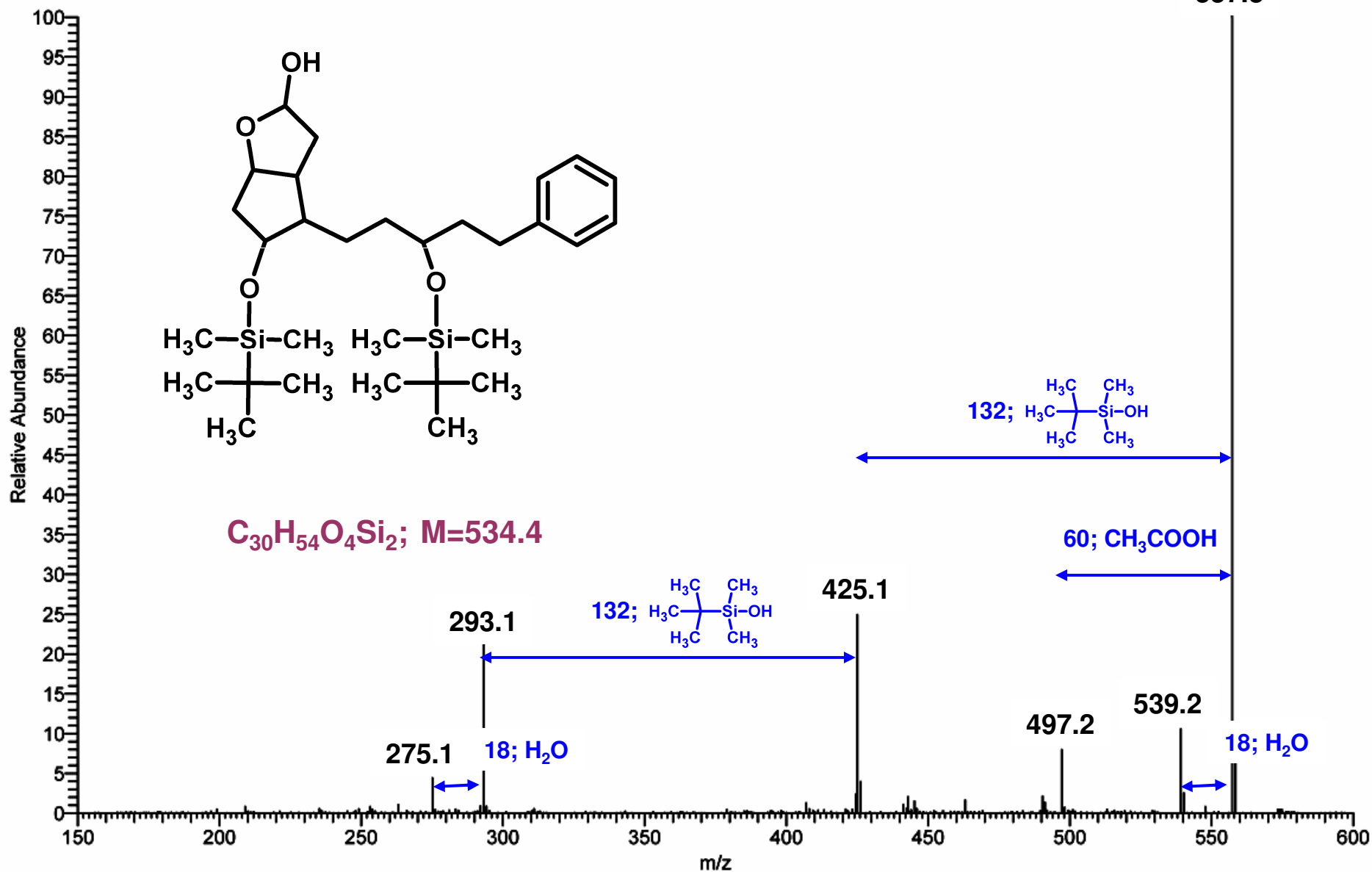
1748jc #1-226 RT: 0.00-1.51 AV: 226 NL: 5.26E1
T: ITMS + c ESI Full ms2 329.20@cid27.00 [90.00-350.00]



Cyclopentafuranol-derivative, disil

ESI+, MS/MS

1644jc #1-749 RT: 0.00-5.02 AV: 749 NL: 2.22E1
T: ITMS + c ESI Full ms2 557.30@cid27.00 [150.00-600.00]

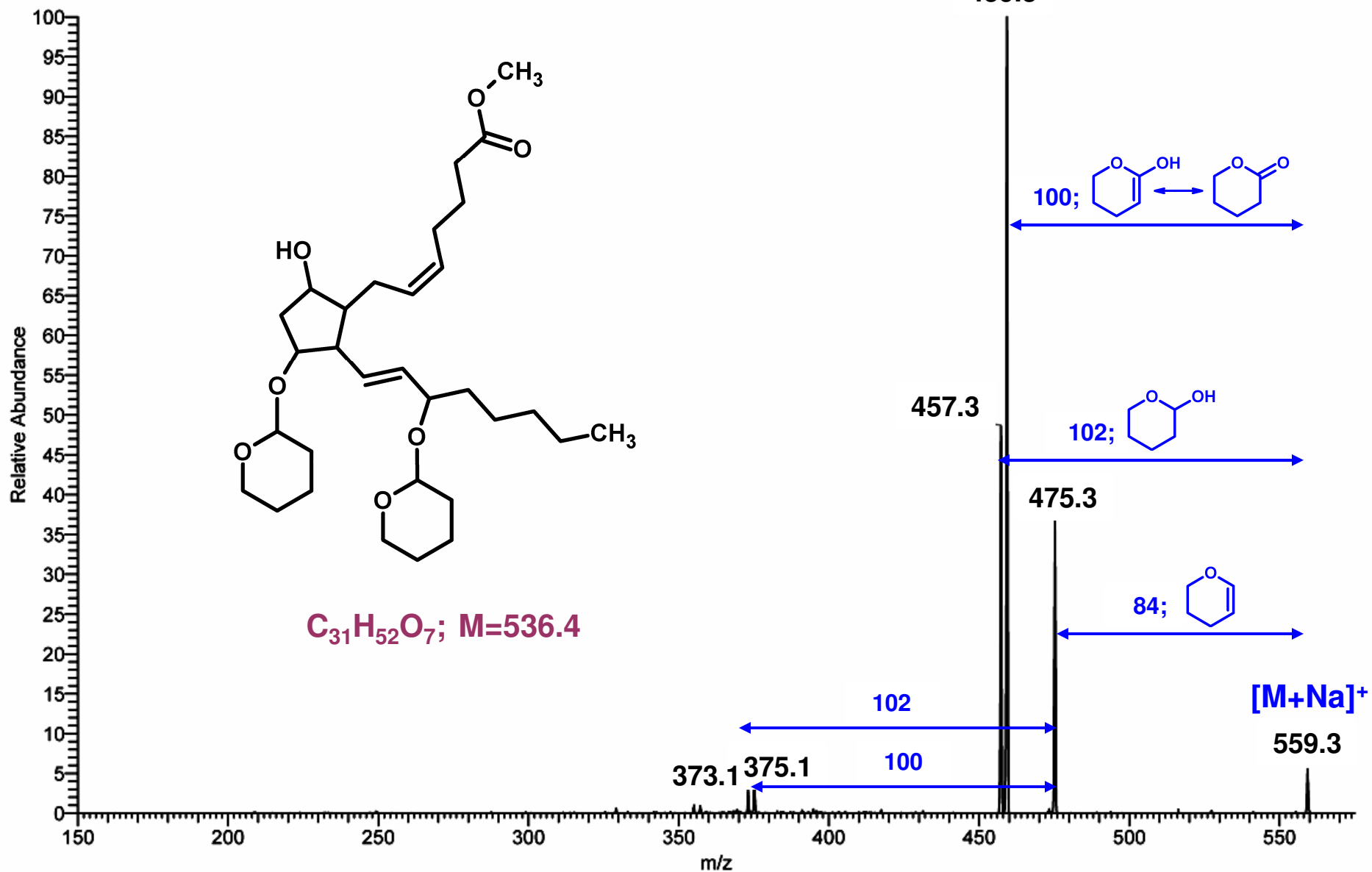


PGF_{2a}-methylester, diTHP

ESI+, MS/MS

3378jc #1-64 RT: 0.01-0.99 AV: 64 NL: 8.64E5

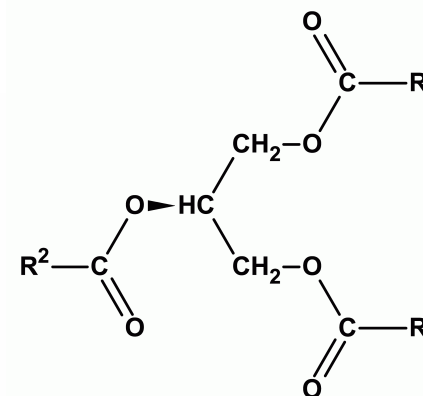
T: + p Full ms2 559.40@cid37.00 [150.00-600.00]



Triacylglycerols

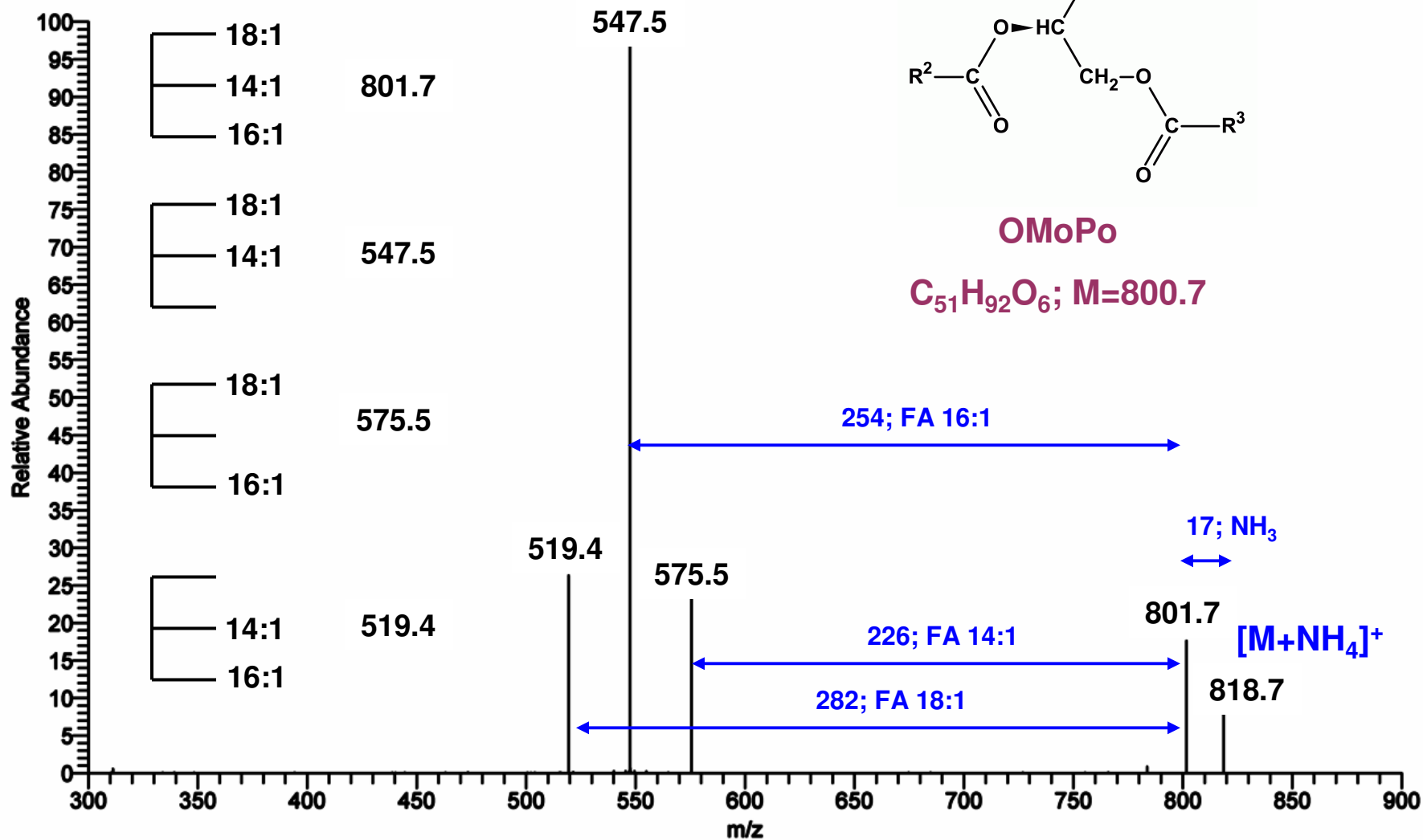
APCI+, MS/MS

JC_TAG_Bter_MSMS01 #6162 RT: 64.99 AV: 1 NL: 3.82E6
T: FTMS + c APCI corona d Full ms2 818.72@cid30.00 [215.00-830.00]



OMoPo

C₅₁H₉₂O₆; M=800.7



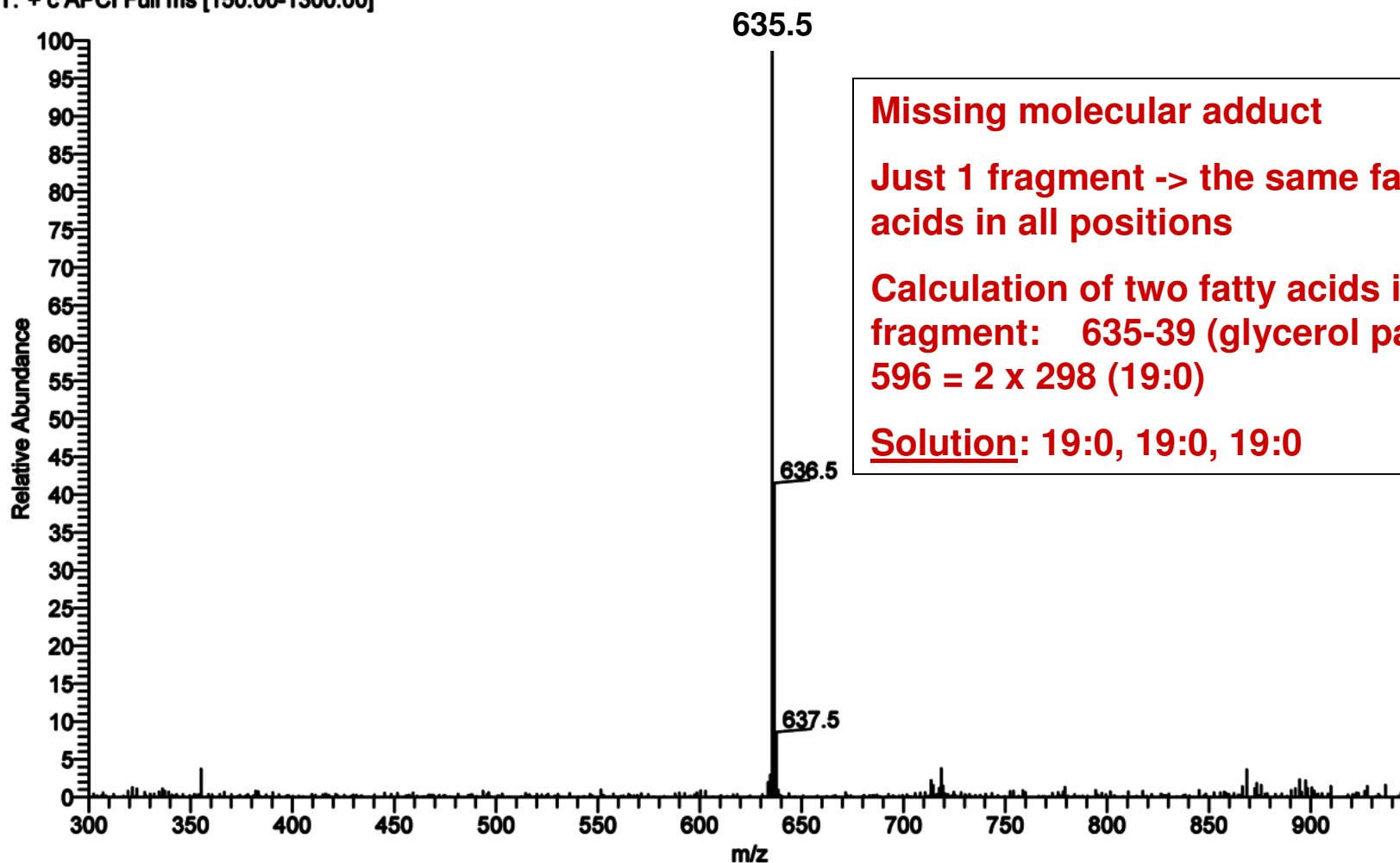
Triacylglycerols



What is the structure?

APCI+, MS

1377ek #4346-4381 RT: 108.29-109.11 AV: 36 SB: 44 106.70-107.89 NL: 4.68E5
T: + c APCI Full ms [150.00-1300.00]



Missing molecular adduct

Just 1 fragment -> the same fatty acids in all positions

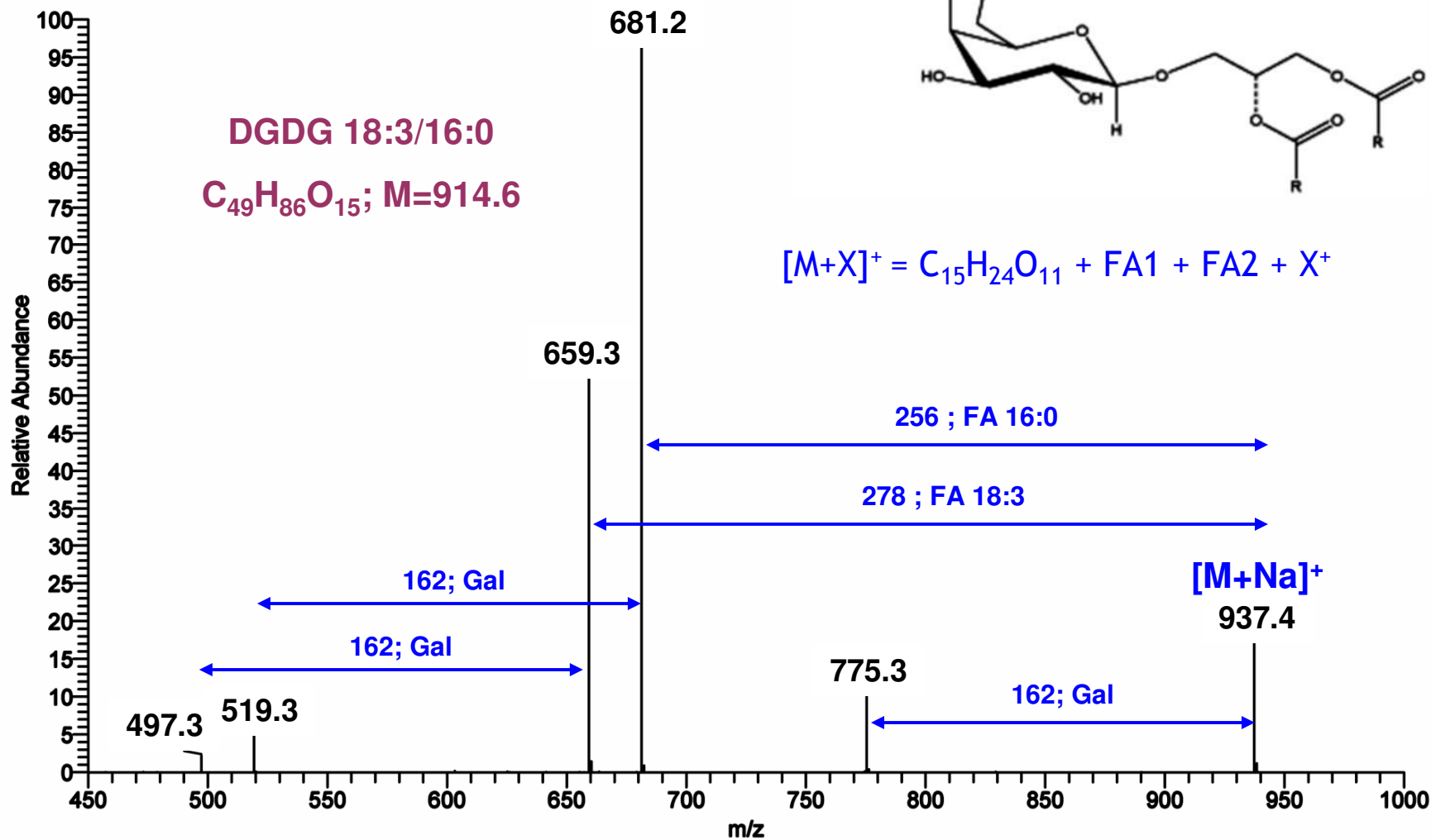
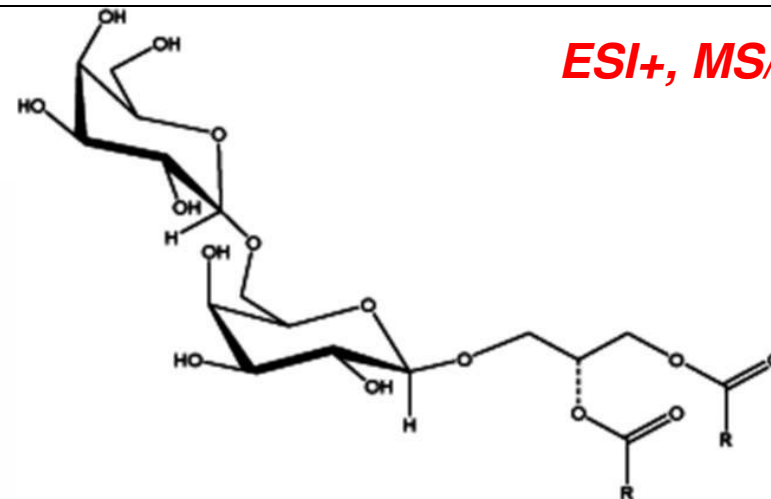
Calculation of two fatty acids in the fragment: $635 - 39$ (glycerol part) = $596 = 2 \times 298$ (19:0)

Solution: 19:0, 19:0, 19:0

Digalaktosyldiacylglycerols

ESI+, MS/MS

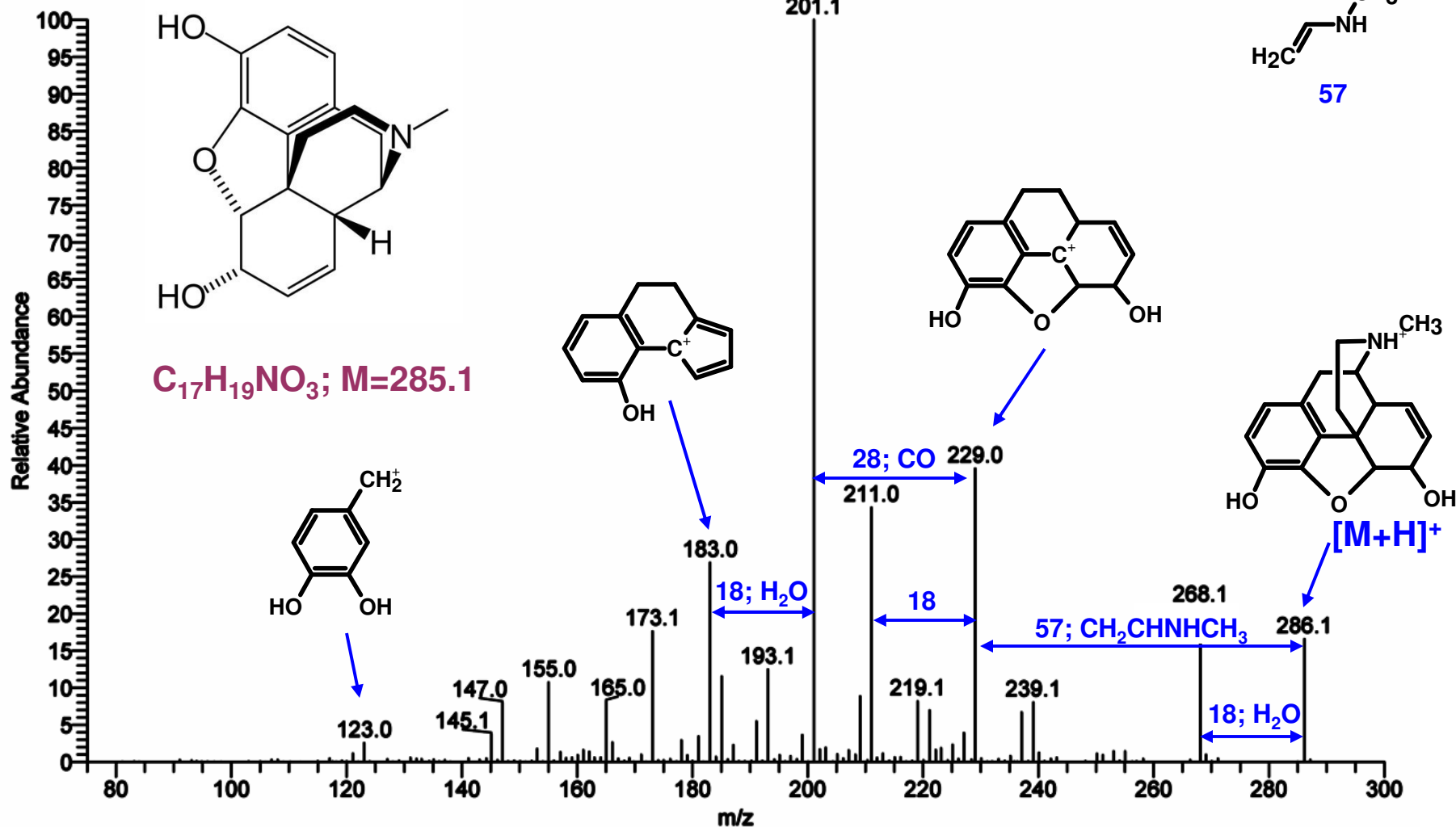
1656MK01 #1342-1375 RT: 21.08-21.58 AV: 17 NL: 1.26E3
 T: ITMS + c ESI d Full ms2 937.47@cid28.00 [245.00-950.00]




Morphine

ESI+, MS/MS

2229jc_080801195950 #1-11 RT: 0.00-0.47 AV: 11 NL: 1.81E2
T: ITMS + c ESI Full ms2 286.10@cid27.00 [75.00-300.00]

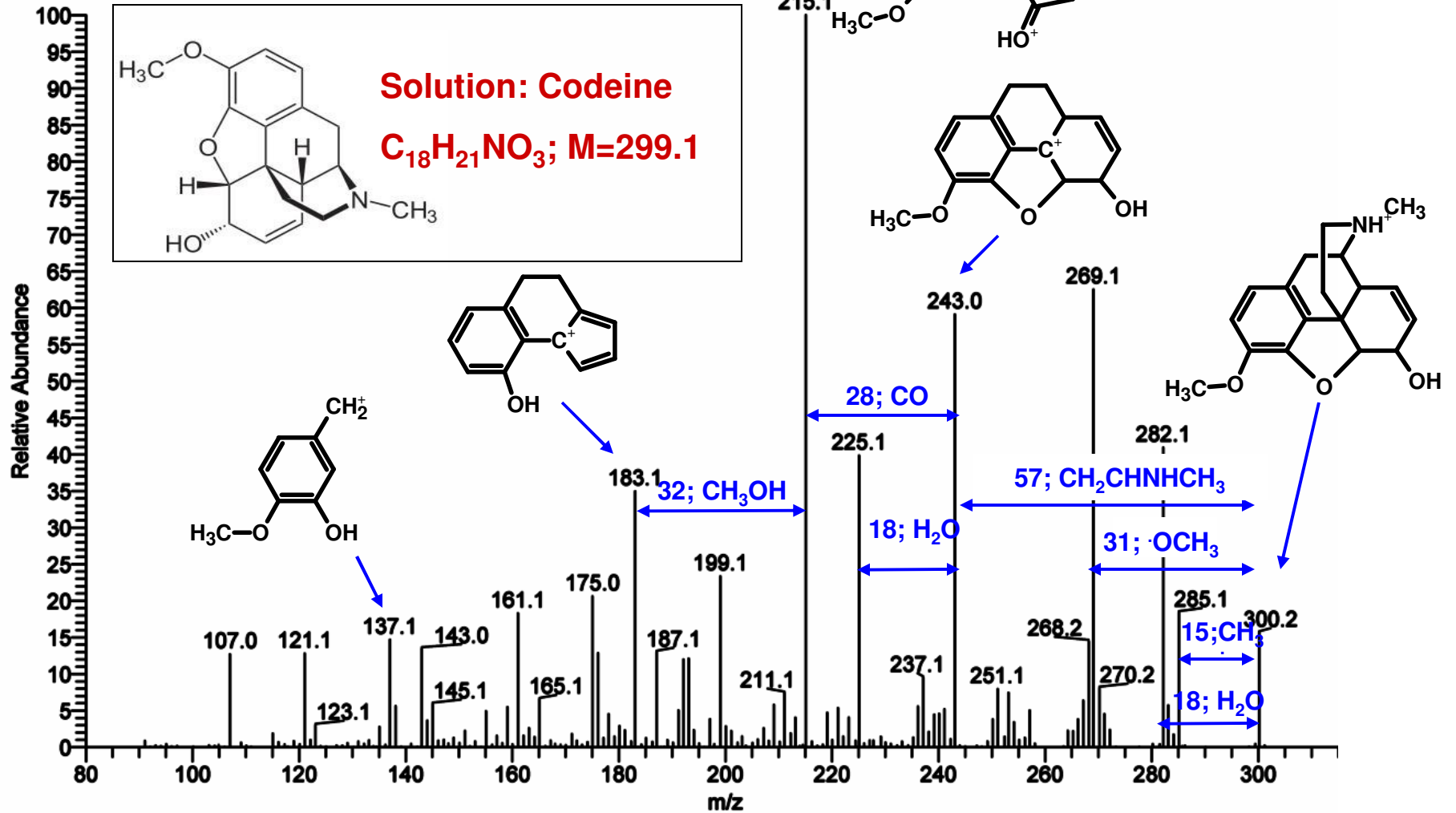
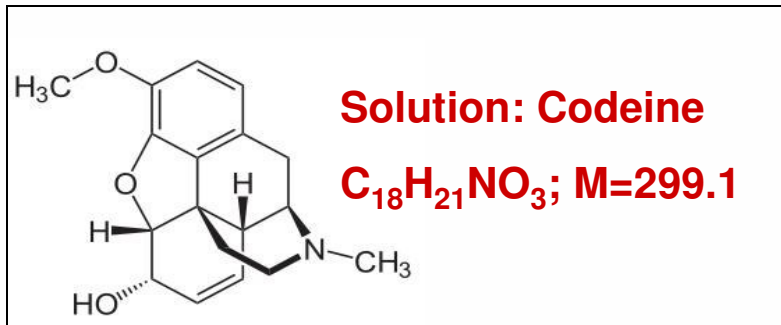


Unknown

 What is the structure?

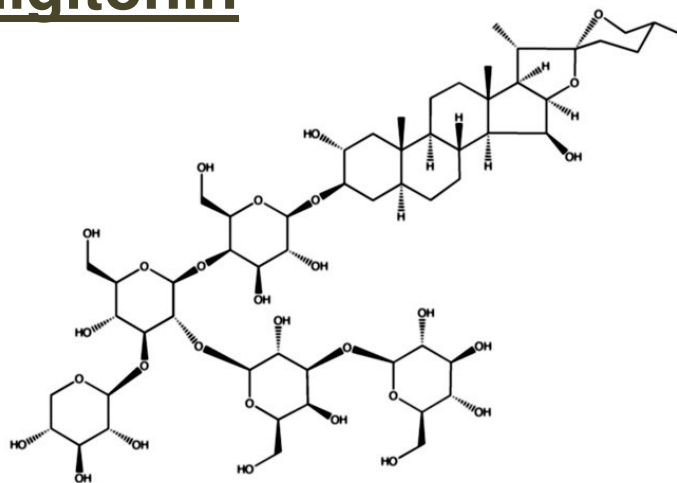
ESI+, MS/MS

2233jc #1-22 RT: 0.00-2.00 AV: 22 NL: 1.51E1
 F: ITMS + c ESI Full ms2 300.10@cid27.00 [80.00-320.00]

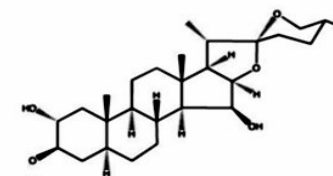


Glycosides - digitonin

Digitonin
 $C_{56}H_{92}O_{29}$



xyl - gluc - gal
|
gluc - gal



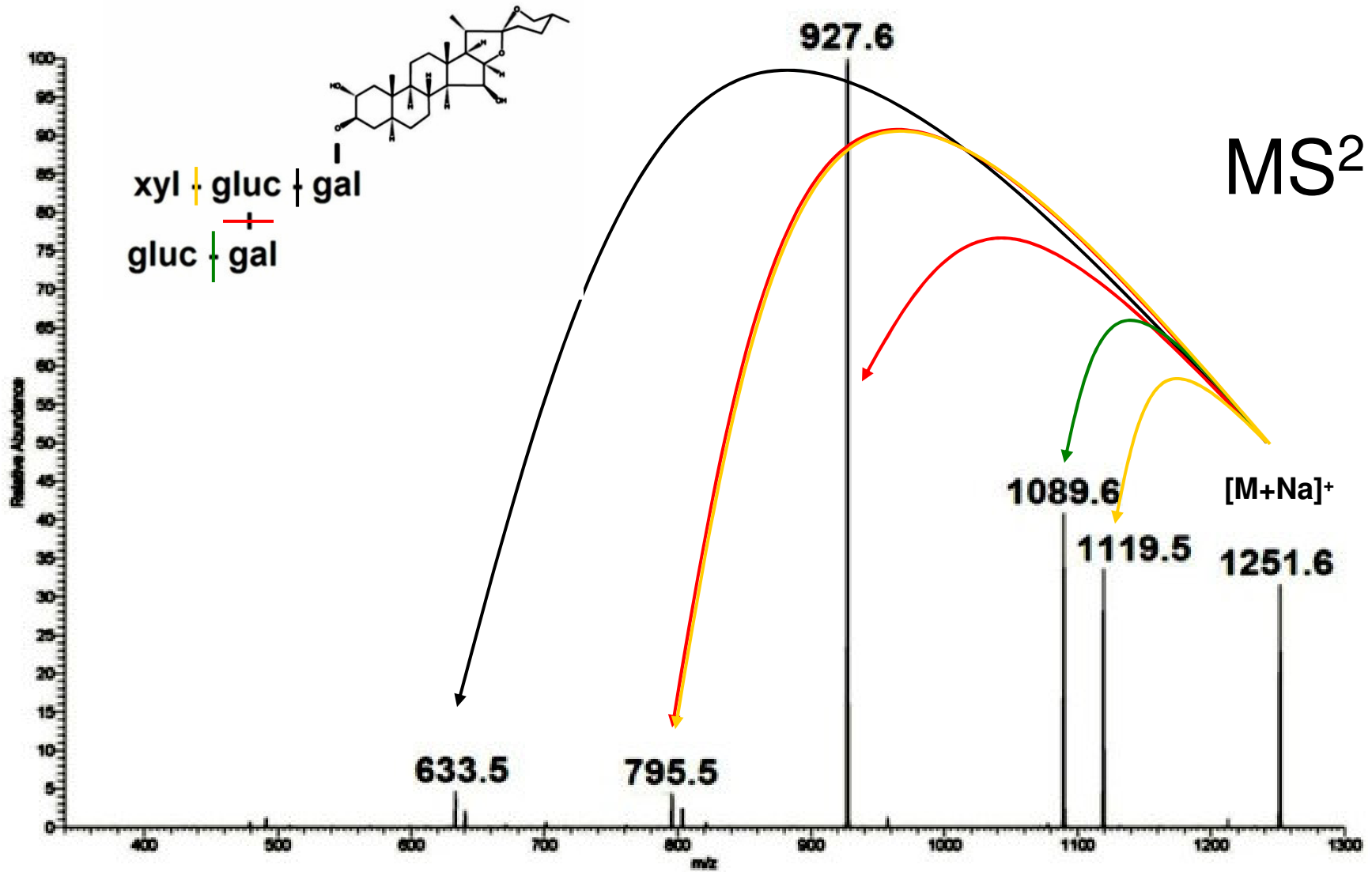
ESI ionization – formation of the molecular adduct $[M+Na]^+$

MS/MS fragmentation – cleavage by elimination of neutral sugars

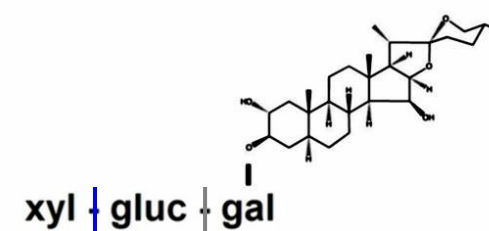
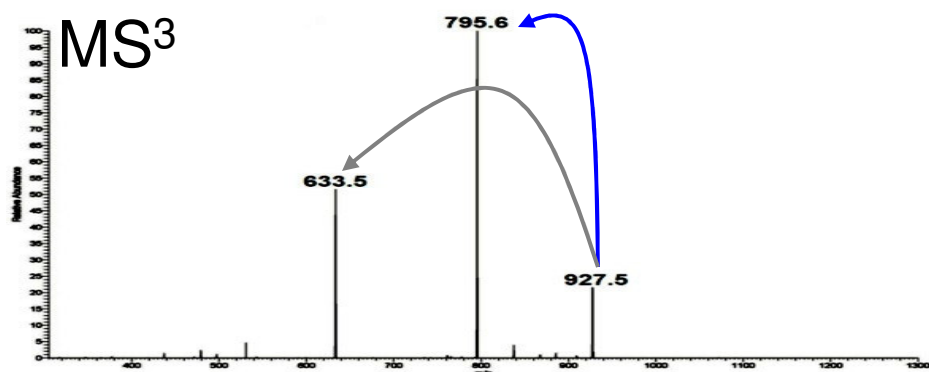
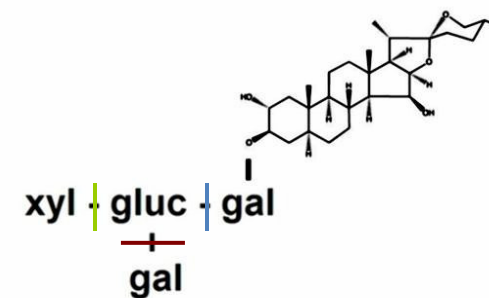
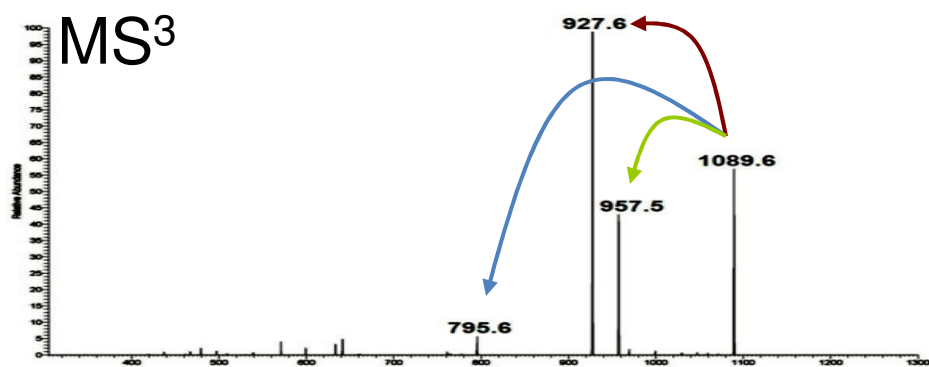
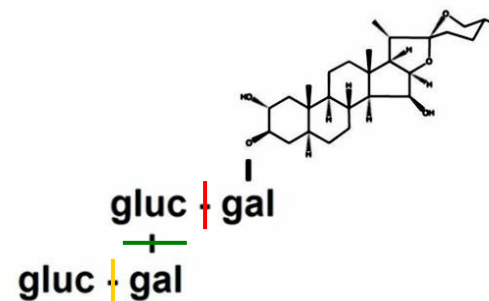
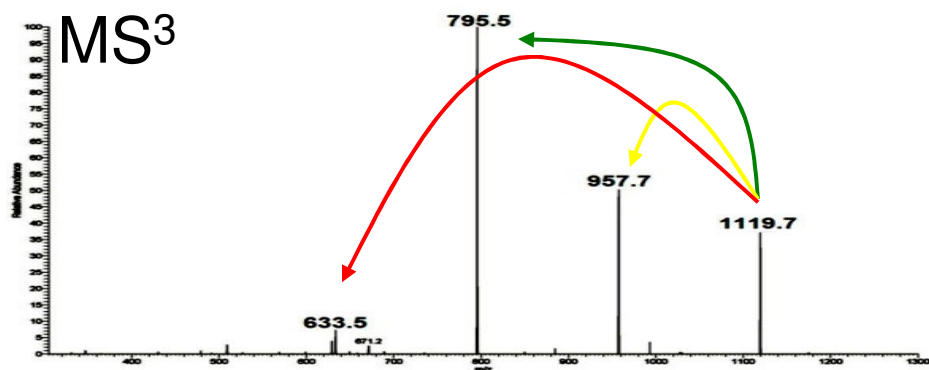
Hexose: loss of **162 u** (the molecule – H_2O)

Pentose: loss of **132 u** (the molecule – H_2O)

Glycosides - digitonin



Glycosides - digitonin



Fragmentation of ions with odd number of electrons (OE^{\bullet})

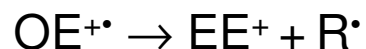
EI

Fragmentation of $OE^{\bullet+}$

EI fragments are formed already in MS step (it is not necessary to use fragmentation techniques such as CID, etc.)

FRAGMENTATION of $OE^{\bullet+}$

I. formation of an ion with even number of electrons and a radical



II. formation of an ion with odd number of electrons and a neutral specie

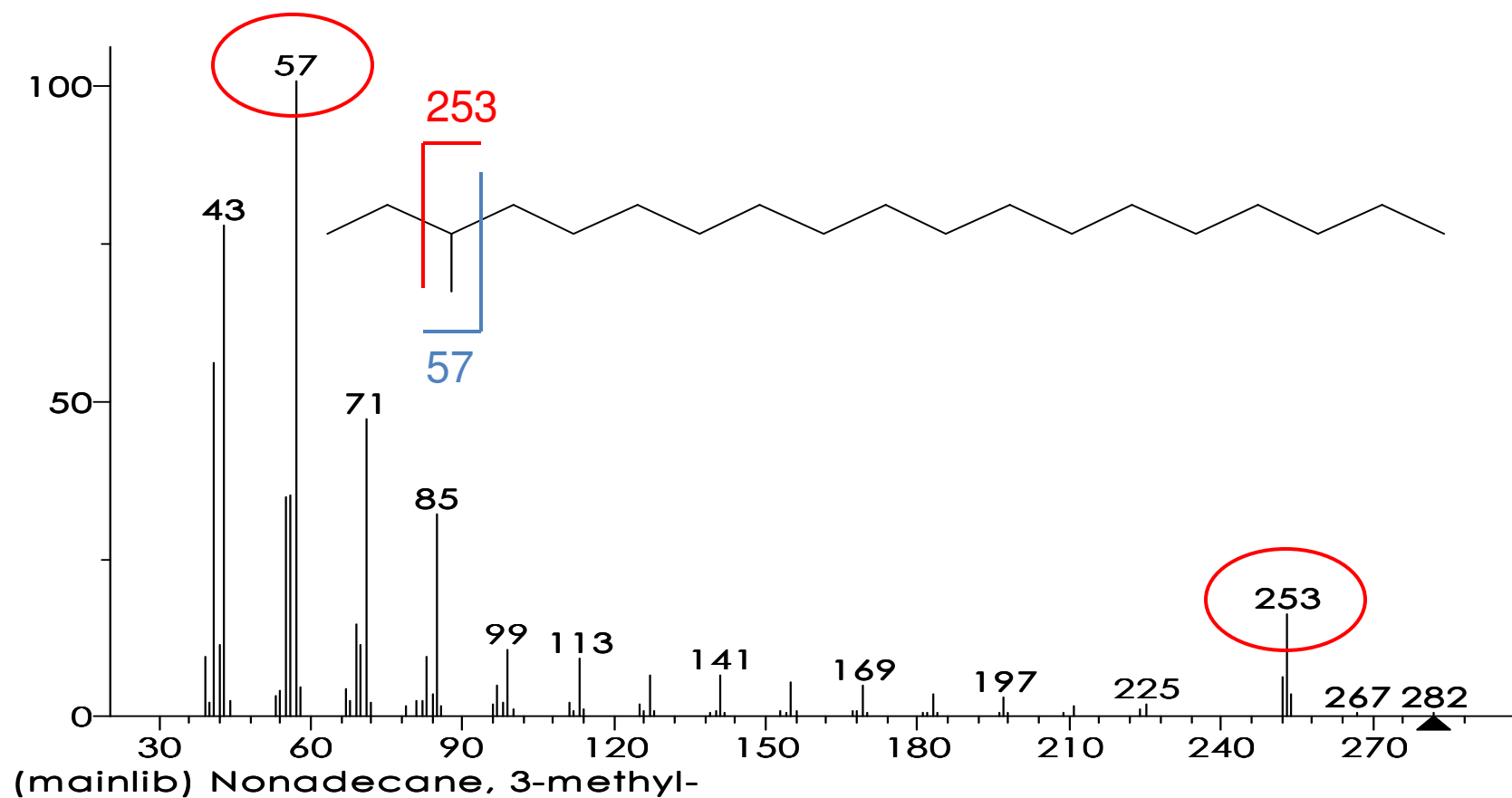


Information-rich spectra are obtained, can be used as a "fingerprint" for the creation of libraries of spectra

Only monomolecular reactions take place

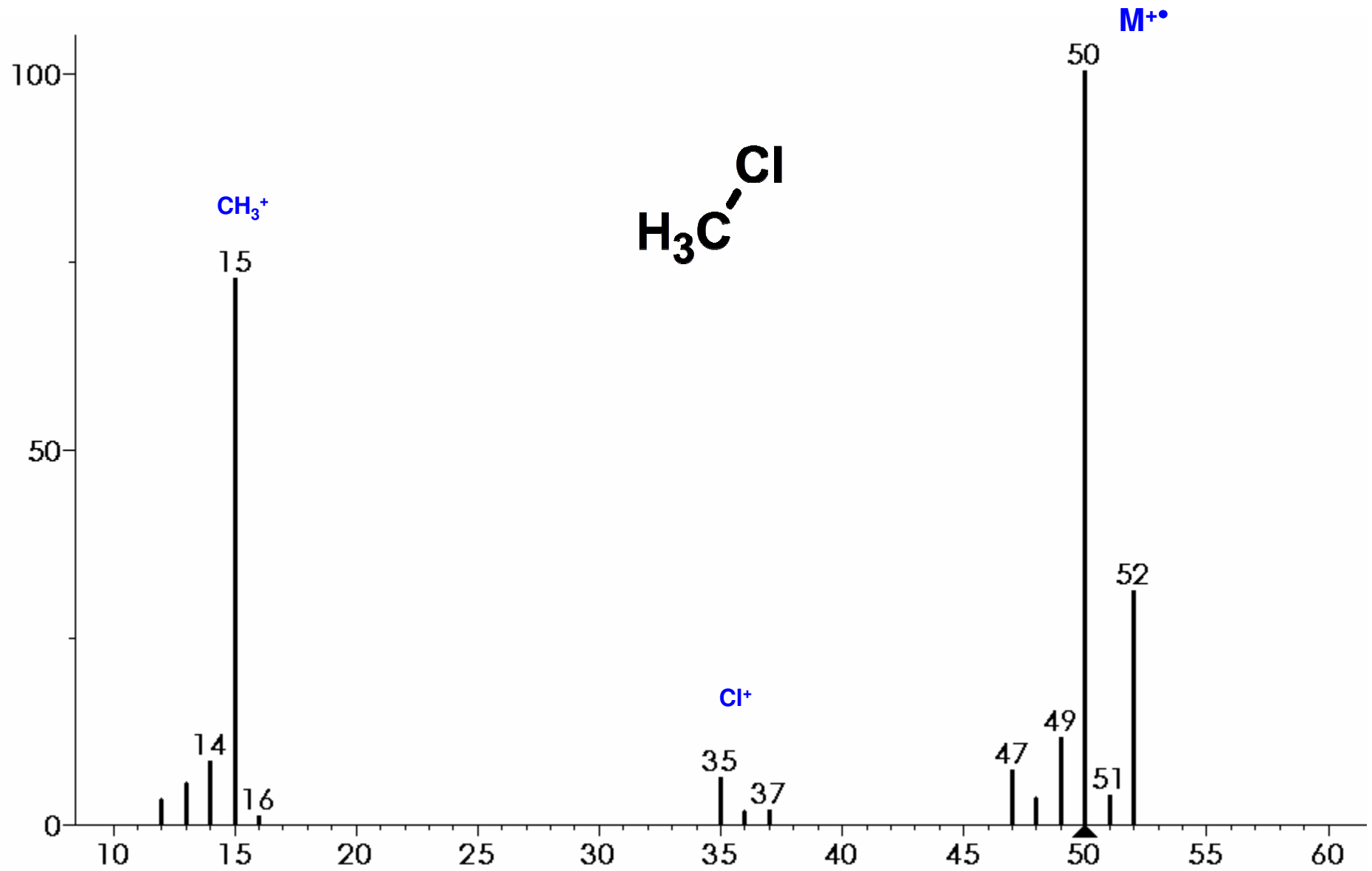
Branched hydrocarbons

- the ion intensity depends on the ability of fragments to stabilize the charge



Chloromethane

EI

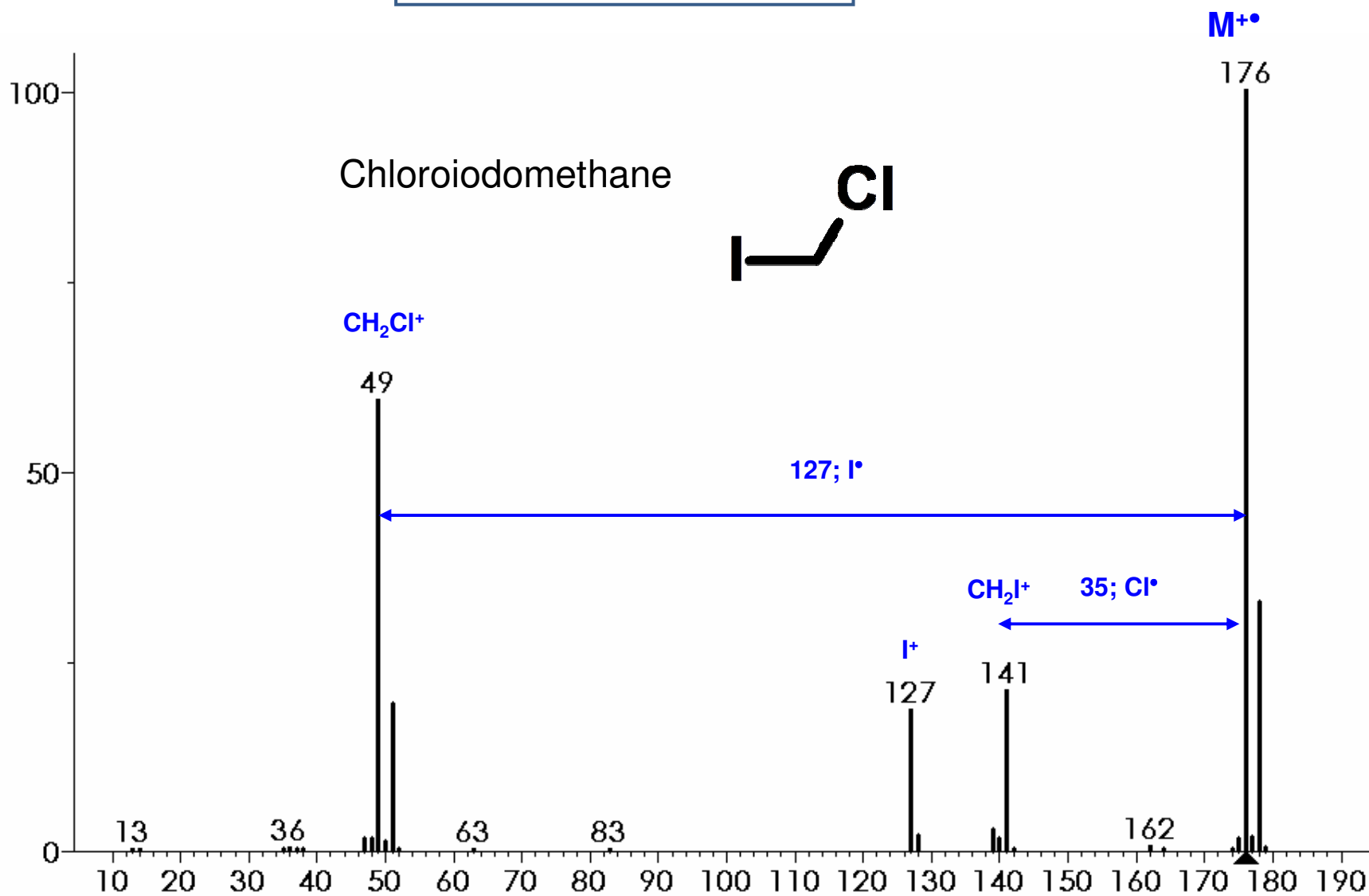


Unknown



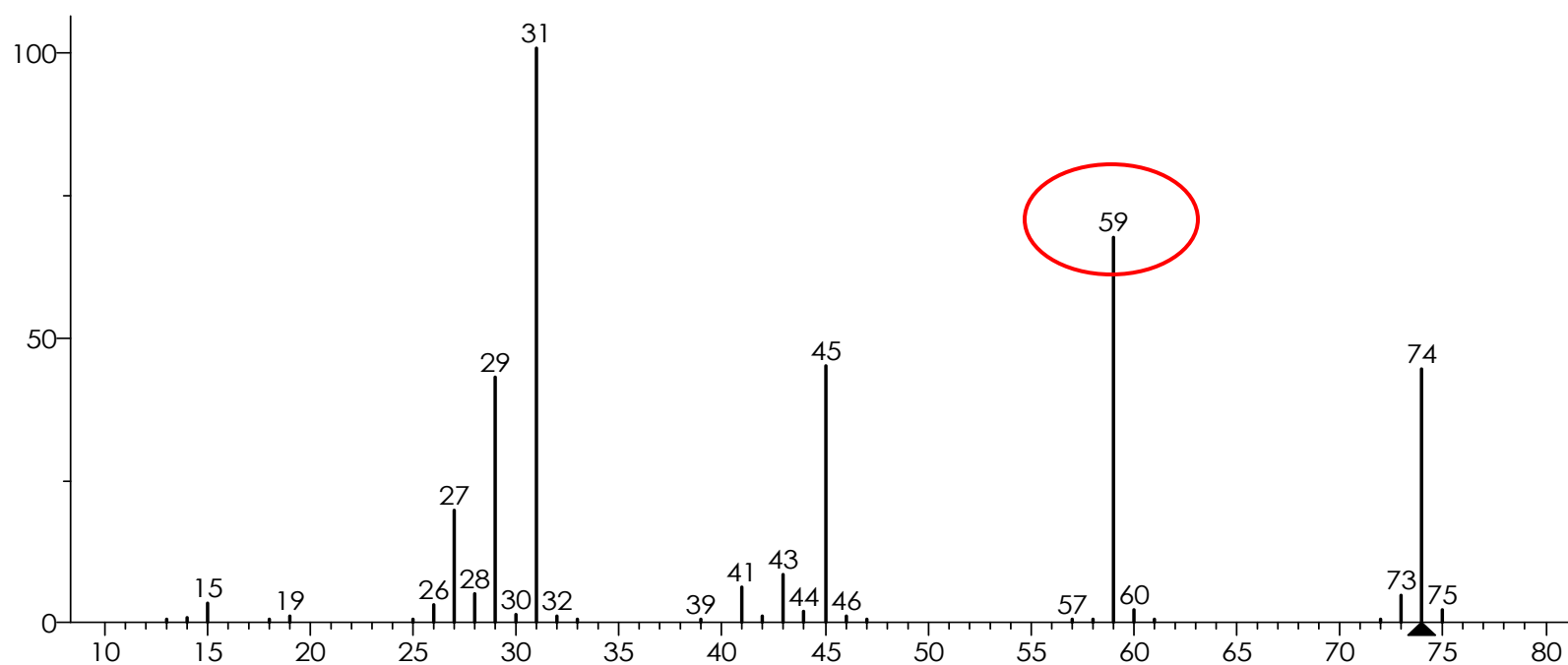
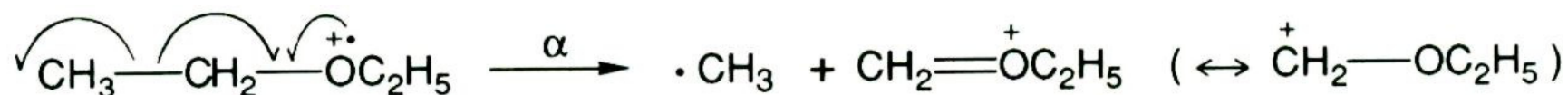
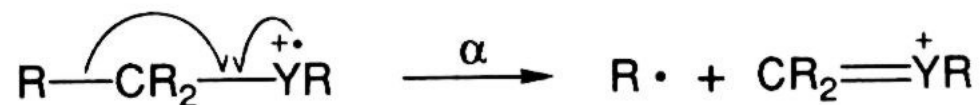
What is the structure?

EI



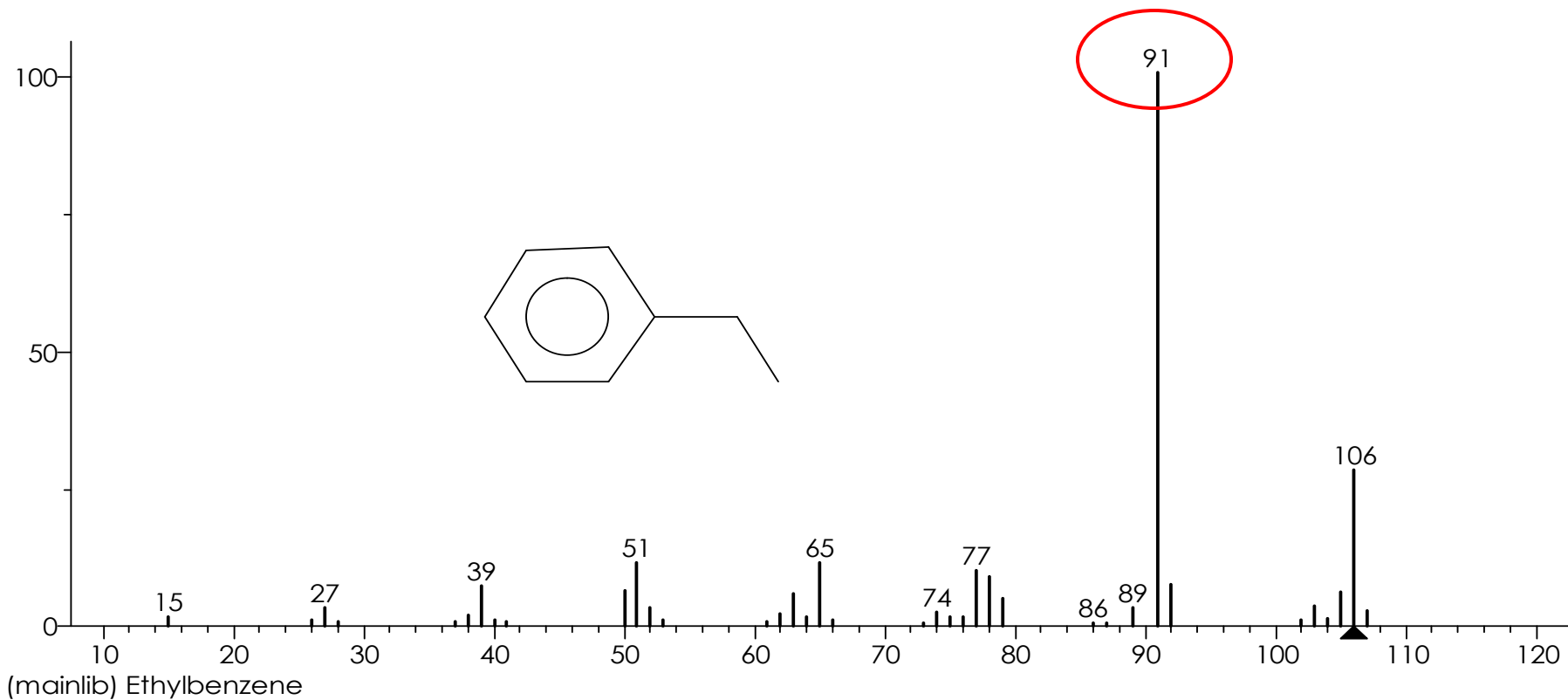
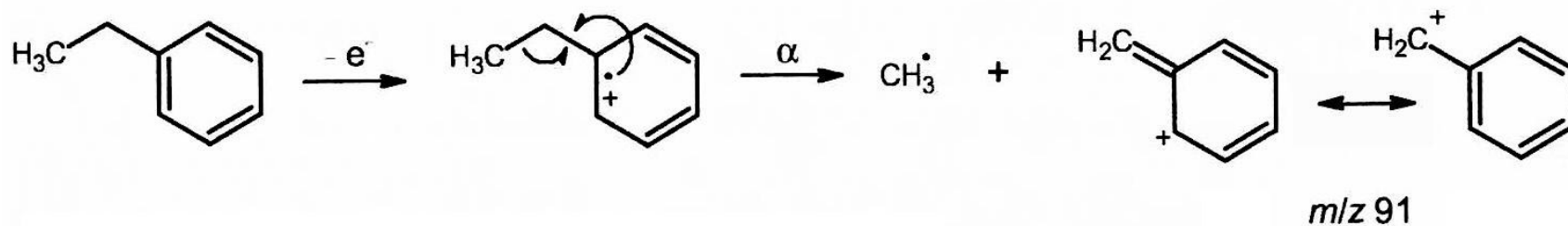
α -cleavage: fragmentation initiated by radical site

- cleavage induced by a strong tendency of electrons to form pairs – the odd electron is provided for the creation of a new bond; the neighboring bond is cleaved



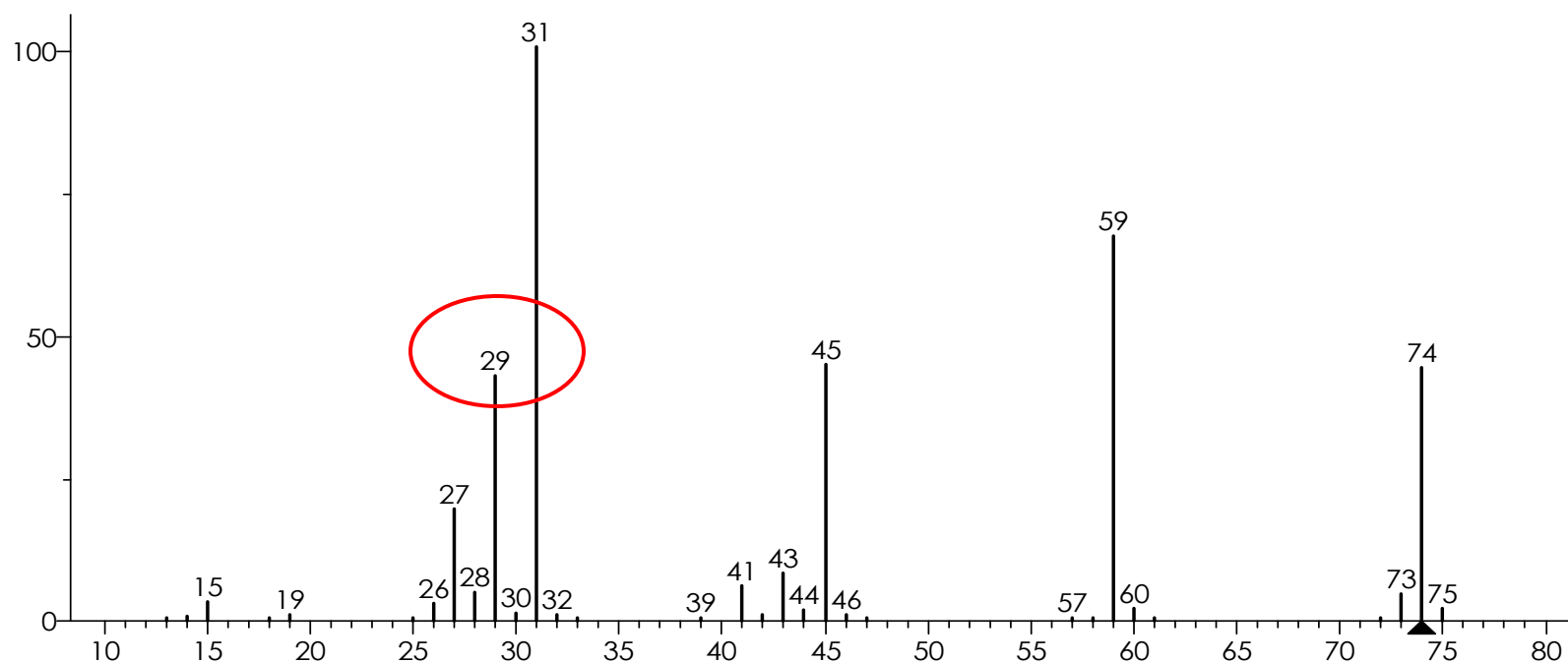
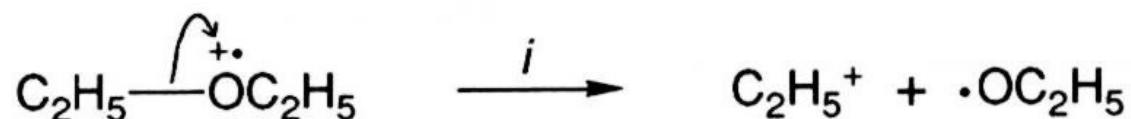
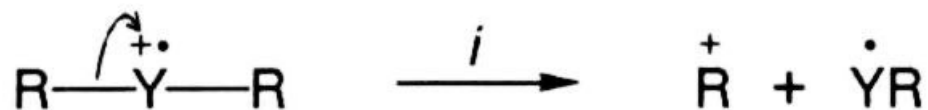
(mainlib) Ethyl ether

α -cleavage: benzylic cleavage



Inductive cleavage: fragmentation caused by a charge

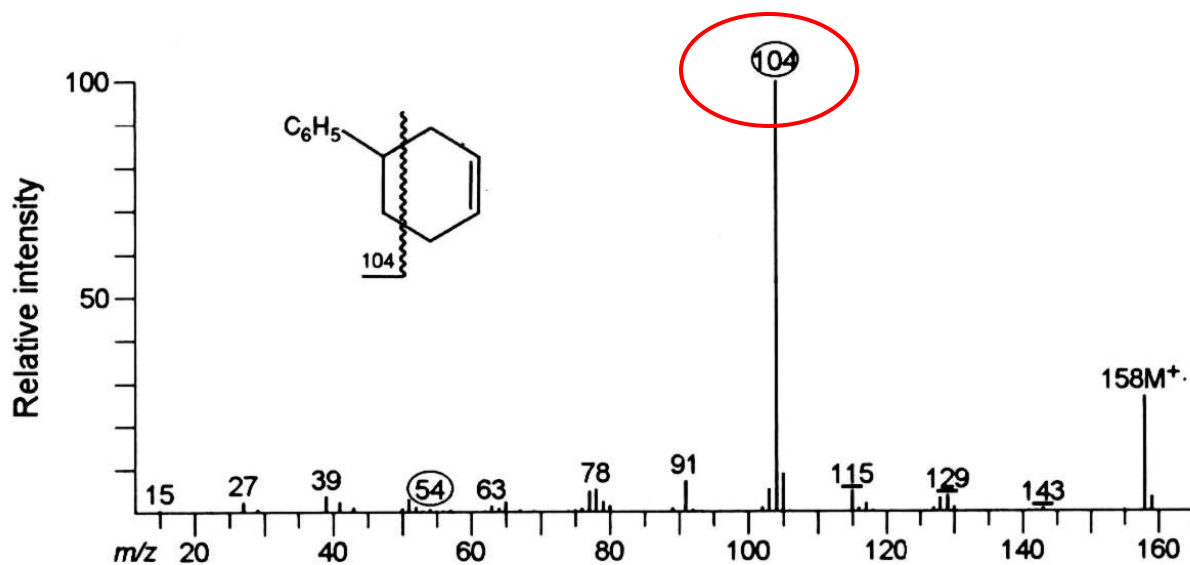
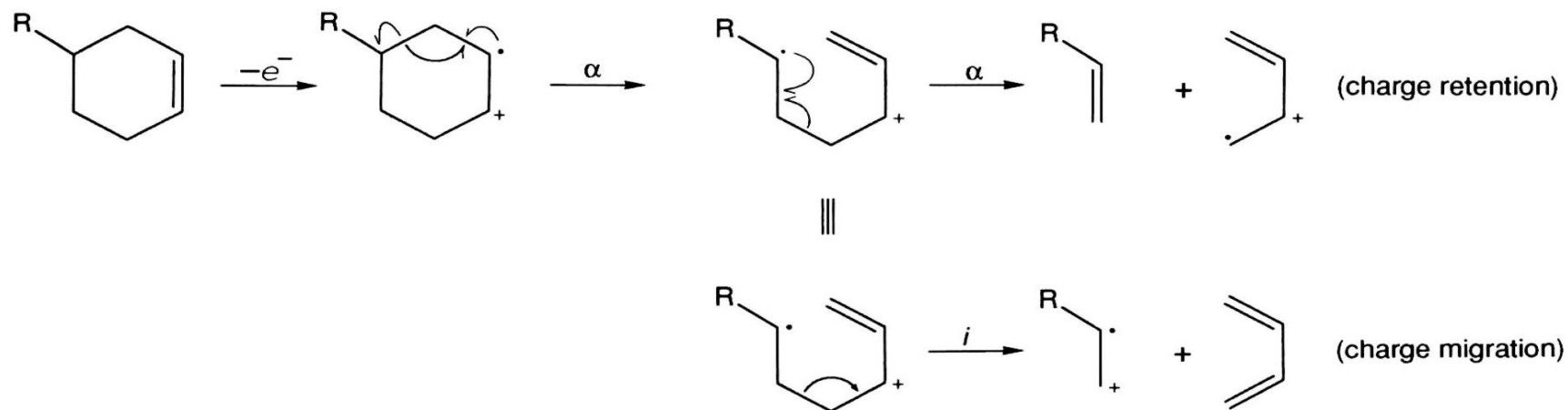
- cleavage initialized by attraction of an electron pair by the charge



(mainlib) Ethyl ether

Fragmentation of cyclic structures – retro Diels-Alder

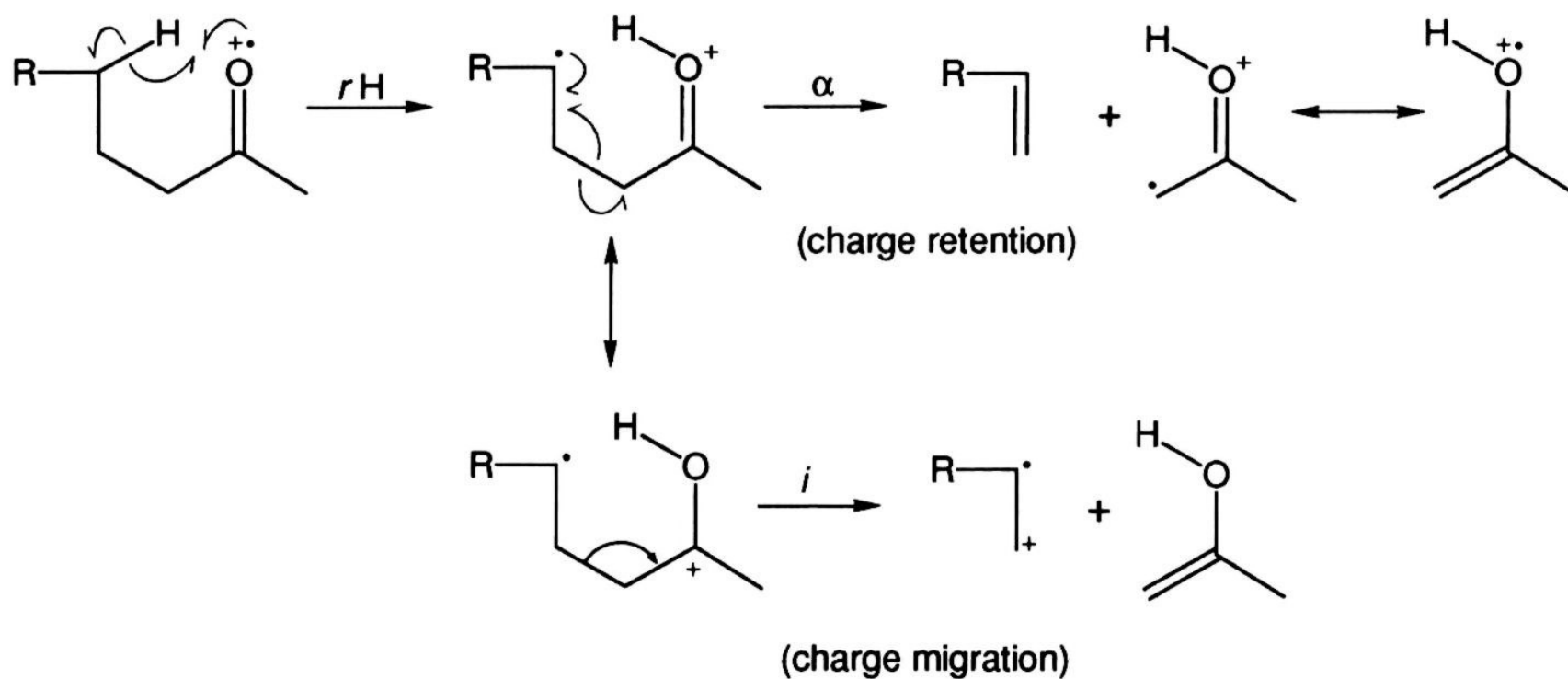
- π -electrons of double bonds in the cyclic structures are the primary site of ionization



4-fenylokyklohexen

Hydrogen rearrangement – McLafferty rearrangement

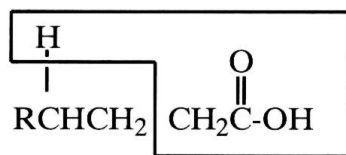
- rearrangement of γ -hydrogen on an unsaturated group over a 6-membered ring. The new radical site **Nové radikálové místo** initializes α -cleavage.



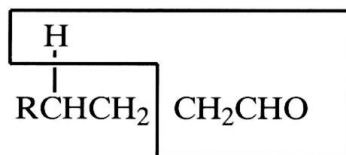
Hydrogen rearrangement – McLafferty rearrangement

- The $OE^{+\bullet}$ fragments are typical for many functional groups - aldehydes, ketones, esters, acids, amides, carbonates, phosphonates, etc.

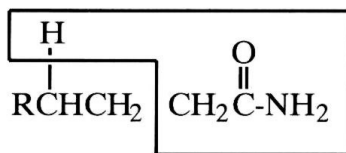
Acids

 m/z 60

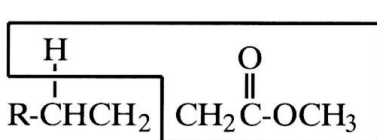
Aldehydes

 m/z 44

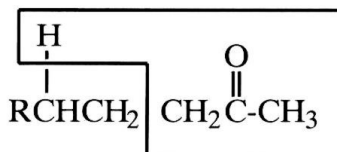
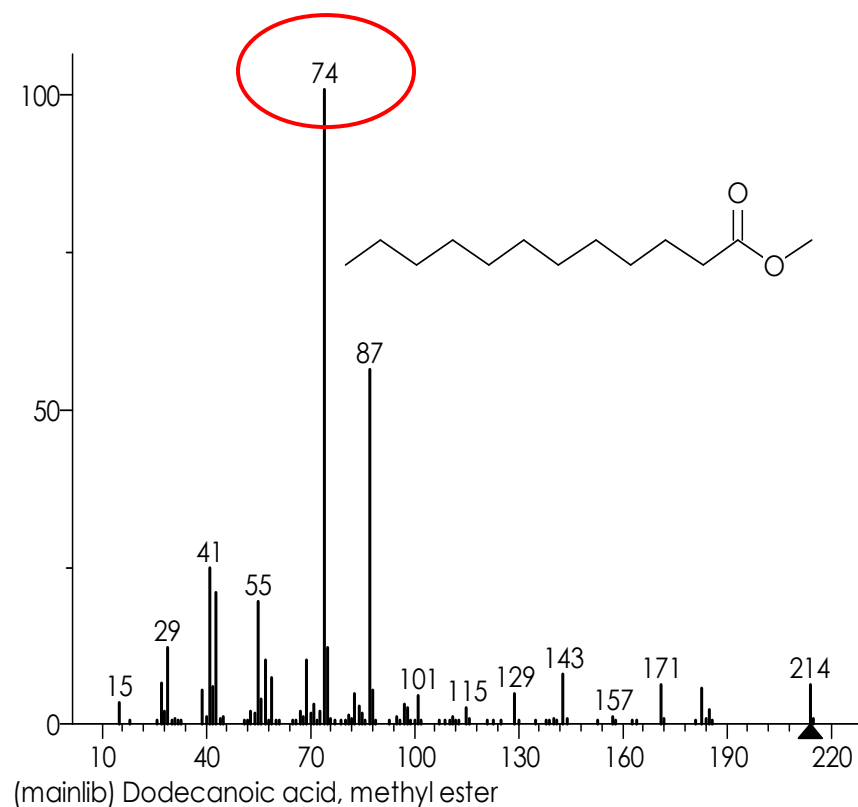
Amides

 m/z 59

Esters

 m/z 74

Ketones

 m/z 58

Characteristic ion series

Homologous series in the low m/z range provide information on structural elements in the molecule.

Ionty	Série	Funkční skupiny
m/z 29, 43, 57, 71, 85, 99, ...	$C_nH_{2n+1}^+$	alkyl („alifatika“)
m/z 31, 45, 59, 73, 87, ...	$C_nH_{2n+1}O^+$	alkoholy, ethery
m/z 33, 47, 61, 75, 89, ...	$C_nH_{2n+1}S^+$	thioly, sulfidy
m/z 30, 44, 58, 72, 86, ...	$C_nH_{2n+2}N^+$	aminy
m/z 29, 43, 57, 71, 85, 99, ...	$C_nH_{2n-1}O^+$	aldehydy, ketony
m/z 45, 59, 73, 87, ...	$C_nH_{2n-1}O_2^+$	kyseliny, estery
m/z 40, 54, 68, 82, 96, ...	$C_nH_{2n-2}N^+$	nitrily
m/z 38, 39, 50-52, 63-65, 75-78, 89-92	-	aromatika

Characteristic ions

m/z	Ion
19	F^+ , H_3O^+
20	HF^+
30	$CH_2NH_2^+$, indikuje aminy
31	indikuje CH_3O- nebo $-CH_2OH$
33,34	HS^+ , H_2S^+
35,36,37,38	Cl^+ , HCl^+
46	NO_2^+ , indikuje nitrosloučeniny
47	CCl^+ , $HC(OH)_2^+$, CH_3S^+
61	$CH_3C(OH)_2^+$ indikuje „dlouhé“ estery kyseliny octové
73	$(CH_3)_3Si^+$, CH_5Si^+
77	fenyl (doprovázen m/z 51 a 50)
105	benzoyl (pokud doprovázen m/z 77)

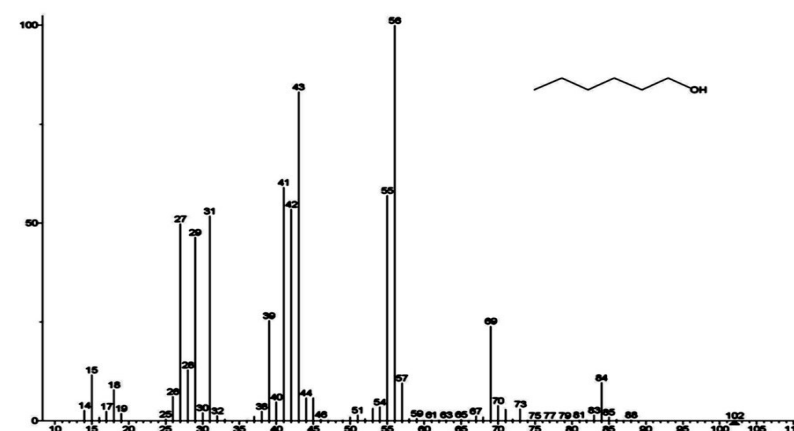
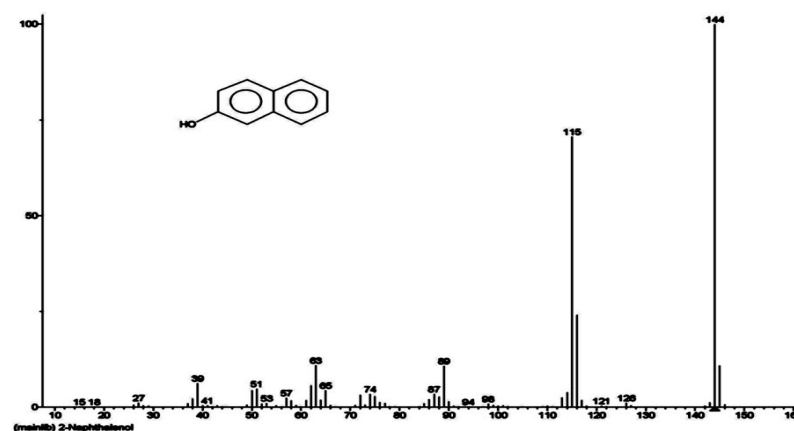
Logical neutral losses

Neutral losses of radicals and neutral molecules must make chemical sense.

Ztráta (u)	Logická ztráta ?	Složení
1	ano	H
2	ano	H ₂
3 - 14	NE	-
15	ano	CH ₃
16	ano	NH ₂ , O
17	ano	OH, NH ₃
18	ano	H ₂ O
19	ano	F
20	ano	HF
21 - 25	NE	-
26	ano	C ₂ H ₂ , CN
27	ano	C ₂ H ₃ , HCN
28	ano	C ₂ H ₄ , CO
29	ano	HCO, C ₂ H ₅ , CH ₃ N
30	ano	CH ₂ O
31	ano	CH ₃ O
32	ano	CH ₃ OH, S
33	ano	SH
34	ano	H ₂ S
35	ano	Cl
36	ano	HCl
37 - 40	NE	-
41 a výše	ano	homologické ztráty

The molecular ion intensity

The molecular ion intensity is related to its stability. The intensity suggests the presence of certain structural elements in the molecule.

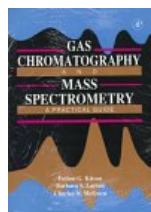


Literature

EI



Fred W. McLafferty and Frantisek Turecek: Interpretation of Mass Spectra. University Science Books (1993). ISBN-10: 0935702253, ISBN-13: 978-0935702255

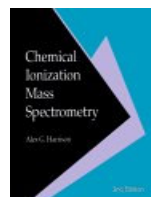


Fulton G. Kitson, Barbara S. Larsen, and Charles N. McEwen: Gas Chromatography and Mass Spectrometry. Academic Press (1996). ISBN-10: 0124833853, ISBN-13: 978-0124833852

ESI
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Levsen et al.: Even-electron ions: a systematic study of the neutral species lost in the dissociation of quasi-molecular ions. J. Mass Spectrom. 42, 1024 – 1044, 2007



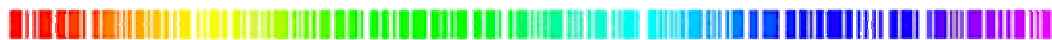
Alex. G. Harrison: Chemical Ionization Mass Spectrometry, CRC(1992). ISBN-10: 0849342546, ISBN-13: 978-0849342547

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Thank you for your attention !
