

High Resolution LC-MS Beta-Agonist Screening Using TurboFlow/Exactive

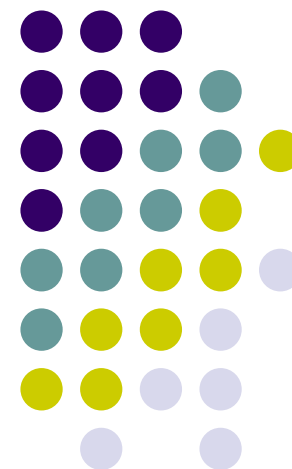
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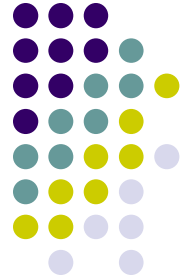


Joseph-Koenig-Strasse 40
D - 48147 Münster, Germany

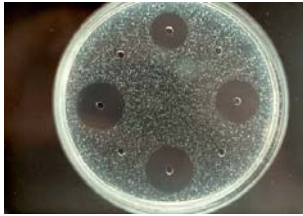
5th Thermo Scientific
High Resolution GC/MS Meeting on POPs
Barcelona, 29-30 April, 2010



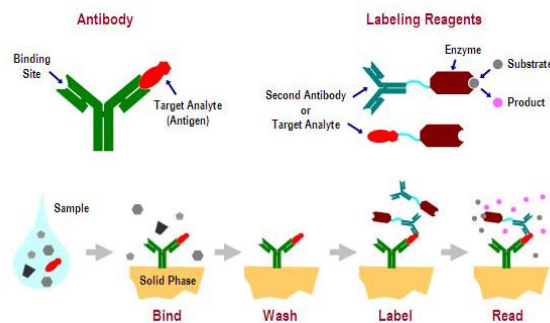
Screening methods



➤ Biological tests



➤ ELISA



➤ GC-MS



➤ LC-MS, LC-MS/MS



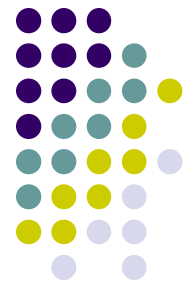
➤ LC-(Q-)TOF/MS



➤ LC-Orbitrap/MS



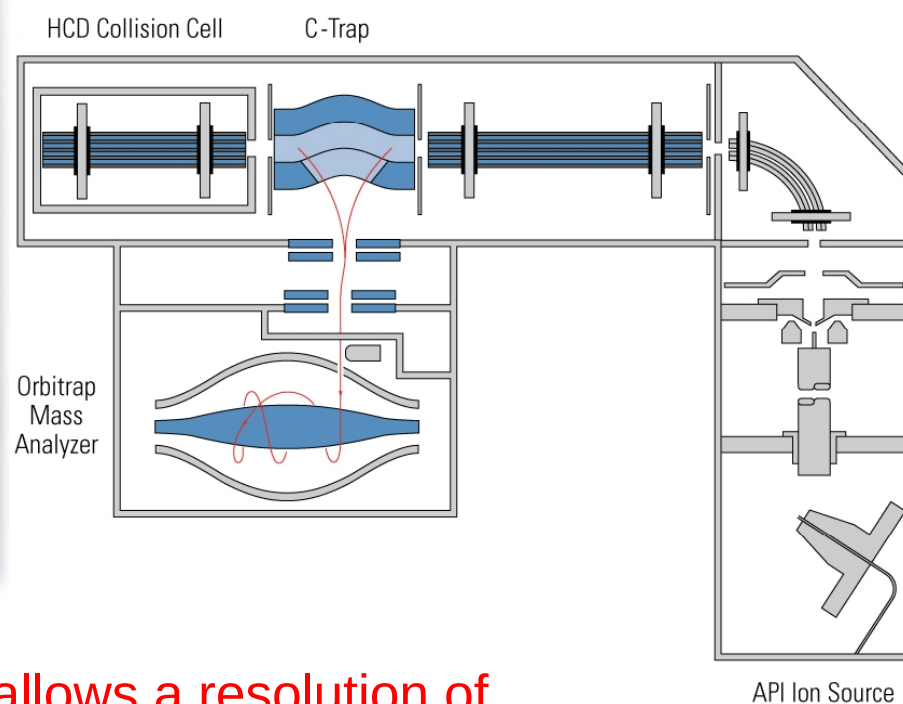
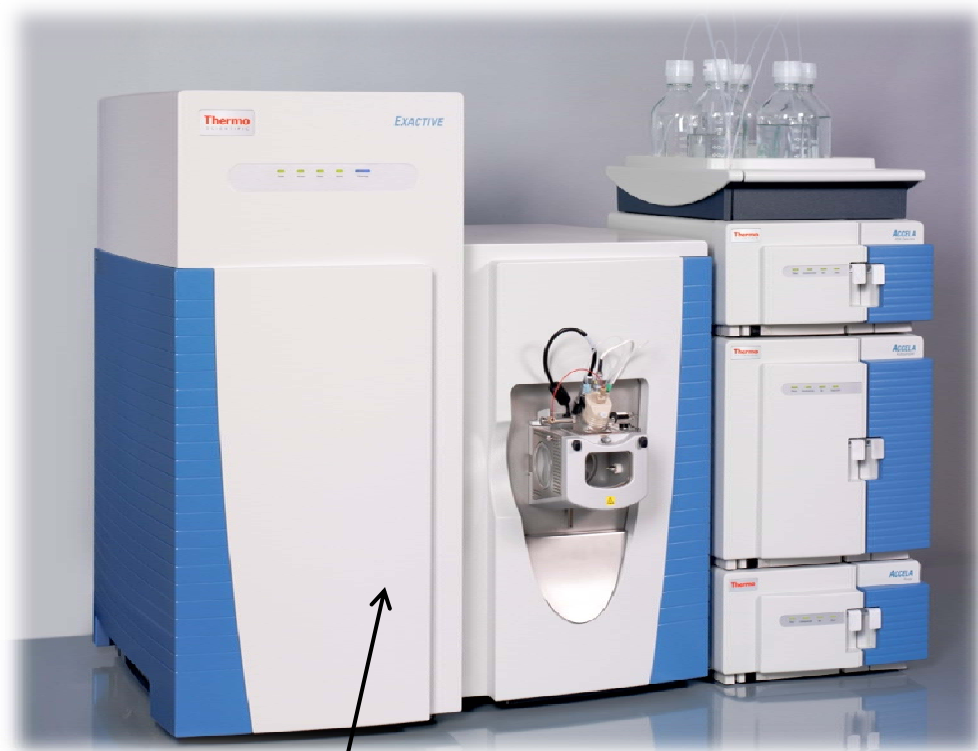
LC-MS/MS – a good screening technique?



- Advantages especially in MRM mode
- **However**, MRM is target analysis
("You see only what you want to see")
 - ⇒ Despite a large number of reference substances "other" compounds of interest which may occur in large concentrations can be easily overlooked
- For the optimization of the mass transitions and the accurate quantification the availability of the corresponding reference compounds is necessary

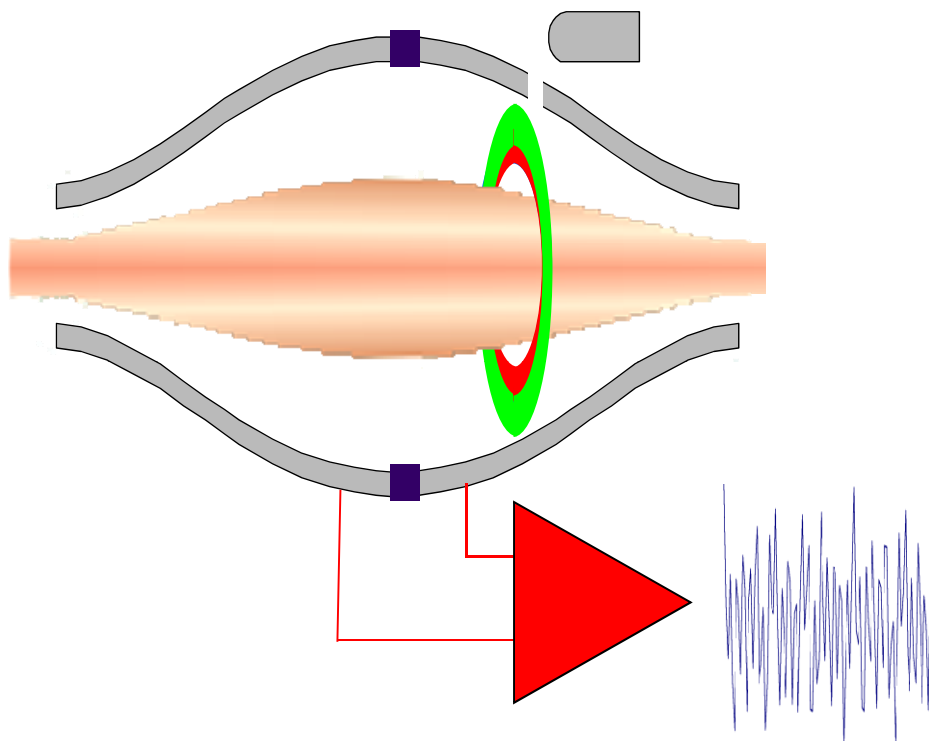


Exactive™ Bench-Top LC-MS-[MS]

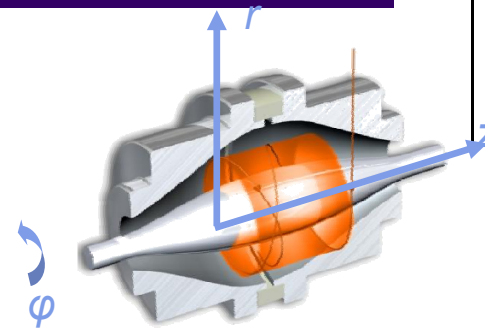


The Exactive allows a resolution of 100,000 with a mass accuracy <1ppm

Orbitrap – Principle of Operation



$$\omega_z = \sqrt{\frac{k}{m/q}}$$



Hyper-logarithmic potential distribution:
“ideal Kingdon trap”

$$U(r, z) = \frac{k}{2} \cdot \left\{ z^2 - r^2 / 2 + R_m^2 \cdot \ln(r / R_m) \right\}$$

Characteristic frequencies:

- Frequency of rotation ω_ϕ
- Frequency of radial oscillations ω_r
- Frequency of axial oscillations ω_z

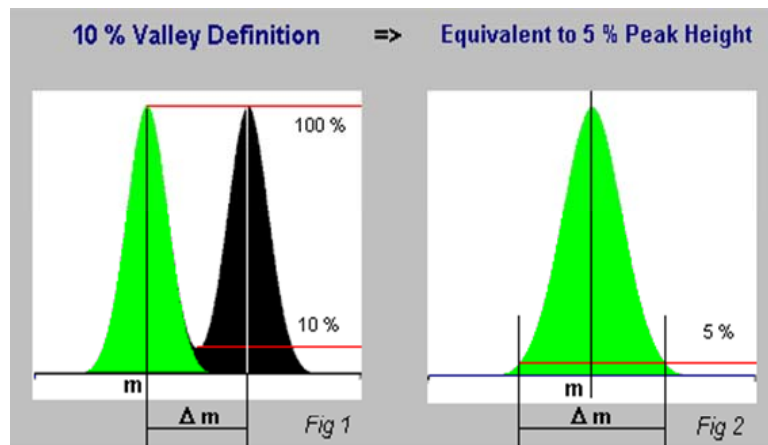
$$\omega_\phi = \frac{\omega_z}{\sqrt{2}} \sqrt{\left(\frac{R_m}{R}\right)^2 - 1} \quad \omega_r = \omega_z \sqrt{\left(\frac{R_m}{R}\right)^2 - 2}$$

Makarov A. *Anal. Chem.* 2000, 72, 1156-1162.

Courtesy of Thermo Scientific

Resolution Definitions

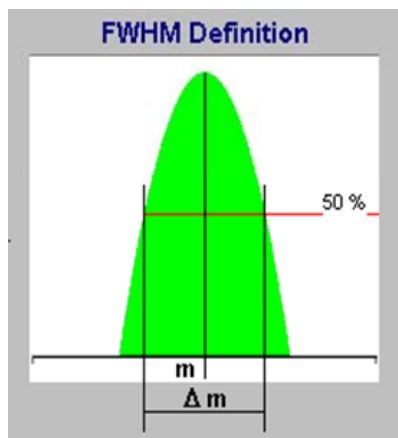
10% Valley



Magnetic Sector

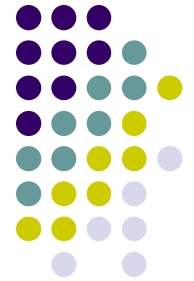
$$2 R_{FWHM} = R_{10\% \text{ valley}}$$

Full Width Half Maximum

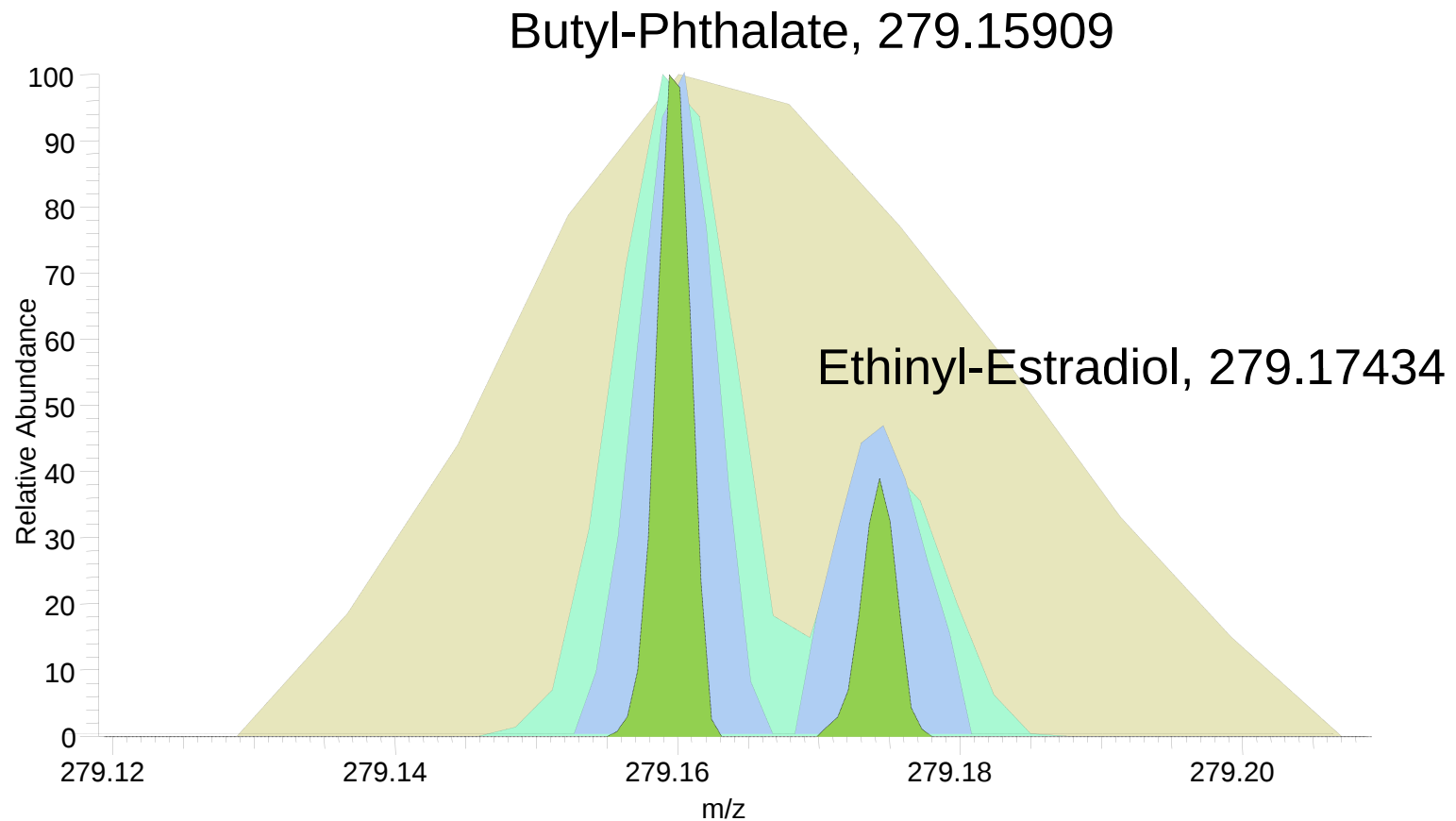


FT ICR
Orbitrap
ToF
Quadrupole

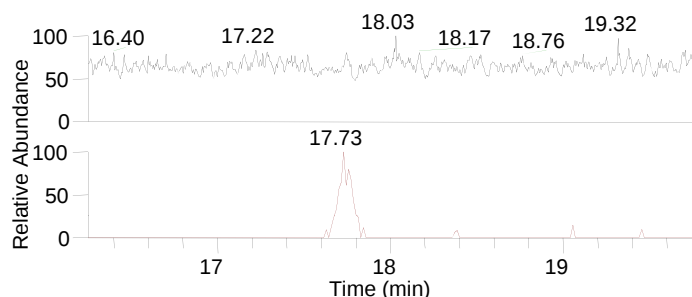
Influence of resolution on mass separation



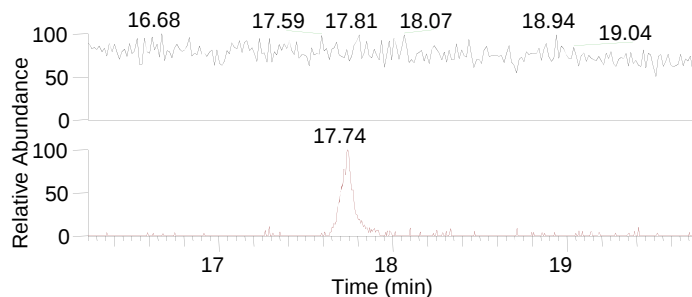
Resolution: 10k, 30k, 50k, 100k



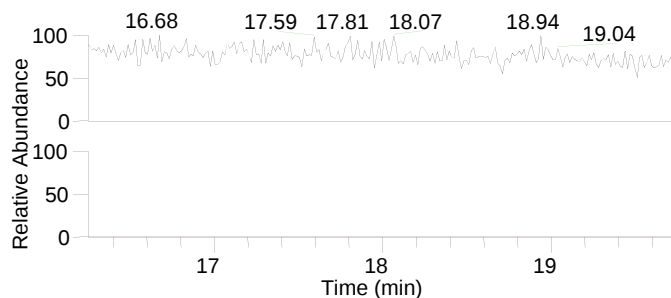
Ethinyl-Estradiol (Standard: 100 ng/ml)



RP = 100,000

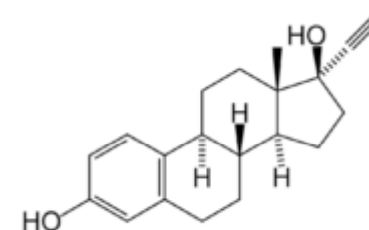


RP = 30,000



RP = 10,000

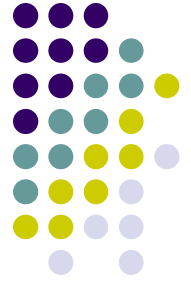
Ethinyl-Estradiol



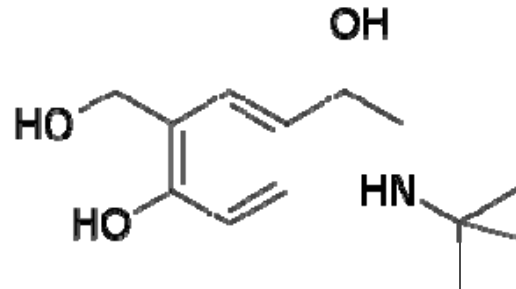
The isobaric phthalate background ion interferes with the Ethinyl-Estradiol ion.

At resolution of 10,000 the steroid mass is shifted because the isobaric ions are not resolved.

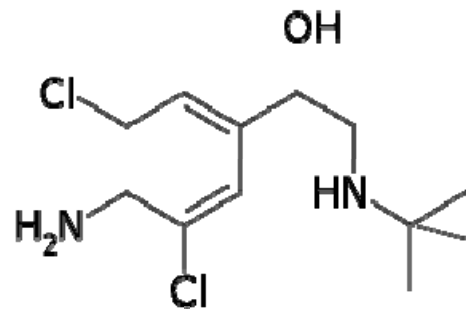
beta-agonists



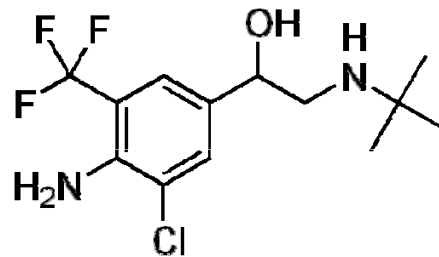
Salbutamol



Clenbuterol



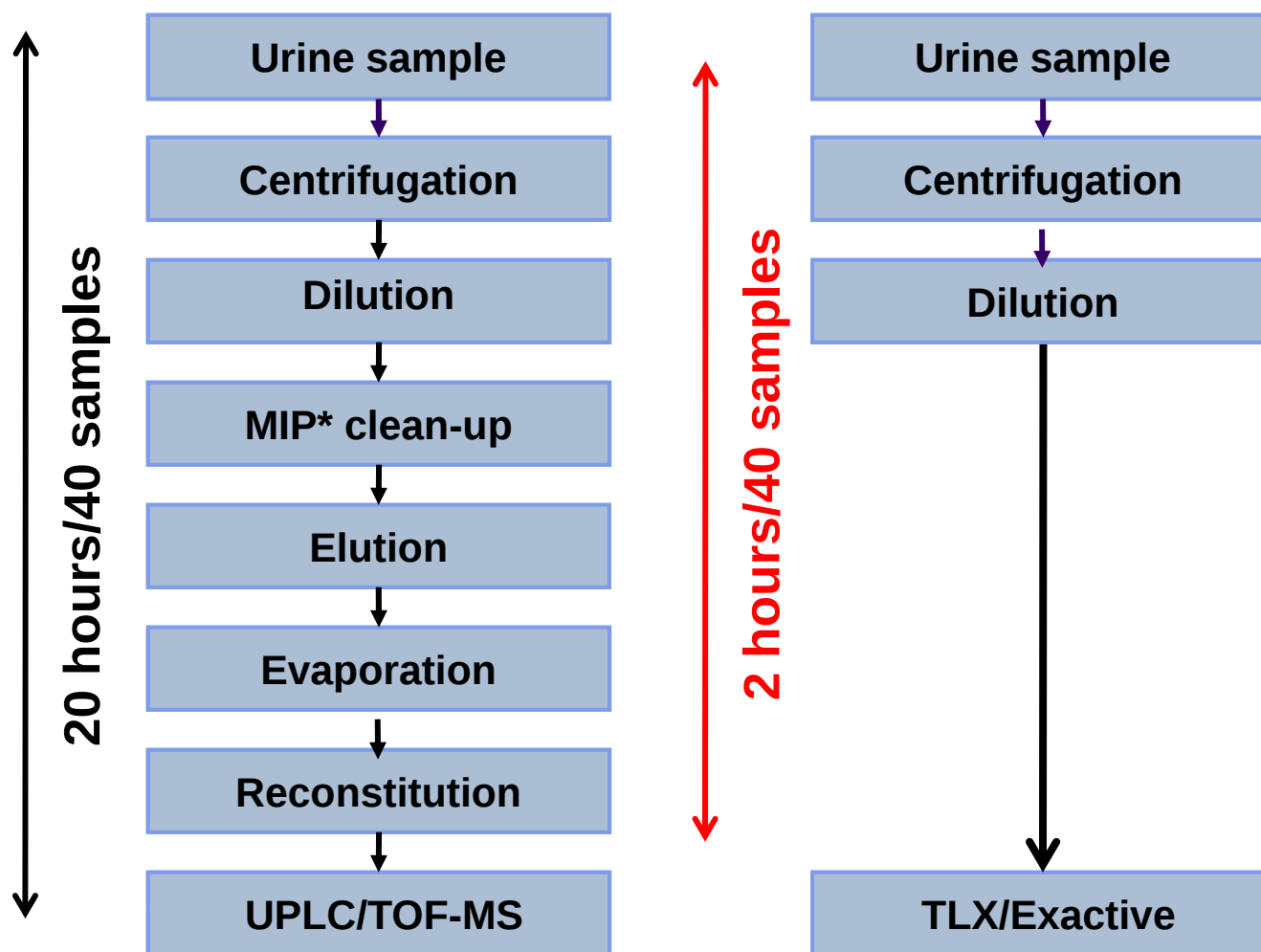
Mabuterol



- are a class of drugs used to treat asthma and other pulmonary diseases
- for growth-promoting purposes in farm animals (illegal)
- different chemical properties e.g. amphoteric
- 24 substances are mentioned



Sample preparation



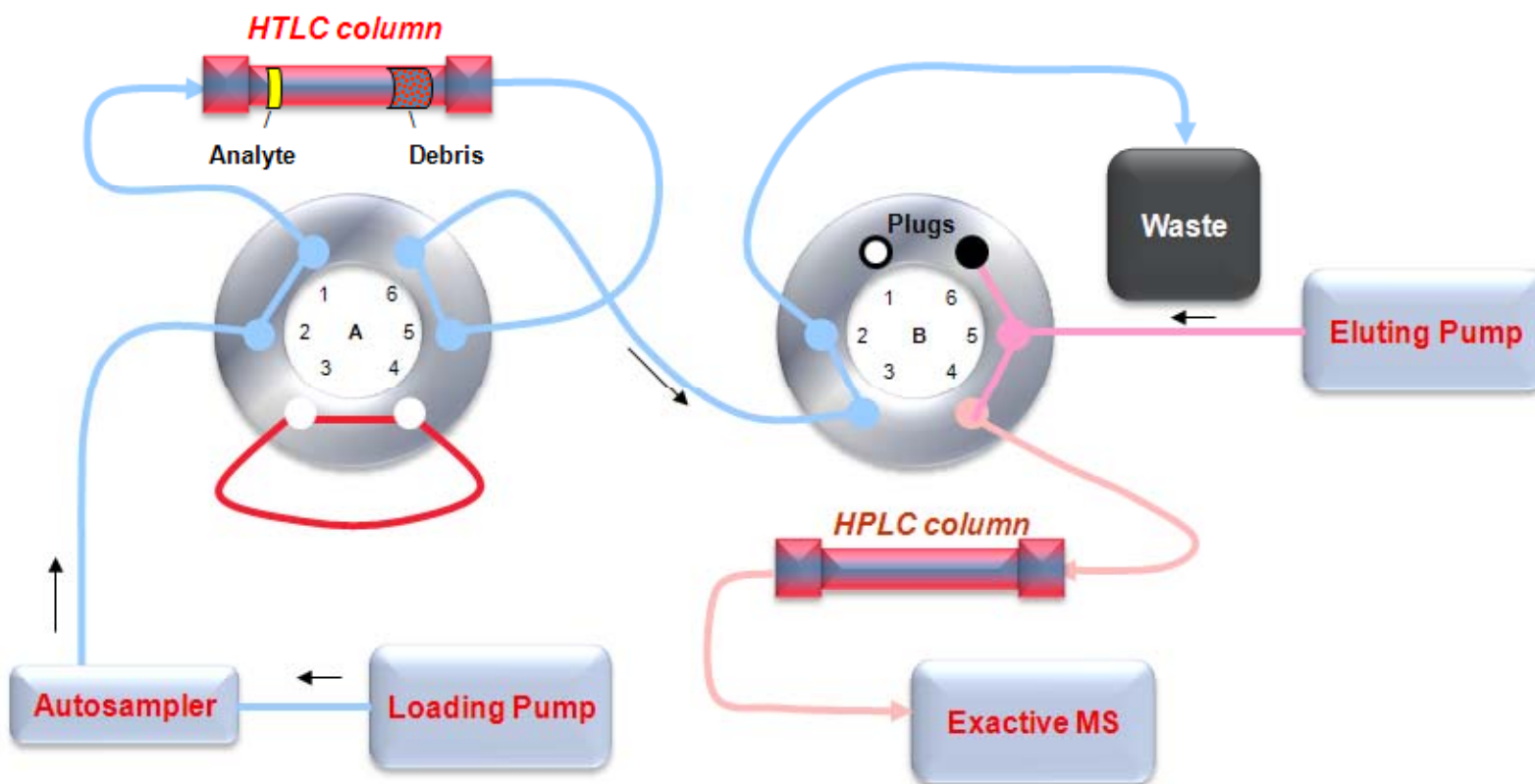
* *Molecularly imprinted polymer SPE column*

CVUAs analytical set up

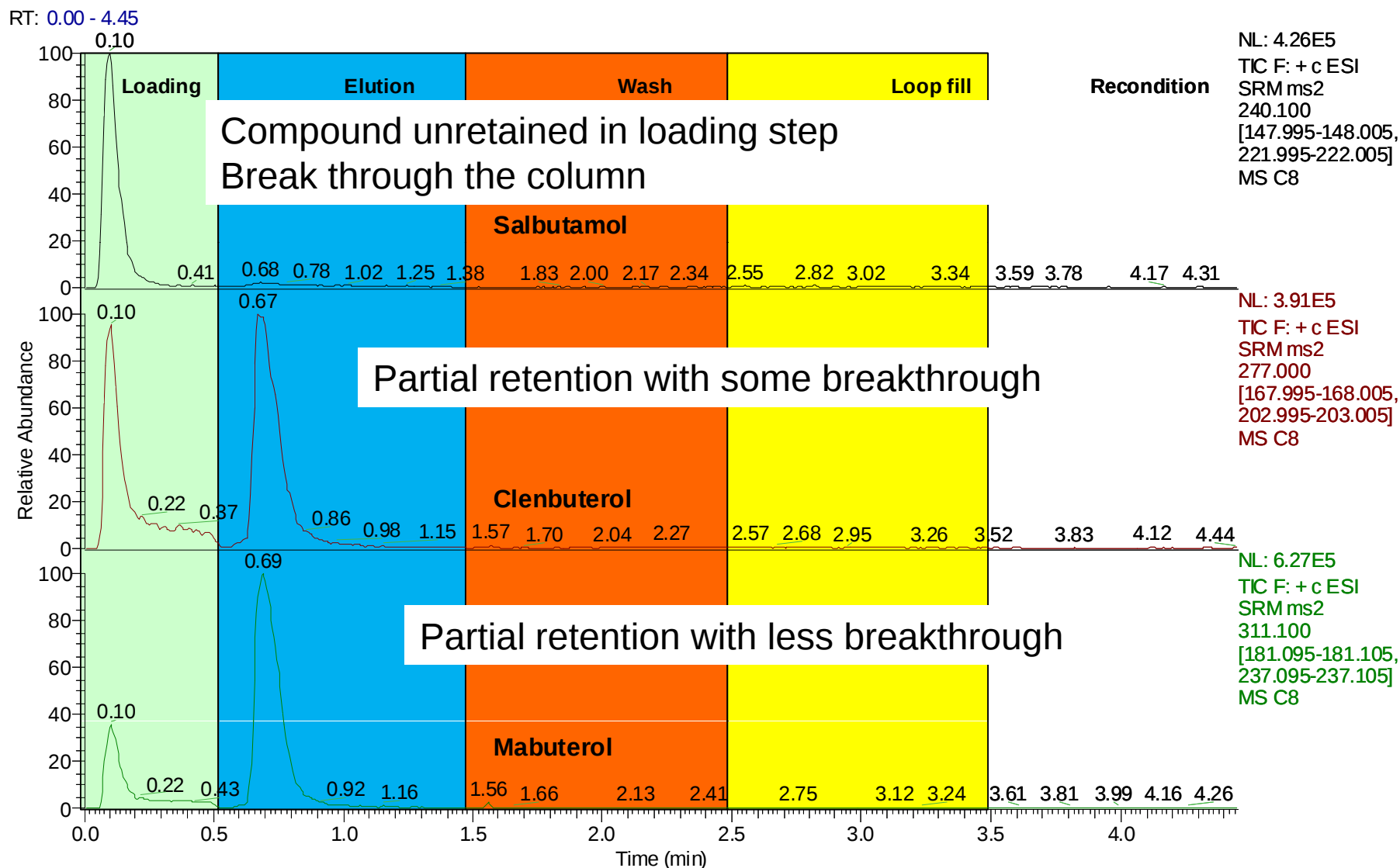
**Thermo TurboFlow® coupled with
Exactive™ High Resolution LC-MS/MS**



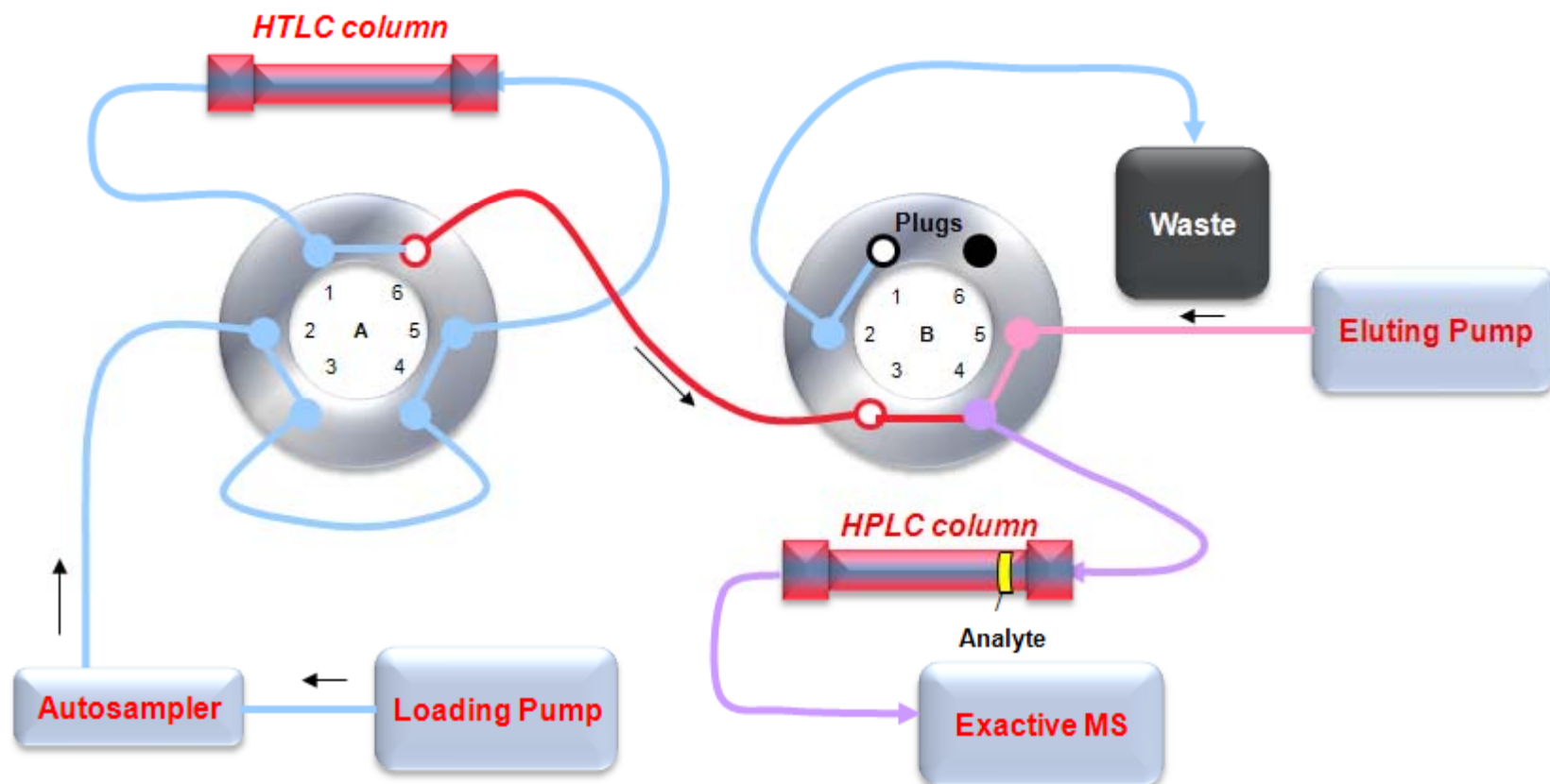
TurboFlow: Loading Step



Loading step: Example of the chromatogram



TurboFlow: Transfer/Focusing Step



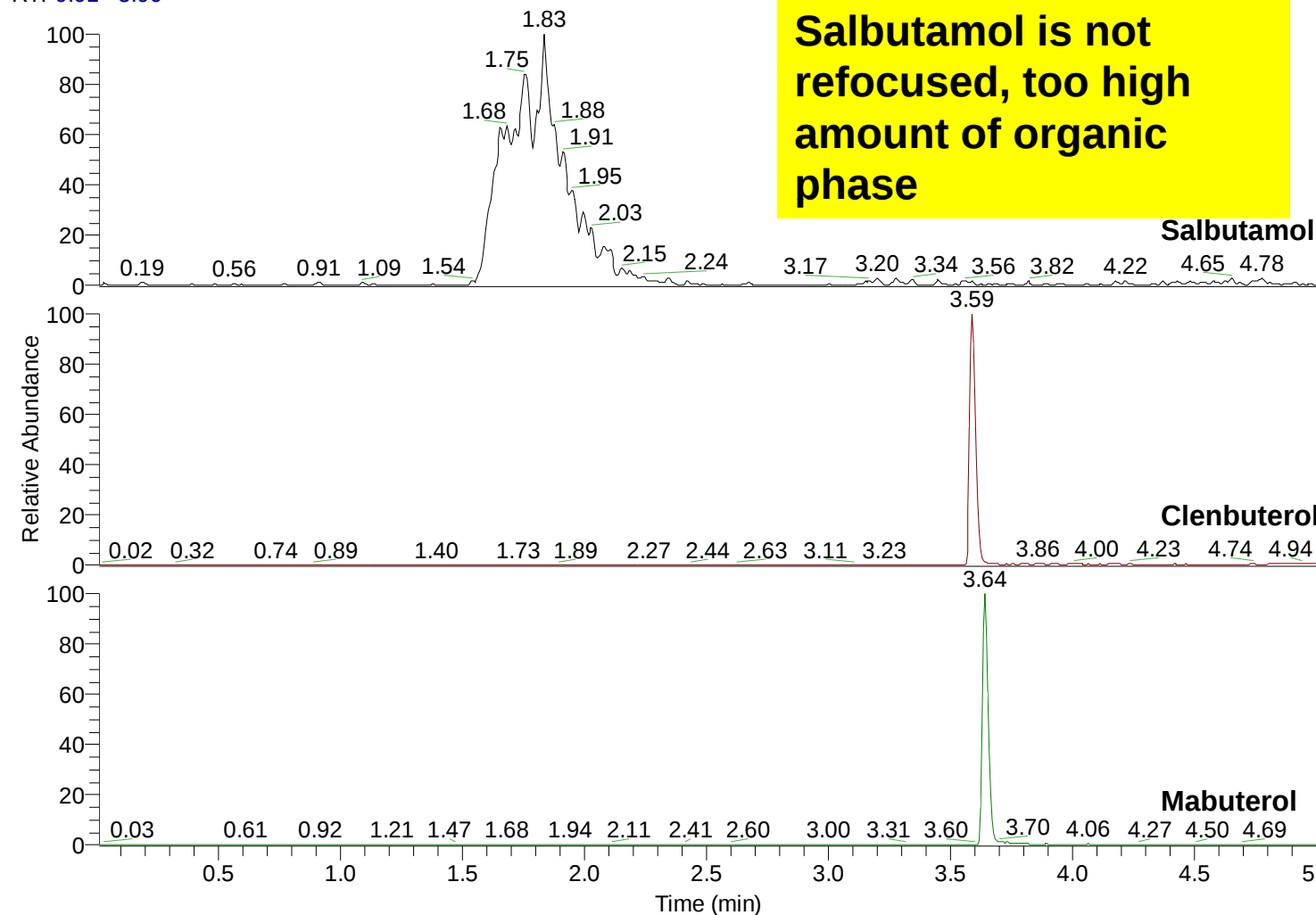
Transfer to analytical column – 250:550, 0% Methanol

C:\Xcalibur\...\CYCLONE-P-PH-0-550-250
None

18.3.2009 12:28:36

CYCLONE-P-PH-0-550-250

RT: 0.01 - 5.00



NL: 3.39E4
m/z= 147.50-148.50 F: + c
ESI SRM ms2 240.100
[147.995-148.005,
221.995-222.005] MS
CYCLONE-P-PH-0-550-250

NL: 6.90E5
TIC F: + c ESI SRM ms2
277.000 [167.995-168.005,
202.995-203.005] MS
CYCLONE-P-PH-0-550-250

NL: 1.05E6
TIC F: + c ESI SRM ms2
311.100 [181.095-181.105,
237.095-237.105] MS
CYCLONE-P-PH-0-550-250

Transfer to analytical column – 100:600, 0% Methanol

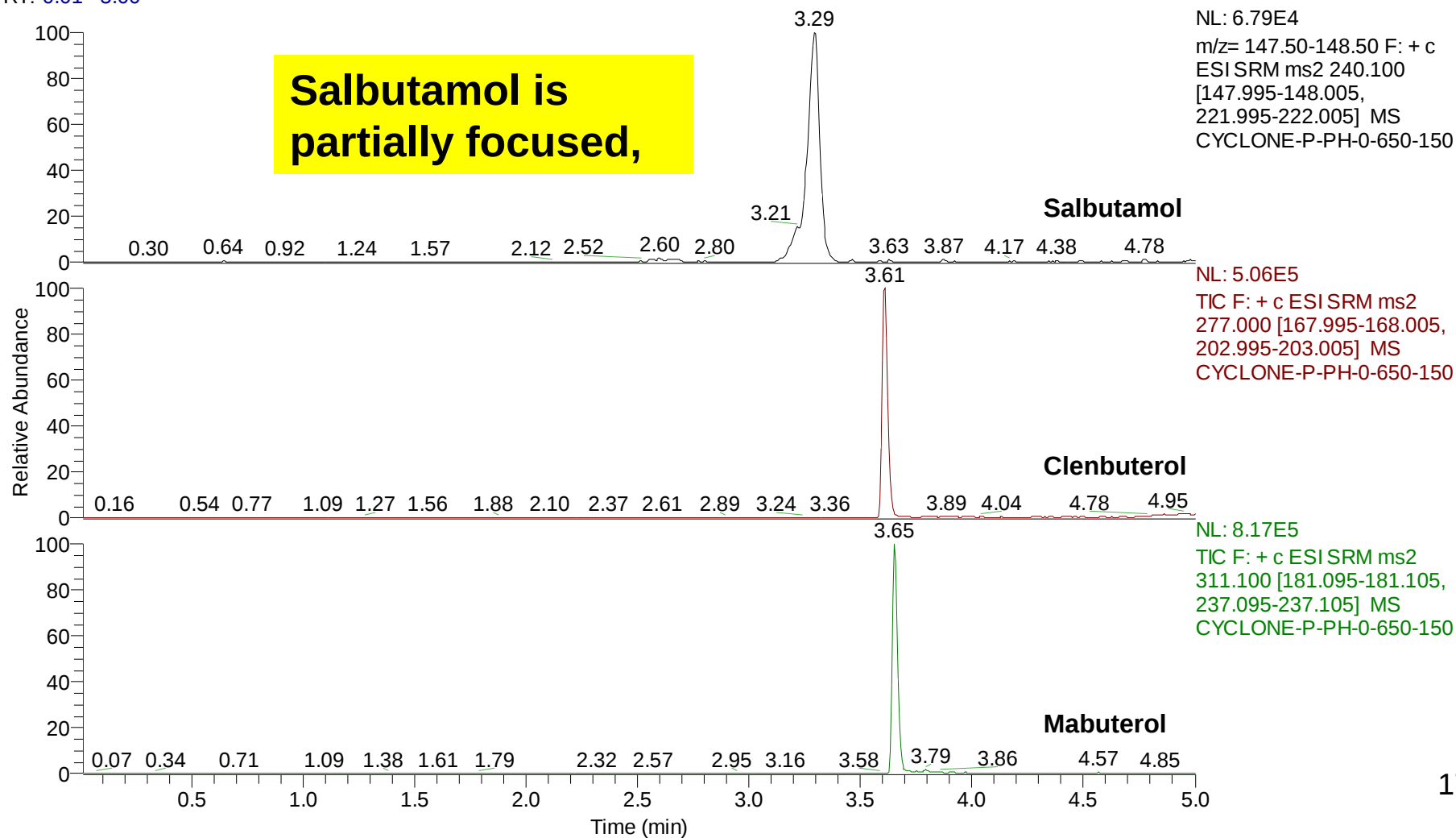
C:\Xcalibur\...\CYCLONE-P-PH-0-650-150

18.3.2009 12:36:00

CYCLONE-P-PH-0-650-150

None

RT: 0.01 - 5.00



Analytical method

TLX Column

Cyclone-P, 0.5 mm x 50 mm

Loading

10 mM NH₄Ac in Water, pH 8, 1.5 ml/min, 30 s

Elution

Methanol/0.1% acetic acid, 1/1, 0.1 ml/min, 120 s

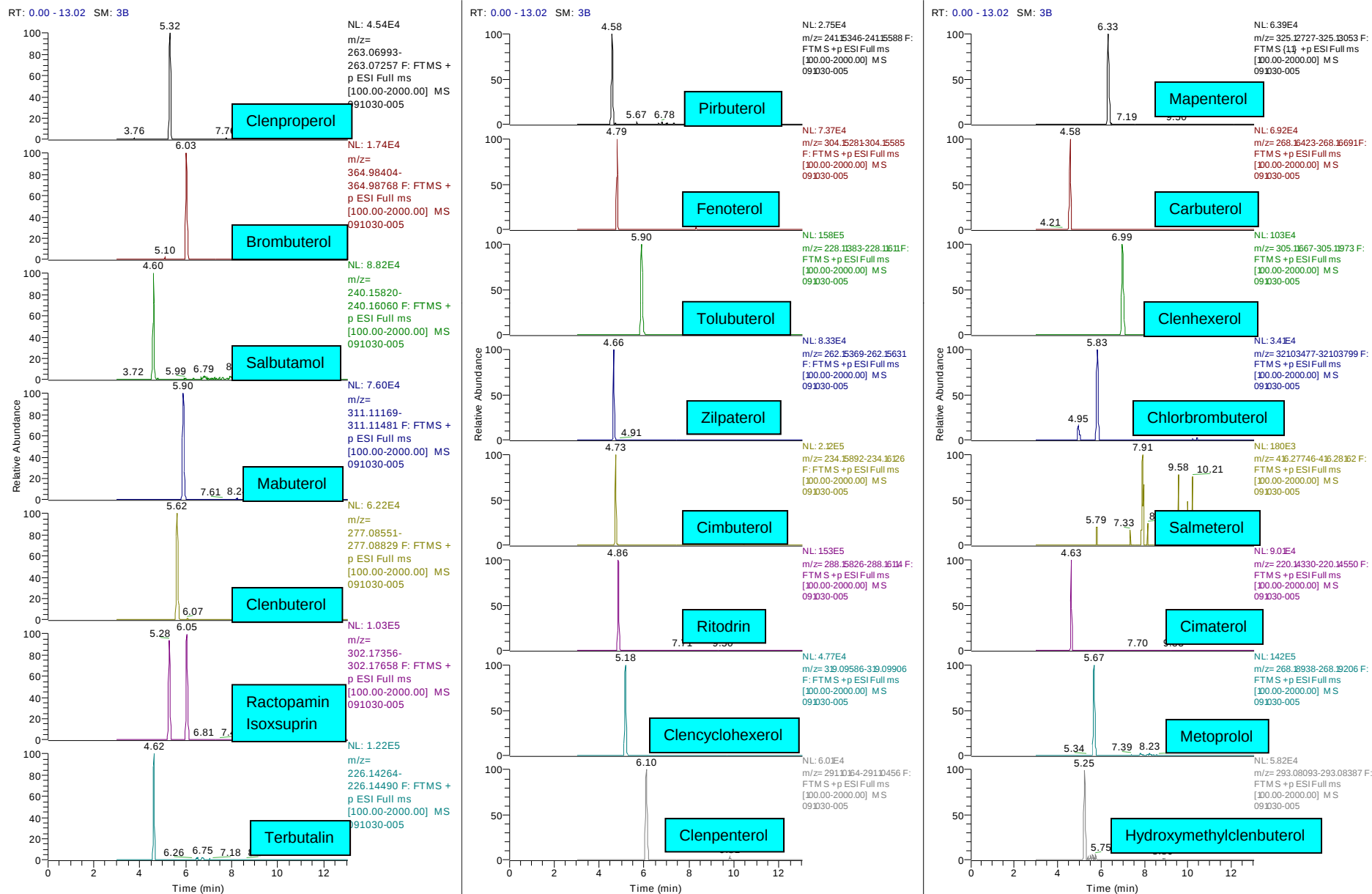
Wash

Methanol/0.1% acetic acid, 8/2, 1.5 ml/min, 60 s
Acetonitrile/acetone/isopropanol, 3/3/4

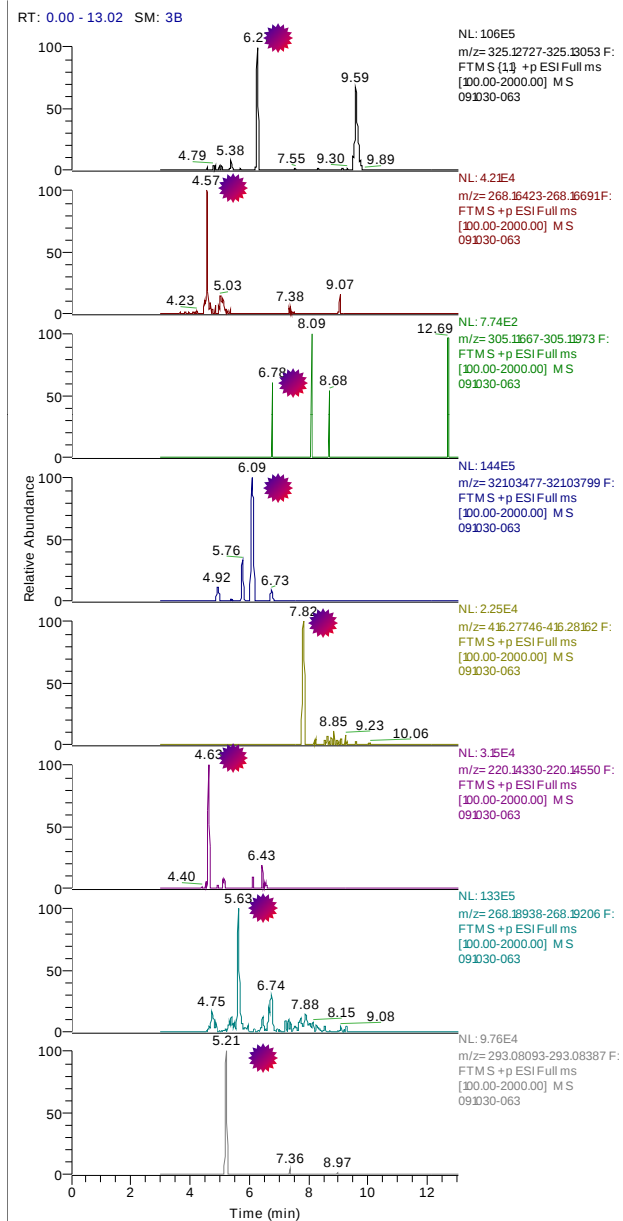
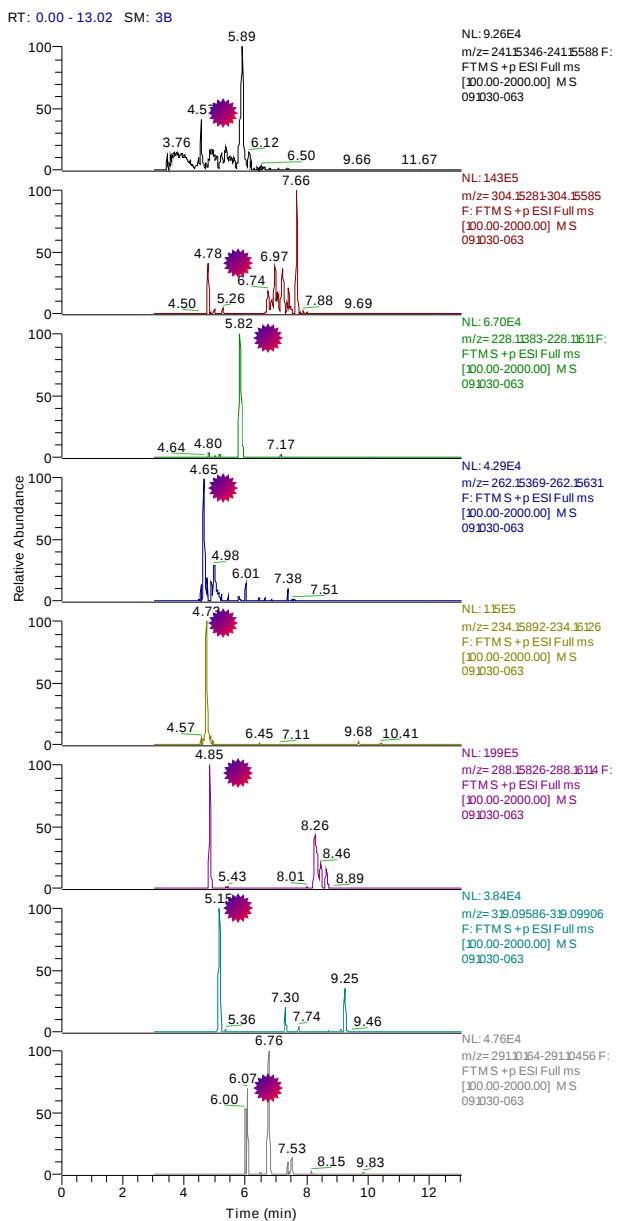
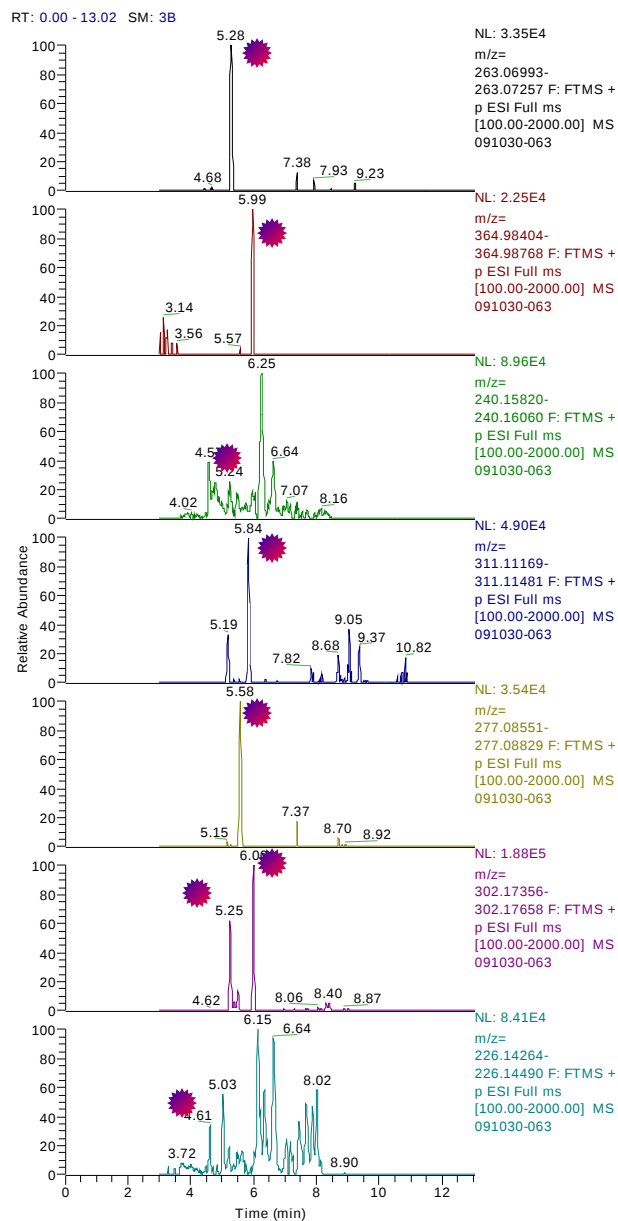
HPLC

Column: Phenyl Column 4.6 x 100 mm
Gradient: A: Methanol, B: 0.1% acetic acid
from 0% B to 95 % B in 300 s

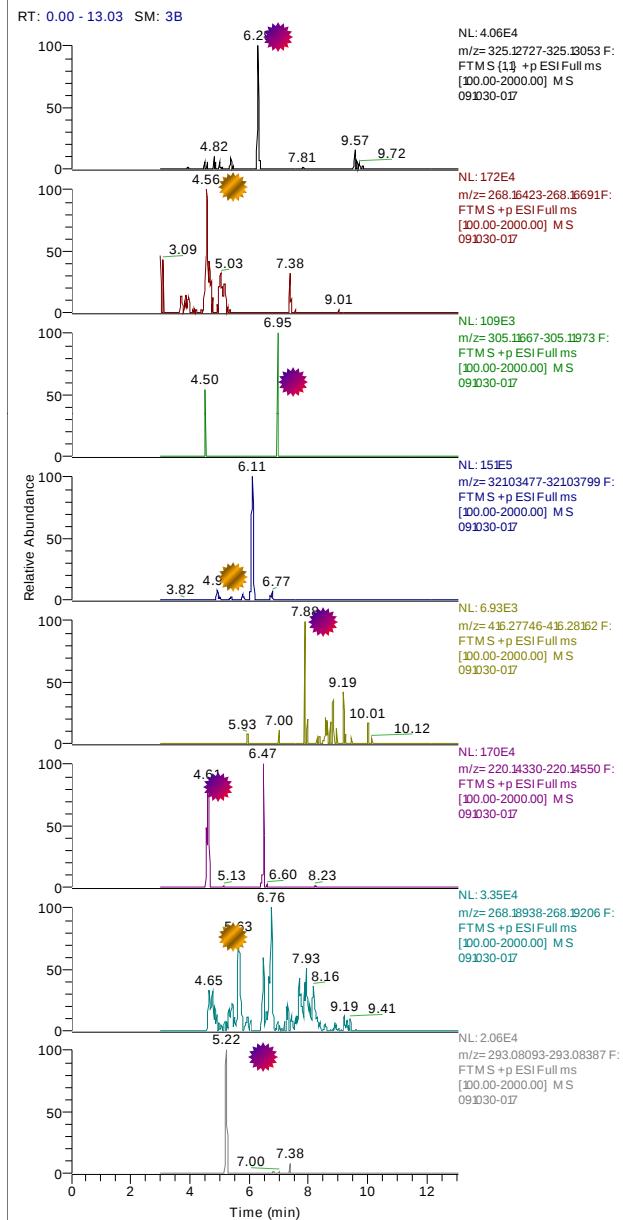
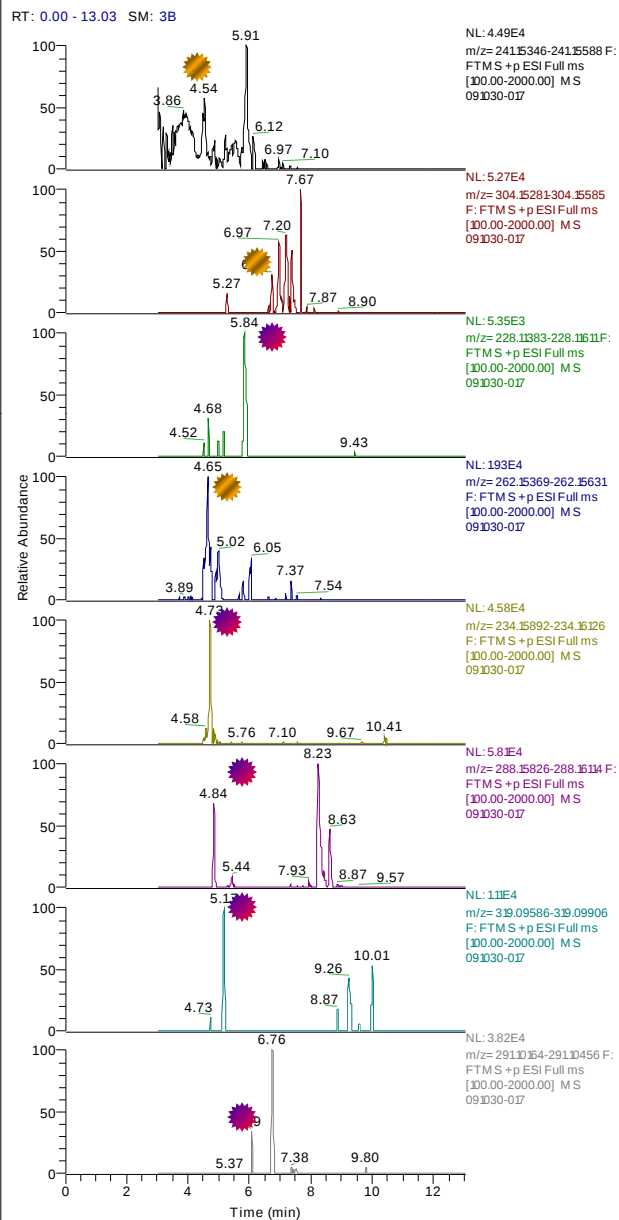
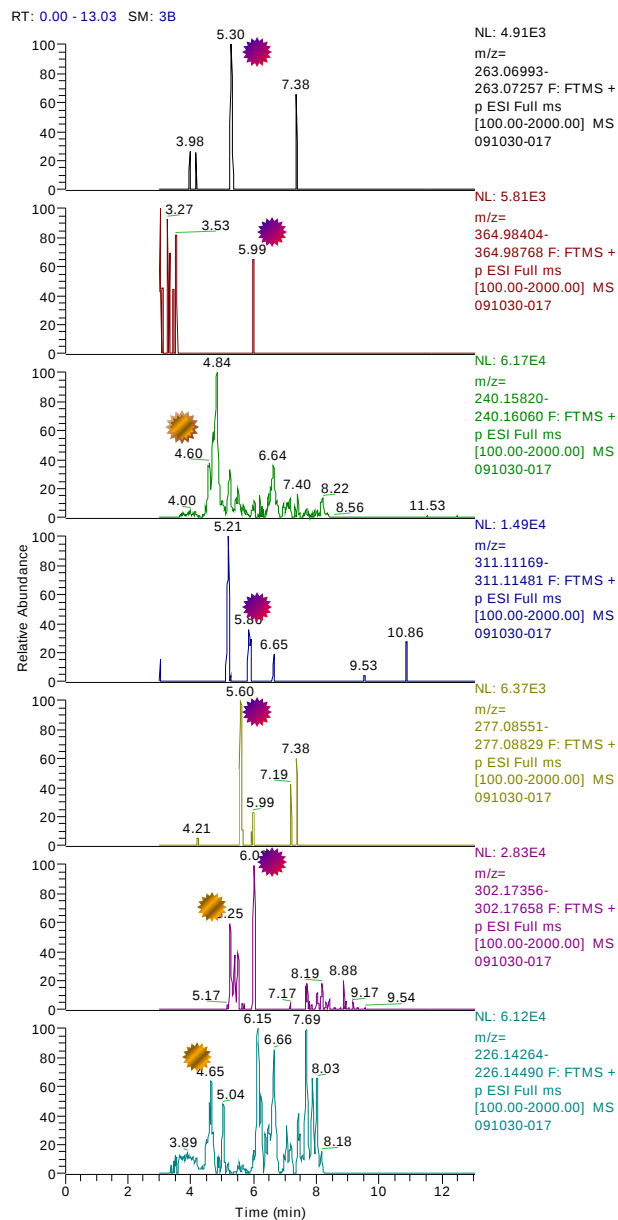
24 beta-agonists (standard; each 0.25 ng/ml)



beta-agonists in urine (spike: 1.0 ng/ml)



beta-agonists in urine (spike: 0.25 ng/ml)



Instruments

Acquisition

Explore

Quantitate



Method



Sequence



Survey



Review All



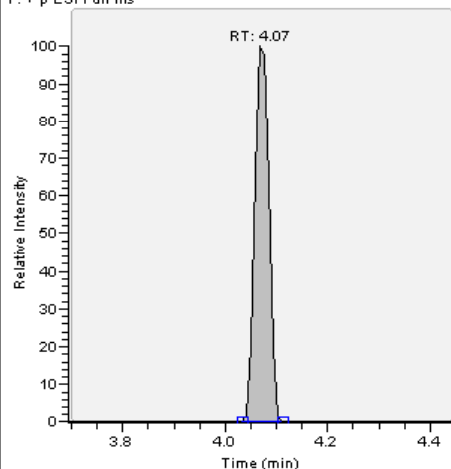
Reports

Review All Results - review standards, QCs, blanks and unknowns

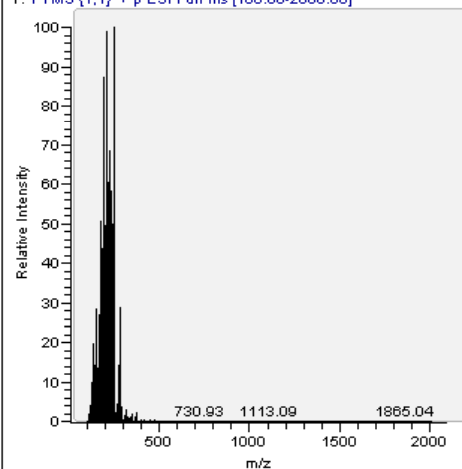
	Filename	Sample ID	Sample Type	Integration	Respo	ISTD Response	Response Ratio	Specified Conc	Calculated Conc	% Diff	% RSD
1	090202-0132	Urin 0.1-01	Standard	Method S	10966	NA	10966.217	0.100	0.085	-15.30	20.79
2	090202-0133	Urin 0.1-02	Standard	Method S	10771	NA	10771.123	0.100	0.083	-16.65	20.79
3	090202-0134	Urin 0.1-03	Standard	Method S	16427	NA	16427.022	0.100	0.123	22.53	20.79
4	090202-0135	Urin 0.1-04	Standard	Method S	13485	NA	13484.827	0.100	0.102	2.15	20.79
5	090202-0136	Urin 0.1-05	Standard	Method S	7714	NA	7713.508	0.100	0.062	-37.83	20.79
6	090202-0137	Urin 0.1-06	Standard	Method S	15336	NA	15335.605	0.100	0.115	14.97	20.79
7	090202-0138	Urin 0.1-07	Standard	Method S	16971	NA	16970.564	0.100	0.126	26.29	20.79
8	090202-0139	Urin 0.1-08	Standard	Method S	14080	NA	14079.536	0.100	0.106	6.27	20.79
9	090202-0140	Urin 0.1-09	Standard	Method S	18227	NA	18226.540	0.100	0.135	34.99	20.79
10	090202-0141	Urin 0.1-10	Standard	Method S	13845	NA	13844.663	0.100	0.105	4.64	20.79
11	090202-0142	Urin 0.1-11	Standard	Method S	14340	NA	14339.739	0.100	0.108	8.07	20.79
12	090202-0143	Urin 0.1-12	Standard	Method S	12592	NA	12591.644	0.100	0.096	-4.04	20.79
13	090202-0144	Urin 0.1-13	Standard	Method S	16801	NA	16800.781	0.100	0.125	25.12	20.79
14	090202-0145	Urin 0.1-14	Standard	Method S	16594	NA	16594.055	0.100	0.124	23.69	20.79
15	090202-0146	Urin 0.1-15	Standard	Method S	14749	NA	14749.050	0.100	0.111	10.91	20.79
16	090202-0147	Urin 0.1-16	Standard	Method S	10619	NA	10619.373	0.100	0.082	-17.70	20.79
17	090202-0148	Urin 0.1-17	Standard	Method S	10040	NA	10039.813	0.100	0.078	-21.71	20.79
18	090202-0149	Urin 0.1-18	Standard	Method S	8218	NA	8217.683	0.100	0.066	-34.34	20.79
19	090202-0150	Urin 0.1-19	Standard	Method S	14366	NA	14366.350	0.100	0.108	8.25	20.79
20	090202-0151	Urin 0.1-20	Standard	Method S	17730	NA	17729.568	0.100	0.132	31.55	20.79
21	090202-0153	Urin 0.25-01	Standard	Method S	36718	NA	36718.130	0.250	0.263	5.23	8.61
22	090202-0154	Urin 0.25-02	Standard	Method S	28963	NA	28963.414	0.250	0.209	-16.25	8.61
23	090202-0155	Urin 0.25-03	Standard	Method S	32929	NA	32929.079	0.250	0.237	-5.26	8.61
24	090202-0156	Urin 0.25-04	Standard	Method S	34352	NA	34352.264	0.250	0.247	-1.32	8.61
25	090202-0157	Urin 0.25-05	Standard	Method S	35579	NA	35579.479	0.250	0.255	2.08	8.61

Carbuterol
Pirbuterol
Salbutamol
Terbutalin
Cimaterol
Zilpaterol
Cimbuterol
Fenoterol
Ritodrin
Clencyclohexerol
Isosuprine
Hydroxymethylclenbuterol
Clenproperol
Clenbuterol
Netoprolol
Chlorbrombuterol
Mabuterol
Tulobuterol
Ractopamin
Brombuterol
Clenpenterol
Mapenterol
Clenhexerol
Salmeterol

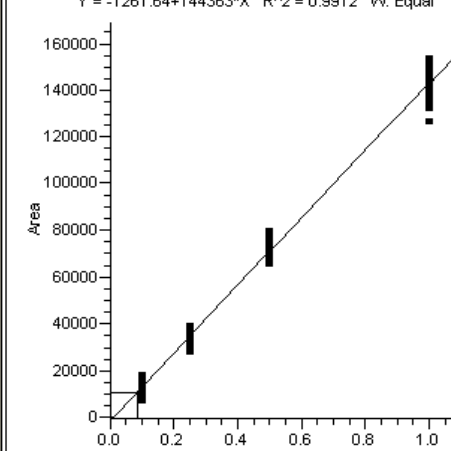
Selected Chromatogram

090202-0132 - m/z= 268.17-268.17 SM: 5 RT: 3.69 - 4.44 NL:
F: + p ESI Full ms

Component Spectrum

090202-0132 #151 RT: 1.22 NL: 2.31E7
T: FTMS {1,1} + p ESI Full ms [100.00-2000.00]

Calibration Curve

Carbuterol
Y = -1261.64+144363*X R^2 = 0.9912 W: Equal

Validation parameters



Compound name	CC α ($\mu\text{g/l}$)	CC β ($\mu\text{g/l}$)	Repeatability RSD (%) (0.5ppb)	In-house repro- ducibility RSD (%) (0.5ppb)	Recovery (%)
Brombuterol	0.19	0.46	20	24	97
Carbuterol	0.22	0.58	25	27	100
Chlorbrombuterol	0.14	0.38	14	26	99
Cimaterol	0.13	0.35	15	19	100,0
Cimbuterol	0.08	0.20	9	13	100
Clenbuterol	0.14	0.33	13	25	98
Clencyclohexerol	0.08	0.21	9	12	100
Clenhexerol	0.22	0.49	22	23	98
Clenpenterol	0.20	0.49	22	25	99
Clenproperol	0.10	0.24	11	15	100
Fenoterol	0.11	0.24	11	14	99
Hydroxymethylclenbuterol	0.09	0.21	9	14	100
Isoxsuprine	0.10	0.25	11	18	100
Mabuterol	0.09	0.23	9	17	100

Many thanks to



Michal Godula

Thermo Fisher Scientific

and our team

**Thank you for
your attention**