

MS SERVICE OF SMALL MOLECULES



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MS SERVICE WHICH WE PROVIDE

- a confirmation of a molecular structure by determination of exact mass and assigning a suitable formula
- MALDI of high molecular compounds such as peptides, proteins, oligonucleotides and some polymers
- a help with an identification of a structure according to fragmentation by EI or CI and measuring MS/MS analysis
- LC-MS analysis providing a previous agreement
- GC-MS and LC-MS self-service - after training and making a schedule

THE MASS SPECTROMETRY SERVICE

Where can you find us and your results ?

- the basement, number of the door 33.

- results and MS spectra on Novell
I:/MISC/MS/DATA/your surname

What should you do ?

- enrol at the sample book and bring a complete form



SAMPLE BOOK

sequence number	date	name	team code	sample label	method (source)	date of measuring	measured by
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- each sample has to be written under its own sequence number separately

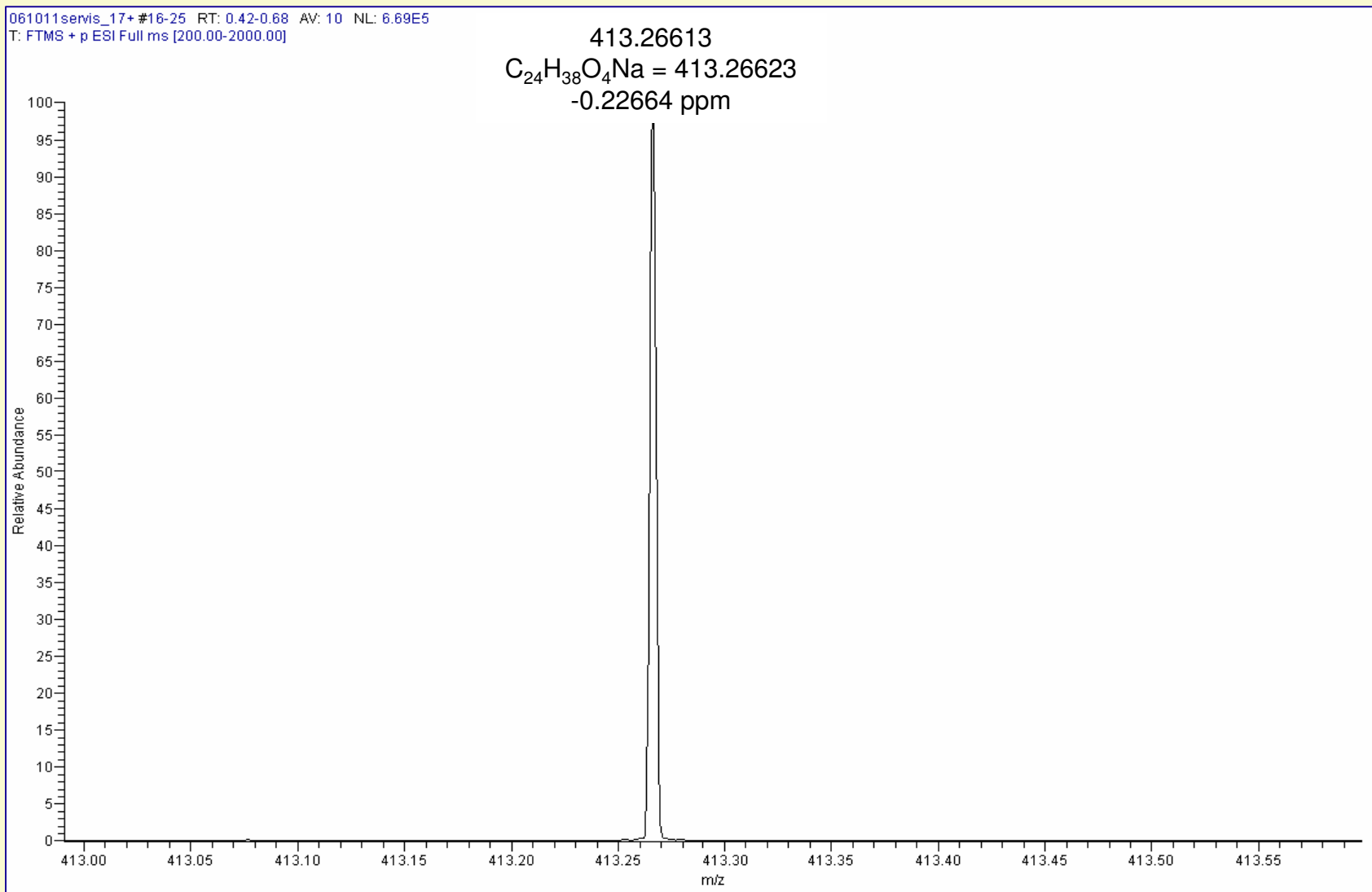


Skupina hmotnostní spektrometrie
Ústav organické chemie a biochemie AV ČR
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Tel.: 220183302, -481, -303 Fax: 220183583

ŽÁDANKA O MS ANALÝZU - SAMPLE SUBMISSION FORM

Jméno / Name:	Označení vzorku / Sample label:		
Pracoviště / Department:	Tel. linka / Phone:	Pořadové číslo ! / Sequence number !:	Datum / Date:
Navrhovaný způsob ionizace / Suggested ionisation technique: <input type="checkbox"/> GC/EI or CI <input type="checkbox"/> Direct probe EI <input type="checkbox"/> ESI <input type="checkbox"/> APCI <input type="checkbox"/> MALDI			
MS analýza / MS analysis: <input type="checkbox"/> LR - Hmotnostní spektrum - jednotkové rozlišení / Mass spectrum - unit resolution <input type="checkbox"/> HR - Přesná hmotnost - vysoké rozlišení / Accurate mass - high resolution <input type="checkbox"/> Peptidické mapování / Peptide mass fingerprinting (Please contact MS staff)			
Předpokládaná struktura / Assumed structure:	Sumární vzorec / formula:		
	Monoizotopická hmotnost / Monoisotopic mass:		
	Rozpouštědla / Solvents: <input type="checkbox"/> H ₂ O <input type="checkbox"/> MeOH <input type="checkbox"/> ACN <input type="checkbox"/> ACN/H ₂ O+FA <input type="checkbox"/> CHCl ₃ <input type="checkbox"/> Et ₂ O <input type="checkbox"/> Hexan <input type="checkbox"/> Aceton		
	Stabilita v kyselinách / Stability in acids: <input type="checkbox"/> Ano / Yes <input type="checkbox"/> Ne / No		
	Stabilita v bázích / Stability in bases: <input type="checkbox"/> Ano / Yes <input type="checkbox"/> Ne / No		
Poznámky / Notes:			

HOW DOES THE LR and HR ESI SPECTRA LOOK LIKE



HOW DOES THE LR and HR EI SPECTRA LOOK LIKE

Elemental Composition Report

Single Mass Analysis (displaying only valid results)

Tolerance = 10.0 PPM / DBE: min = -100.0, max = 1000.0

Element prediction: Off

Monoisotopic Mass, Odd and Even Electron Ions

17 formula(e) evaluated with 1 results within limits (up to 50 closest results for each mass)

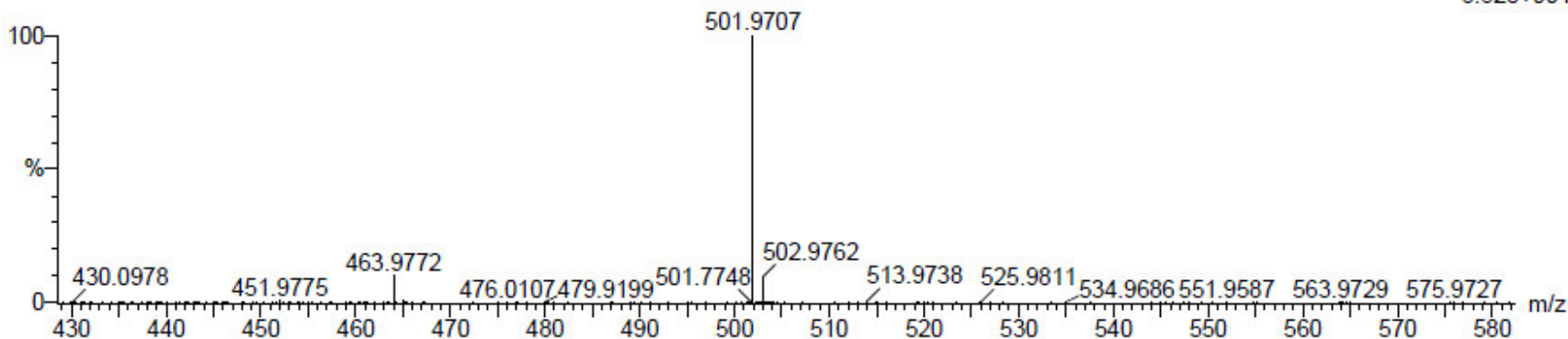
Elements Used:

C: 3-15 H: 0-1 N: 1-1 F: 0-30

heptacosia

heptacosia 162 (2.701) Cm (160:191)

TOF MS EI+
8.82e+004



Minimum: -100.0
Maximum: 5.0 10.0 1000.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Formula
501.9707	501.9711	-0.4	-0.8	0.5	3495.1	C9 N F20

SAMPLES

Notes

- if the samples are either **toxic, unstable or moisture, light and high temperature sensitive** you have to put it down in the form.
- if you have **a small amount of sample**, you can certainly get it back, providing it is written in the form.

Solid samples:

It is better to submit **a solid sample** (100 µg - 1.0 mg). Please do not forget mention a suitable solvent.

Desolved samples:

Please use only solvents for LC/MS or distilled solvents.

- samples should be **fully dissolved**.

All instruments are usually capable of detecting at least 1 µg/mL.

However, it is good habit to submit around 1 mg/mL.

RECOMMENDED SOLVENTS

Which solvent can I use? - it depends on the type of ionization and whether you want to use GC chromatography before MS or not.

- **ESI:** the most preferred solvents are **MeOH**, **H₂O** and **ACN** - a solution should be free from nonvolatile buffers and another additives.
CHCl₃ and **acetone** are also acceptable in mixture with MeOH or ACN.
- **APCI:** **MeOH**, **ACN**, **IPA**, **CHCl₃**, **EtAc**, **EtOH** and **acetone**
- **EI/CI with direct probe:** solid samples or dissolved in volatile solvents.
- **EI/CI coupled with GC:** hexan, ether, CHCl₃, CH₂Cl₂, MeOH, EtOH. The samples for GC/MS must be free from strong acid, bases and oxidizing compounds.
- **MALDI:** in a solution free from nonvolatile buffers, solvents and surfactants

! Please, avoid solvents with high boiling points (DMSO, DMF) !

USUAL CONTAMINANTS

- **Phthalates** - from plastics, contaminated solvents
diisobutylphthalate (masses EI: 149, 205, 223, 278, ESI: 279, 301)
diisooctylphthalate (masses EI: 149, 167, 279, 390, ESI: 391, 413)
- **Antioxidants** - from plastics : **irganox, irgafos**
(the most common masses EI: 316 ,591, 647, 642, ESI: 663, 685)
- **Polysiloxans** - from silicone rubber, teflon lined caps from vials
(the most common masses EI: 73, 147, 221, 295, 355, ESI: 297, 371, 445, 519)
- **PEGs** - extracted polymer from teflon/silicon septum (+44 series)
- **Amides** - from plastics - **oleamide** (ESI: 282), **stearamide** (ESI: 284), **erucamide** (ESI: 338)
- **Detergents** - **Triton X-100**
- **Fatty acids** - **palmitic and oleic acid** from skin (masses negESI : 255, 283)

SALTS, BUFFERS and ANOTHER ADDITIVES

- **volatile** - TFA, FA, acetic acid, ammonium acetate, ammonium formate etc.

- can be used in lower concentrations (up to 10mM)

- **nonvolatile** - phosphate, sulfate buffers, SDS, CHAPS, TRIS, HEPES etc.

- should not be used because of decreasing signal, salt clusters creation and unbearable contamination of mass spectrometers

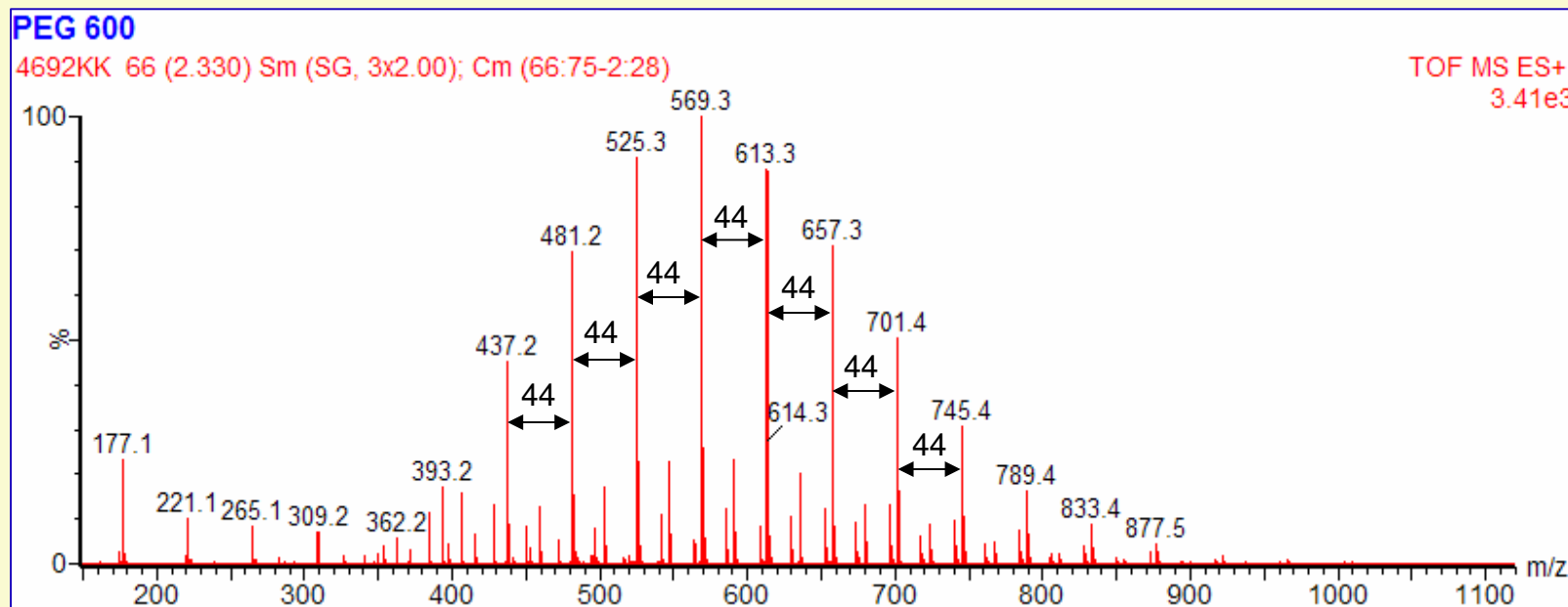
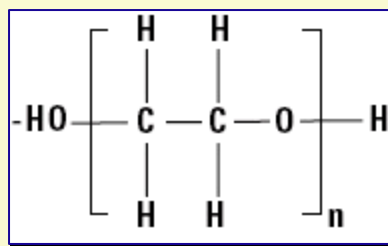
	MW	MALDI	MALDI	ESI	ESI
Surfactant/ buffer/ salt	(g/mol)	(mM)	(wt.%)	(mM)	(wt.%)
TRIS	121	100	1	n.a.	n.a.
HEPES	238	100	2,4	n.a.	n.a.
Urea	60	500	3	n.a.	n.a.
Dithiotreitol	154	500	7,7	n.a.	n.a.
Guanidine	96	250	2,4	n.a.	n.a.
Glycerol	92	130	1,2	n.a.	n.a.
Triton X-100	628	1,6	0,1	1,6	0,1
Tween20	1228	n.a.	n.a.	0,81	0,1
SDS	288	0,35	0,01	0,34	0,01
CHAPS	615	0,16	0,01	1,6	0,1
Sodium Azide	65	15	0,1	3,1	0,02
NaCl	58	50	0,29	n.a.	n.a.
Sodium Acetate	82	50	0,41	n.a.	n.a.
TFA	114	n.a.	n.a.	4,4	0,05
NaHPO ₄	120	10	0,12	10	0,12

<http://masspec.scripps.edu/services/proteomics/saltol.php>

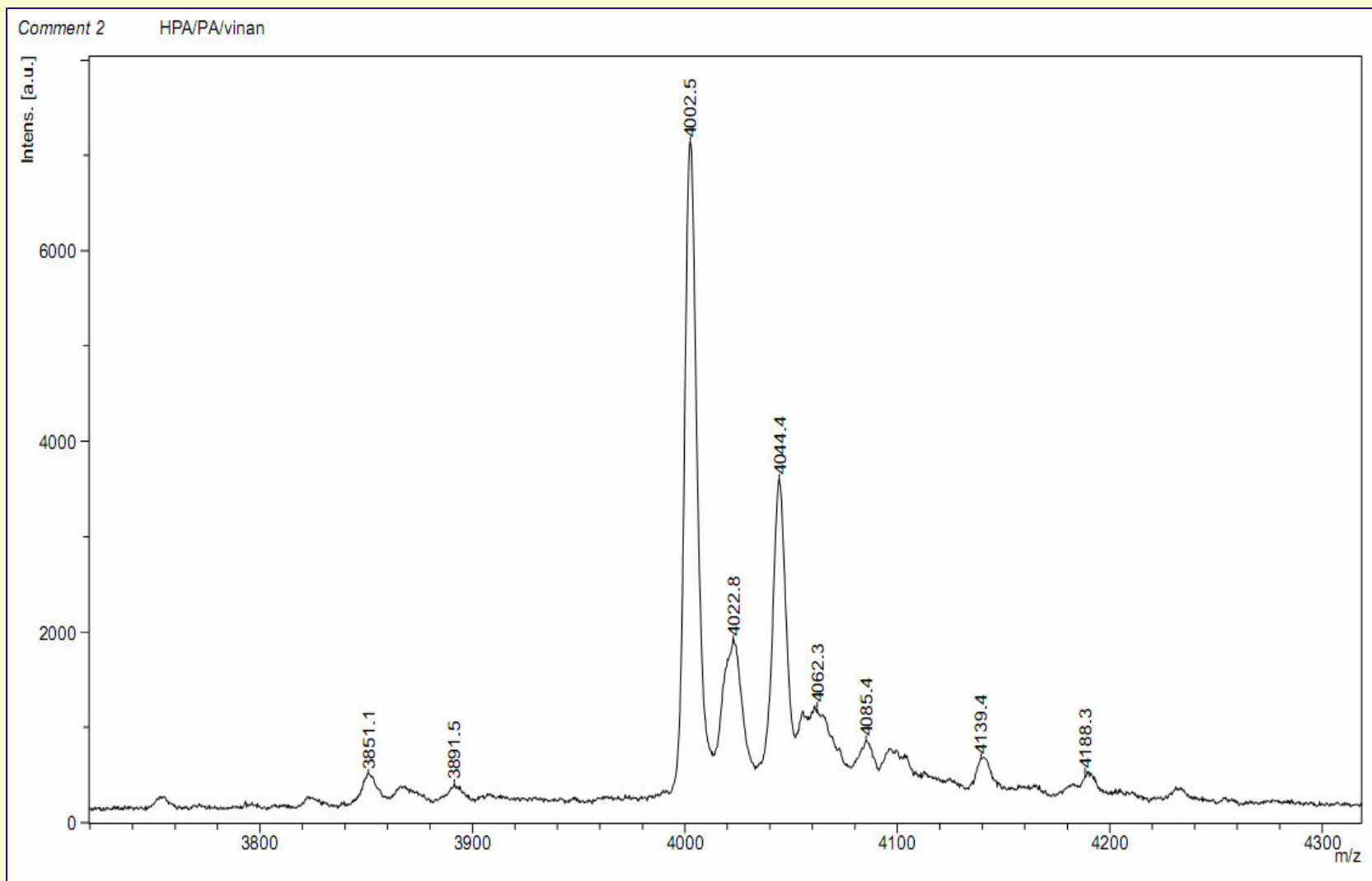
ADDUCTS AND CLUSTER PEAKS IN +/-ESI

- +23n....sodium adducts....+ESI
- +32n....methanol adducts....+ESI
- +39n....potassium adducts....+ESI
- +41n....acetonitrile adducts....+ESI
- +44n.....polyethylene glycol related (-CH₂CH₂O-)n....+ESI
- +53n.....ammonium chloride adducts (NH₄Cl)....+ESI
- +63n.....ammonium formate adducts (HCOONH₄)....+ESI
- +68n.... sodium formate adducts (HCOONa)....+/-ESI
- +74n.....polysiloxanes (Si(CH₃)₂O)....+/- ESI
- +77n.....ammonium acetate salts (CH₃COONH₄)....+ESI
- +82n.....sodium acetate adducts (CH₃COONa)....+/-ESI
- +114n ...TFA (trifluoroacetic acid) adducts (CF₃COOH)....-ESI
- +136n....sodium TFA (trifluoroacetic acid) adducts (CF₃COONa)....+/-ESI
- +288n ...SDS (sodium dodecylsulfate) adducts....-ESI

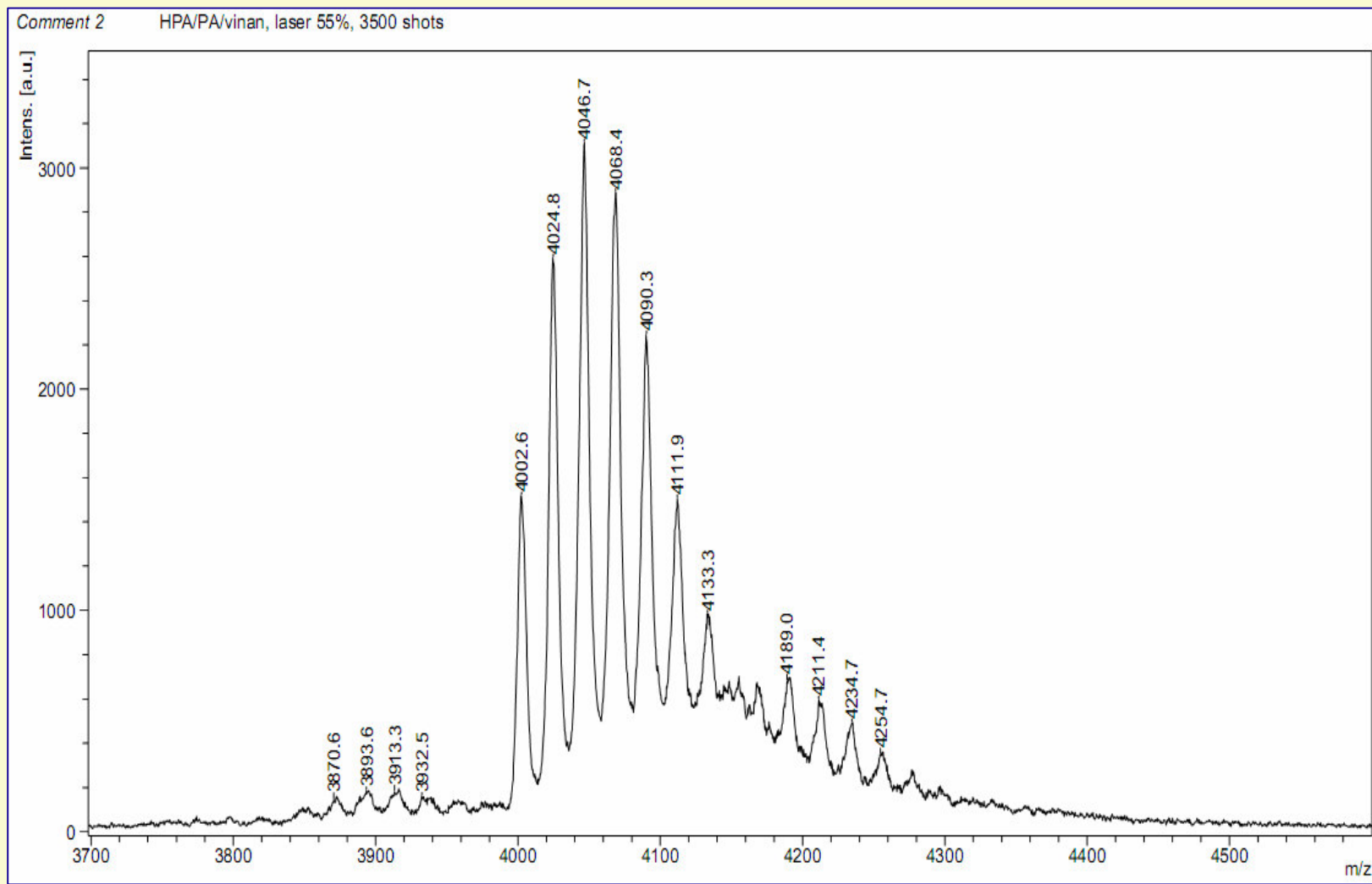
ESI SPECTRUM OF POLYETHYLENE GLYCOL 600



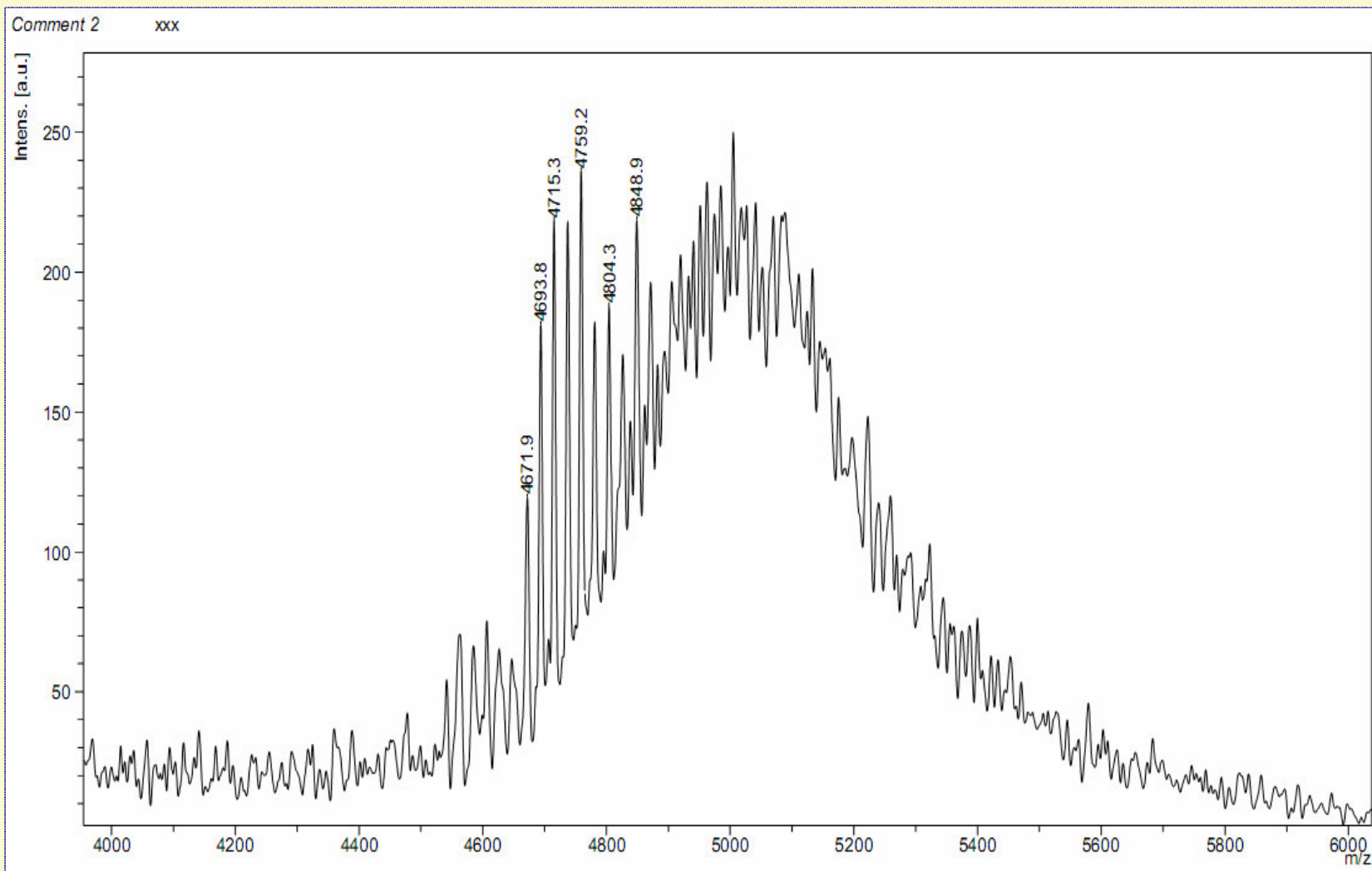
MALDI OF CLEAN SAMPLE



MALDI OF SAMPLE WHICH IS SLIGHTLY CONTAMINATED WITH SALTS



MALDI OF SAMPLE WHICH IS HIGHLY CONTAMINATED WITH SALTS

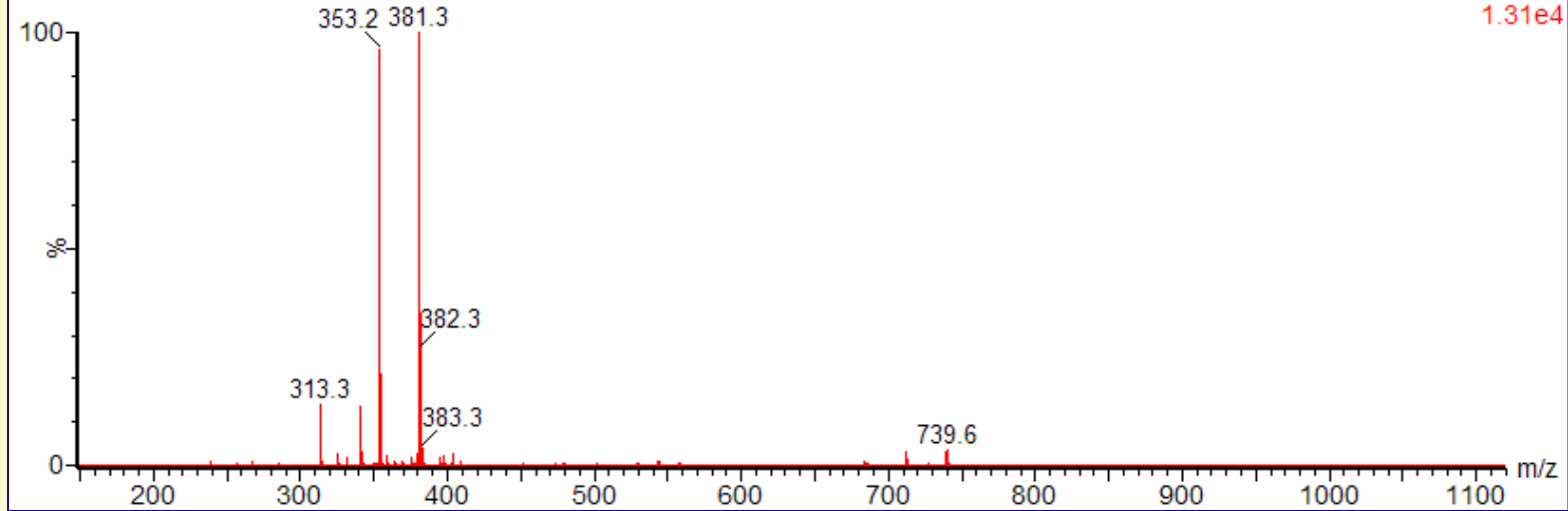


MeOH FROM EPPENDORFS

ORDINARY EPPENDORF

5119EM 34 (1.200) Sm (SG, 3x2.00); Cm (32:40-9:27)

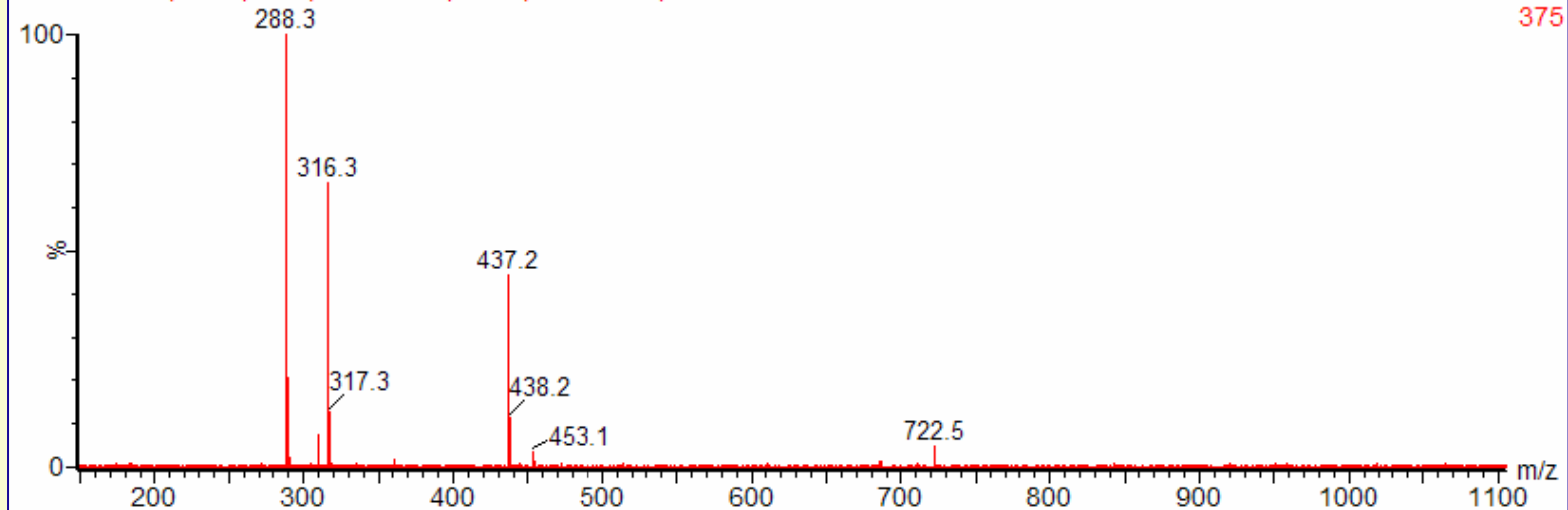
TOF MS ES+
1.31e4



MAXIMUM RECOVERY - EPPENDORF

5117EM 27 (0.953) Sm (SG, 3x2.00); Cm (22:27-1:18)

TOF MS ES+
375

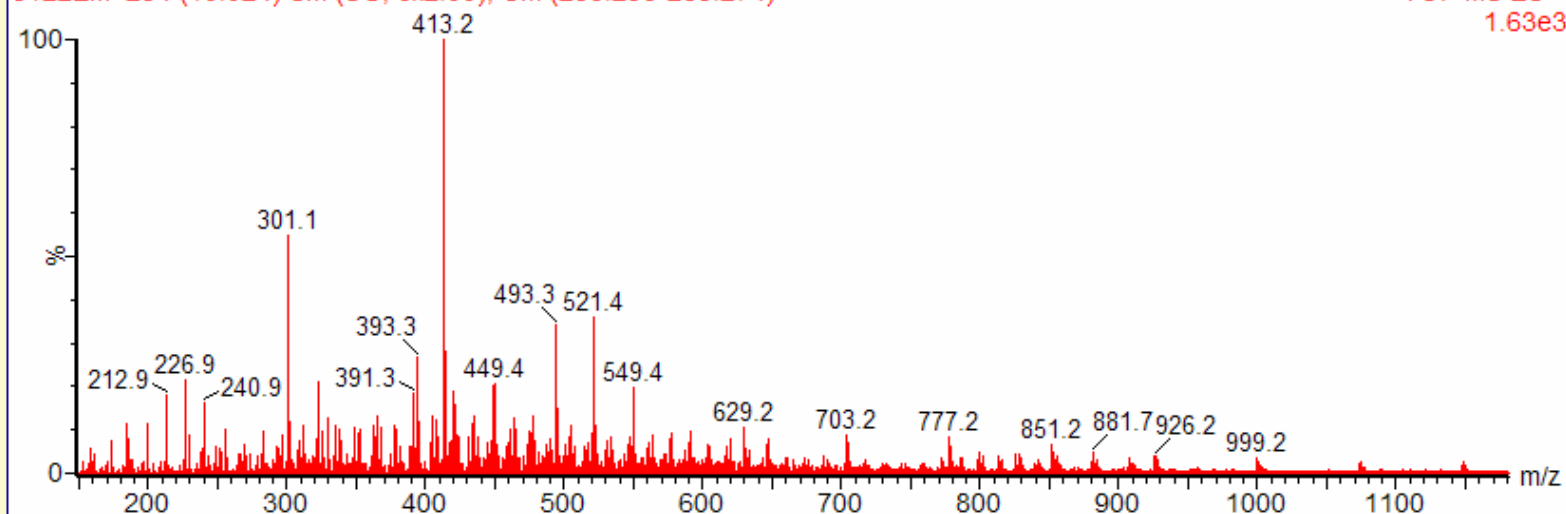


MeOH FROM GLASS VIALS

glass vial from our paper stock

5122EM 284 (10.024) Sm (SG, 3x2.00); Cm (283:293-265:274)

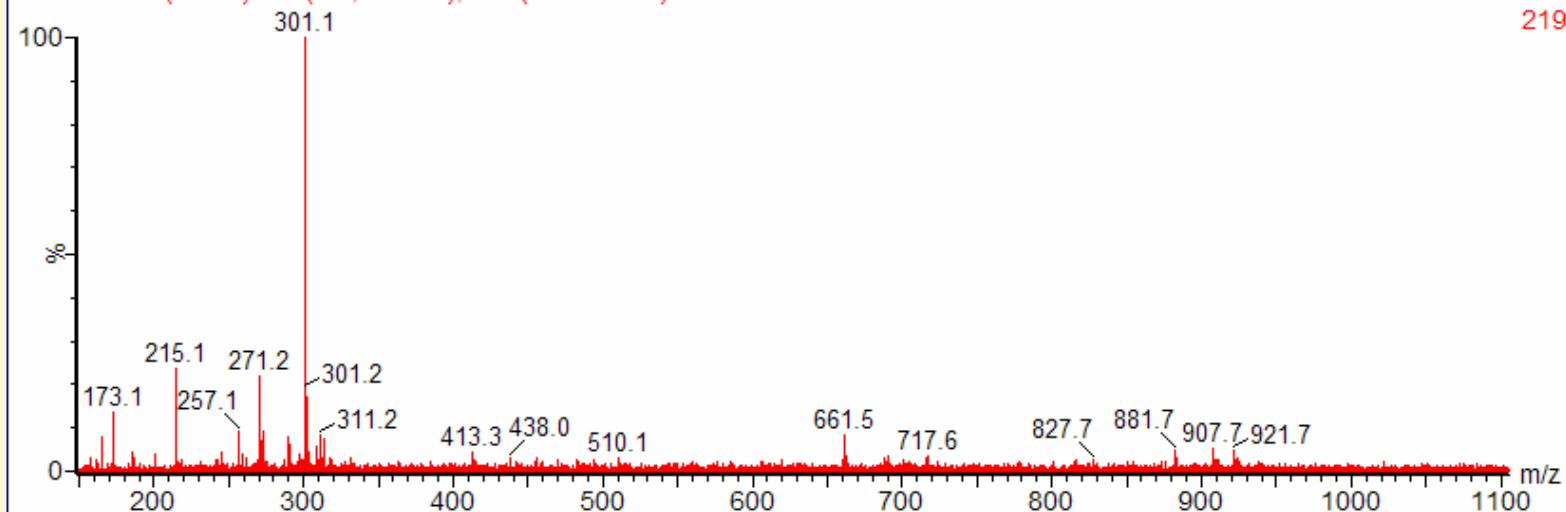
TOF MS ES+
1.63e3



GLASS VIAL - CHROMSERVIS

5116EM 23 (0.812) Sm (SG, 3x2.00); Cm (22:28-1:16)

TOF MS ES+
219



VIALS AND EPPENDORFS

Recommendation:

- glass vials + caps and septa PTFE/silicon, 2ml: *Chromservis s.r.o., P.N.C4000* for organic solvents
- eppendorfs MAXIMUM RECOVERY, 1,5 ml: *P-LAB a.s., P.N. U344701* for polar solvents

Thank you for your attention