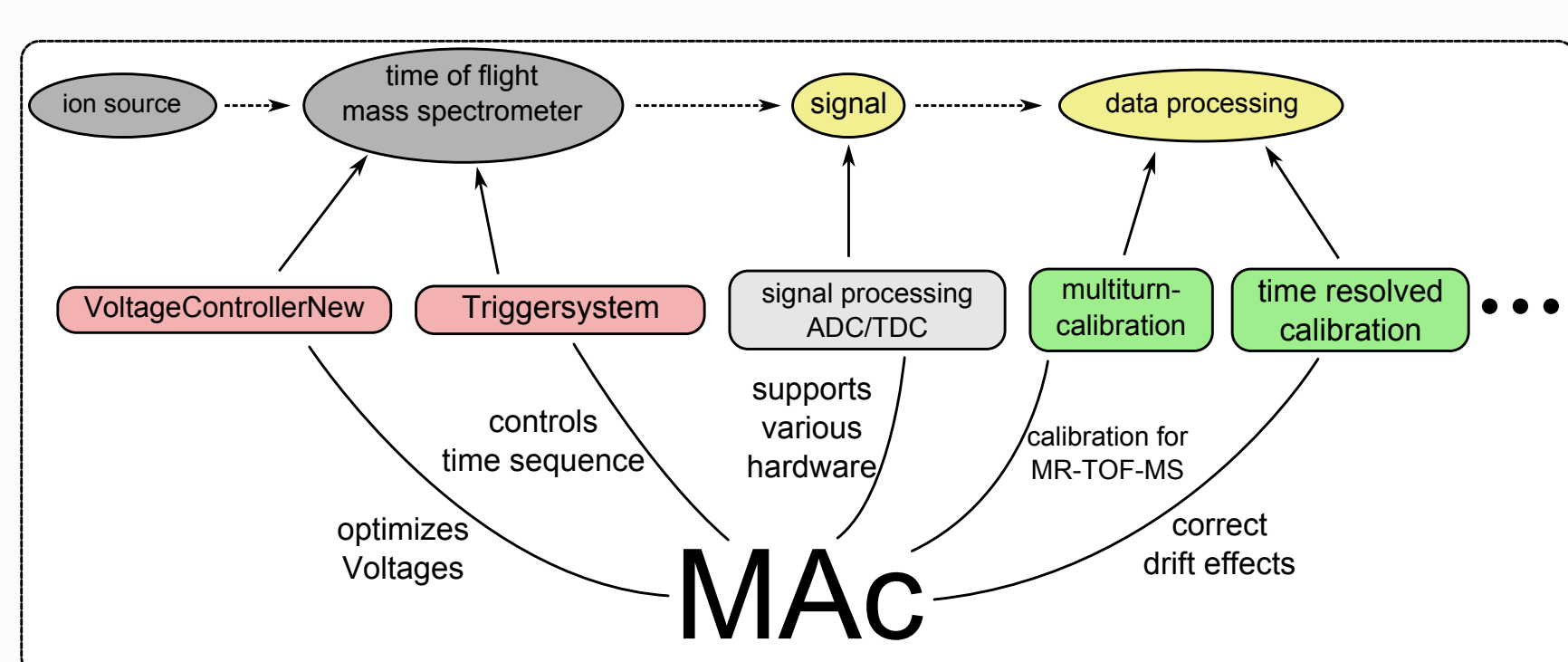


Julian Bergmann<sup>[1]</sup>, Alexander Pikhtelev<sup>[2]</sup>, Samuel Ayet San Andres<sup>[3]</sup>, Timo Dickel<sup>[1,3]</sup>, Jens Ebert<sup>[1,4]</sup>, Hans Geissel<sup>[1,3]</sup>, Christine Hornung<sup>[1]</sup>, Johannes Lang<sup>[1]</sup>, Wayne Lippert<sup>[1]</sup>, Christian Lotze<sup>[1]</sup>, Wolfgang Plaß<sup>[1,3]</sup>, Ann-Kathrin Rink<sup>[1]</sup> and Christoph Scheidenberger<sup>[1,3]</sup>

<sup>[1]</sup>JLU Gießen, Deutschland — <sup>[2]</sup>Institute of Energy Problems of Chemical Physics, Russian Academy of Sciences, Moscow, Russia — <sup>[3]</sup>GSI Darmstadt, Deutschland — <sup>[4]</sup>TU Darmstadt, Deutschland



## 1. MAC Software

- data acquisition
- data evaluation
- hardware control

### Features:

- + ADC/TDC for large dynamic range
- + time sequence controller
- + multi-reflection calibration
- + voltage optimizer
- + time resolved calibration
- + spectra accumulation
- + automatic peak detection
- + import/export various file formats
- + data operations (e.g. smoothing)

## 2. Enlarging dynamic range using TDC and ADC simultaneously.

### TDC:

- + detection of single Ions
- dead time effect

### ADC:

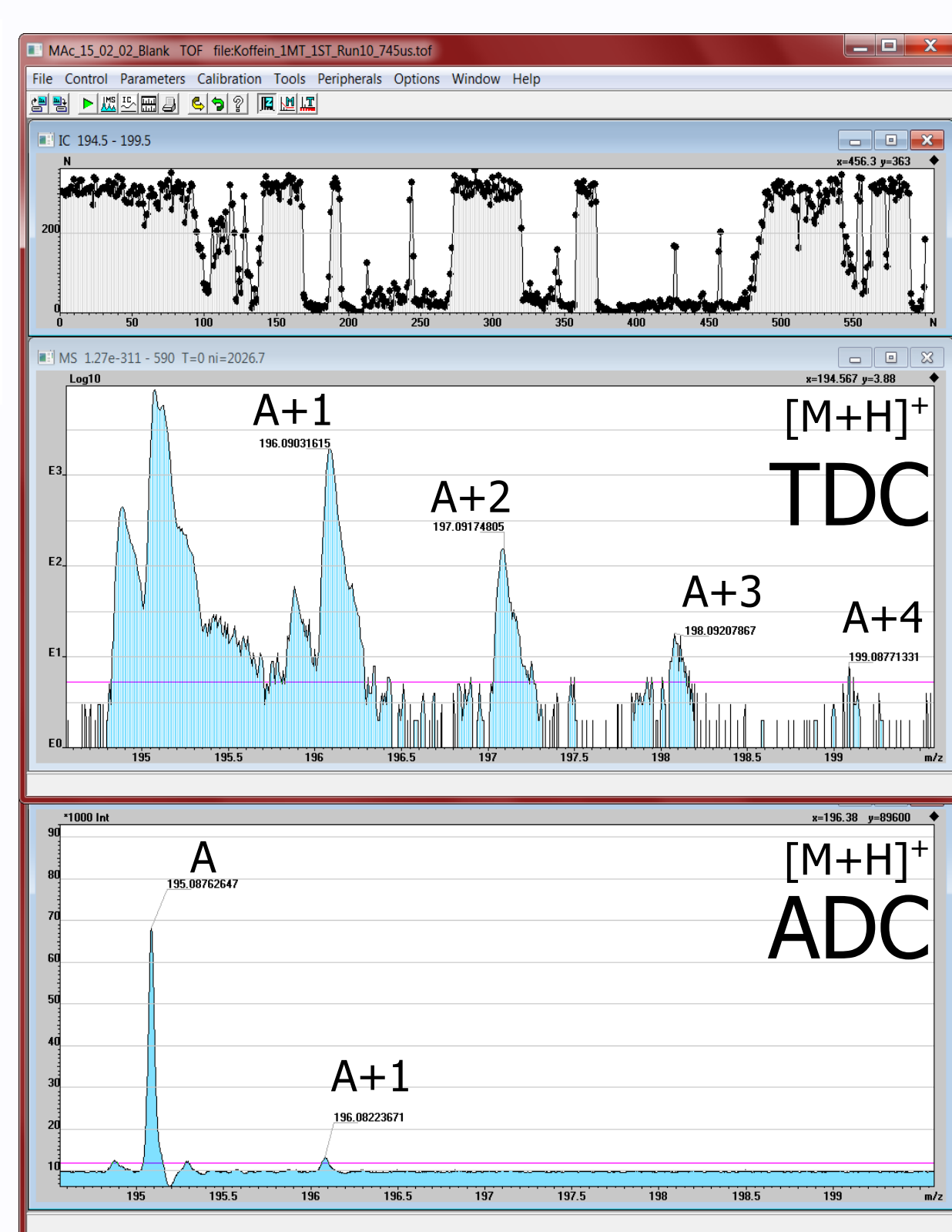
- + detection of high signal rates
- low intensity signals lost

### Combined:

- + dynamic range > 4 Orders of magnitude
- + detection of high and low signal rates

### MAC:

- common user Interface
- common calibration
- common file format and analysis tools



Example: Coffein ( $10^{-4}$  mol)

### TDC (top):

- + A+1 to A+4

### ADC (bottom):

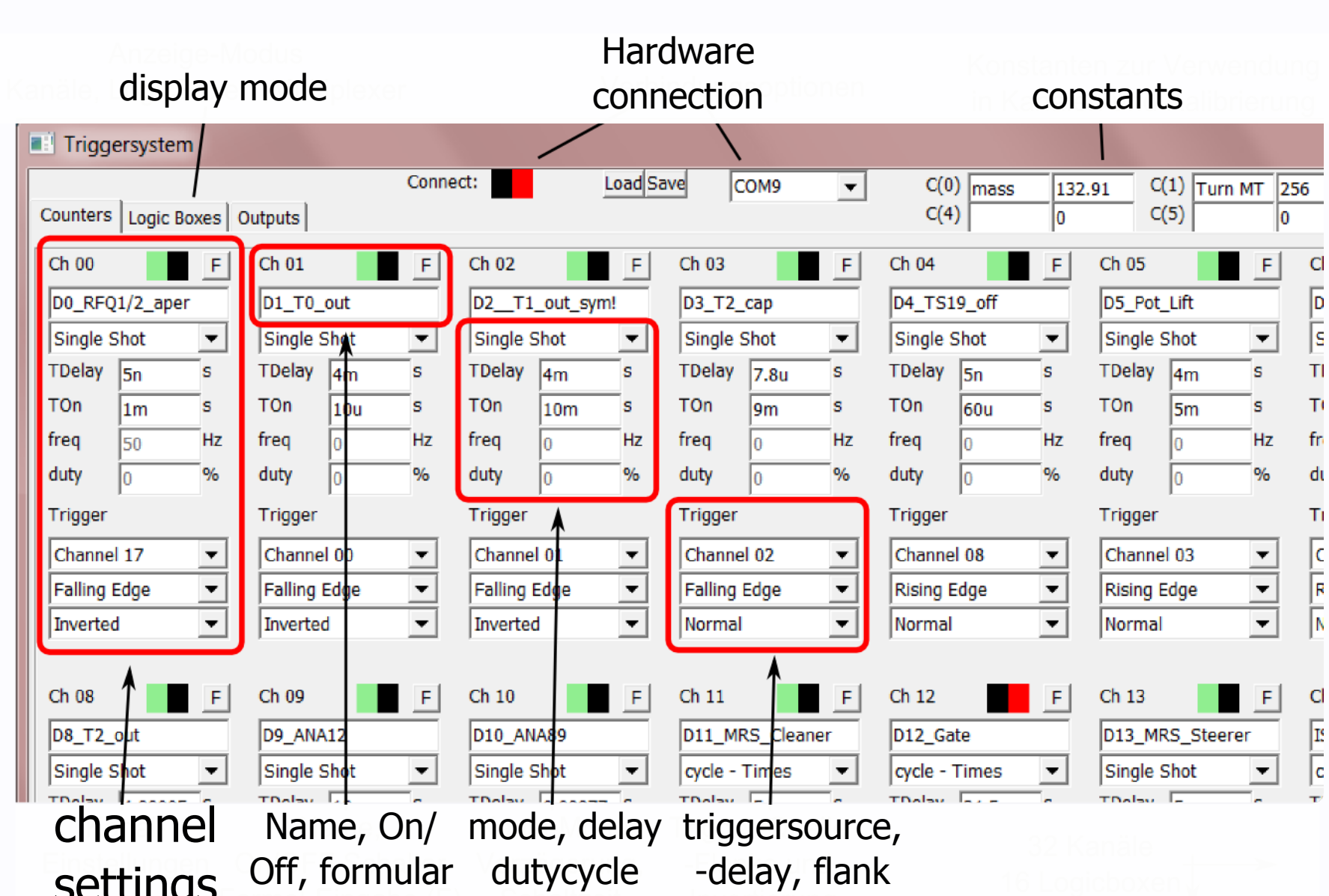
- + A and A+1

| Device | Isotope     | MAC      | Literature | rel. Deviation         |
|--------|-------------|----------|------------|------------------------|
| ADC    | m195 / m196 | 9,374 79 | 9,457 79   | 0,008 78               |
|        | m196 / m197 | 11,7599  | 11,7515    | $7,1480 \cdot 10^{-4}$ |
| TDC    | m197 / m198 | 5,629 95 | 19,1104    | 0,705 40               |
|        | m198 / m199 | 3,404 73 | 33,768     | 0,899 17               |

Excelent Isotop distribution  
unresolve Isobars

## 3. Time Sequence controller

- controls time sequences for traps, electrodes etc.
- multiple modes to set delays and duty cycles



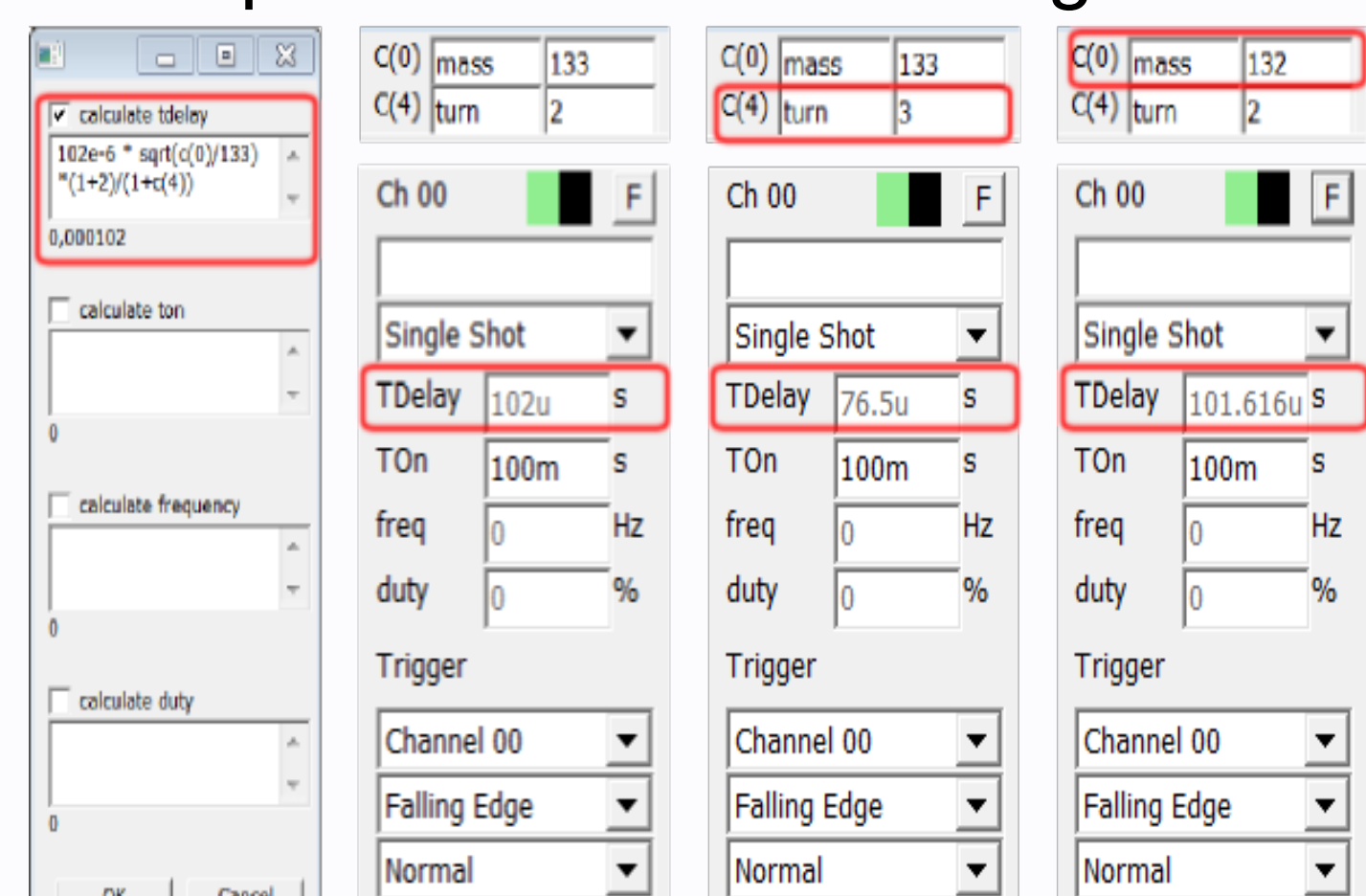
- + link triggers for 32 channels to each other or up to 16 logic channel combinations

- + use formulars to calculate delay and duty cycles

- + link to other channel's times

- + use up to 8 constants

### Example of automatic scaling



- + set times can be used in other MAC parts
- calibration

## 4. Multiturn Calibration

### Calibration Formular

$$m = a \frac{(t - t_0)^2}{(1 + b \cdot N)^2}$$

$$a = \frac{2q \cdot U}{I_{detec}^2} \quad b = \frac{I_{turn}}{I_{detec}}$$

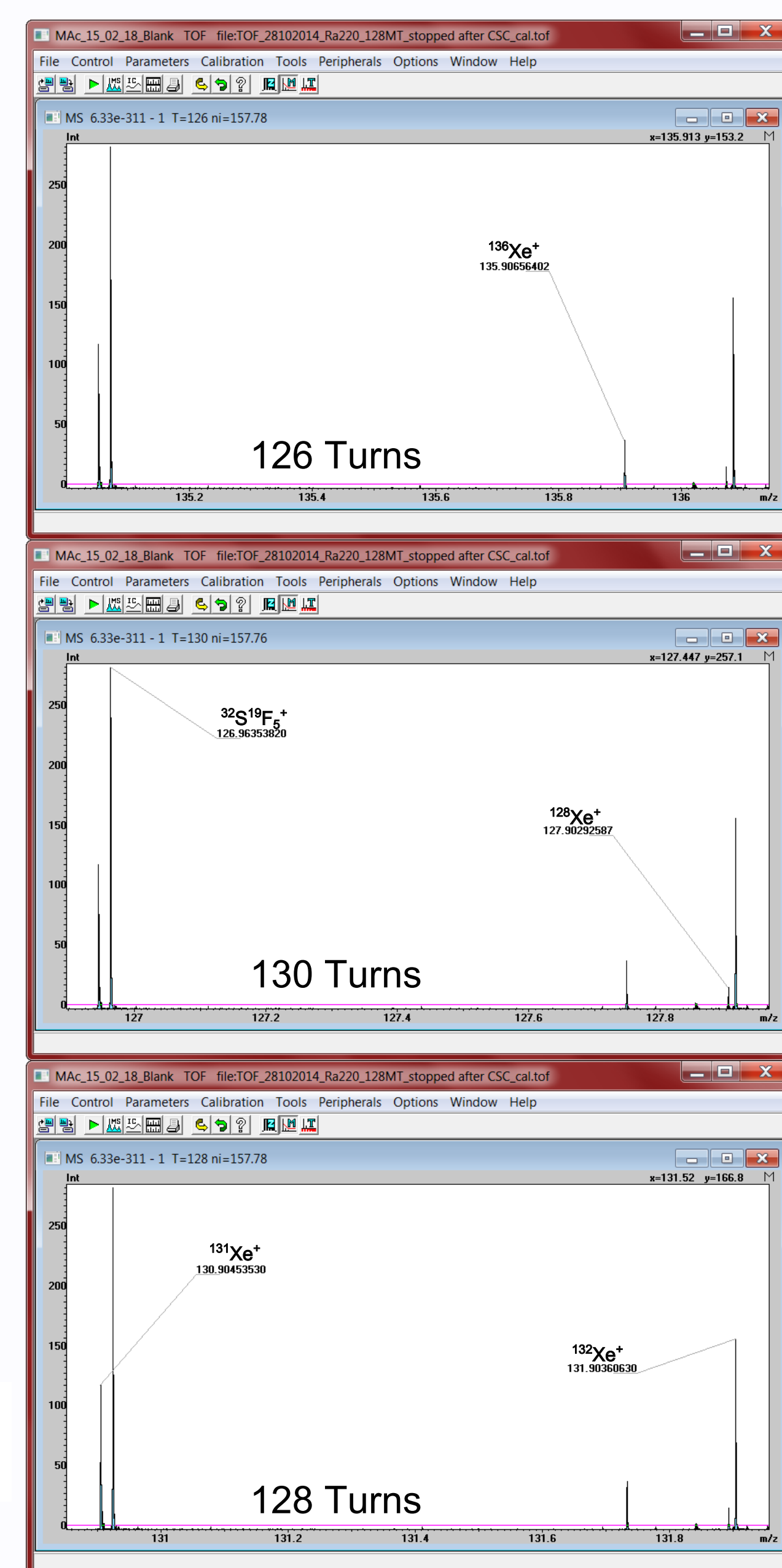
$t_0$ : electronic delay time

- + ≤3 calibrants → analytical
- + >3 calibrants → least square deviation minimizer
- + uses time sequence controller for trap extraction time
- + use and display for different turns in multiturn calibration
- + mass range greatly increased

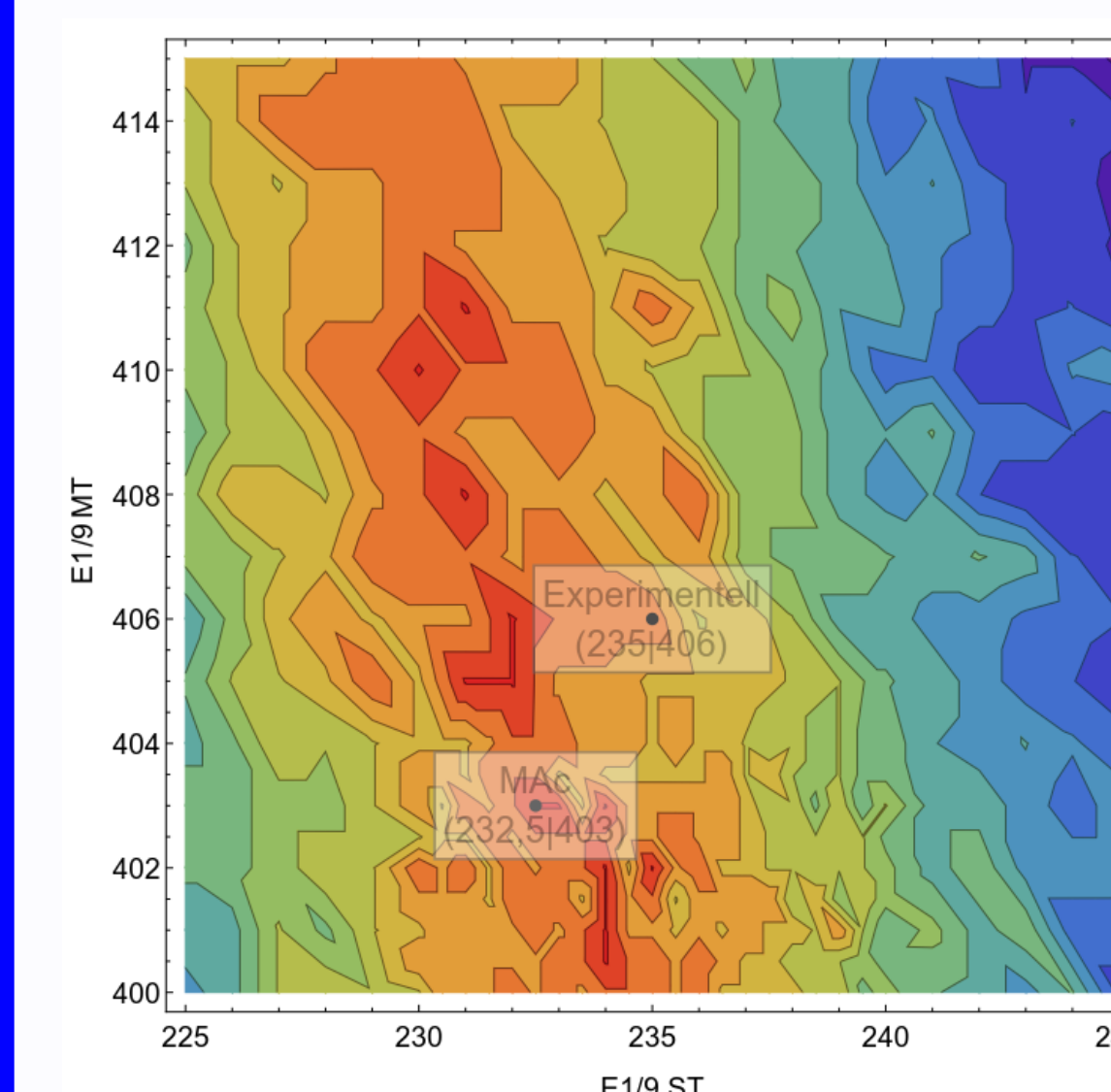
### Example: Xe & SF6

- + between 126 and 130 Turns
- + precision of online identified masses of  $4 \cdot 10^{-7}$

| Time (µs)  | Turns | Mass MAC (u) | Mass Lit (u) | Lit-MAC Lit          | Nuklid              |
|------------|-------|--------------|--------------|----------------------|---------------------|
| 49,3071469 | 130   | 127,9029259  | 127,9029822  | $4,40 \cdot 10^{-7}$ | $^{128}\text{Xe}^+$ |
| 46,5327600 | 126   | 135,906564   | 135,9066657  | $7,48 \cdot 10^{-7}$ | $^{136}\text{Xe}^+$ |



## 5. Voltage Optimizer



- connects to all available power supplies
- free choice for amount of channels
- from adjustable MIN to MAX all voltage combinations are tested
- automatic measurement taken for each combination
- + 20 measurements per minute
- $R^2 + a^2$  automatically calculated
- + factor a to weight Intensity

### Conclusion:

Indispensable for preparing in-situ measurements e.g. for Ambiprobe MR-TOF-MS

## 6. Time resolved calibration

### Problem:

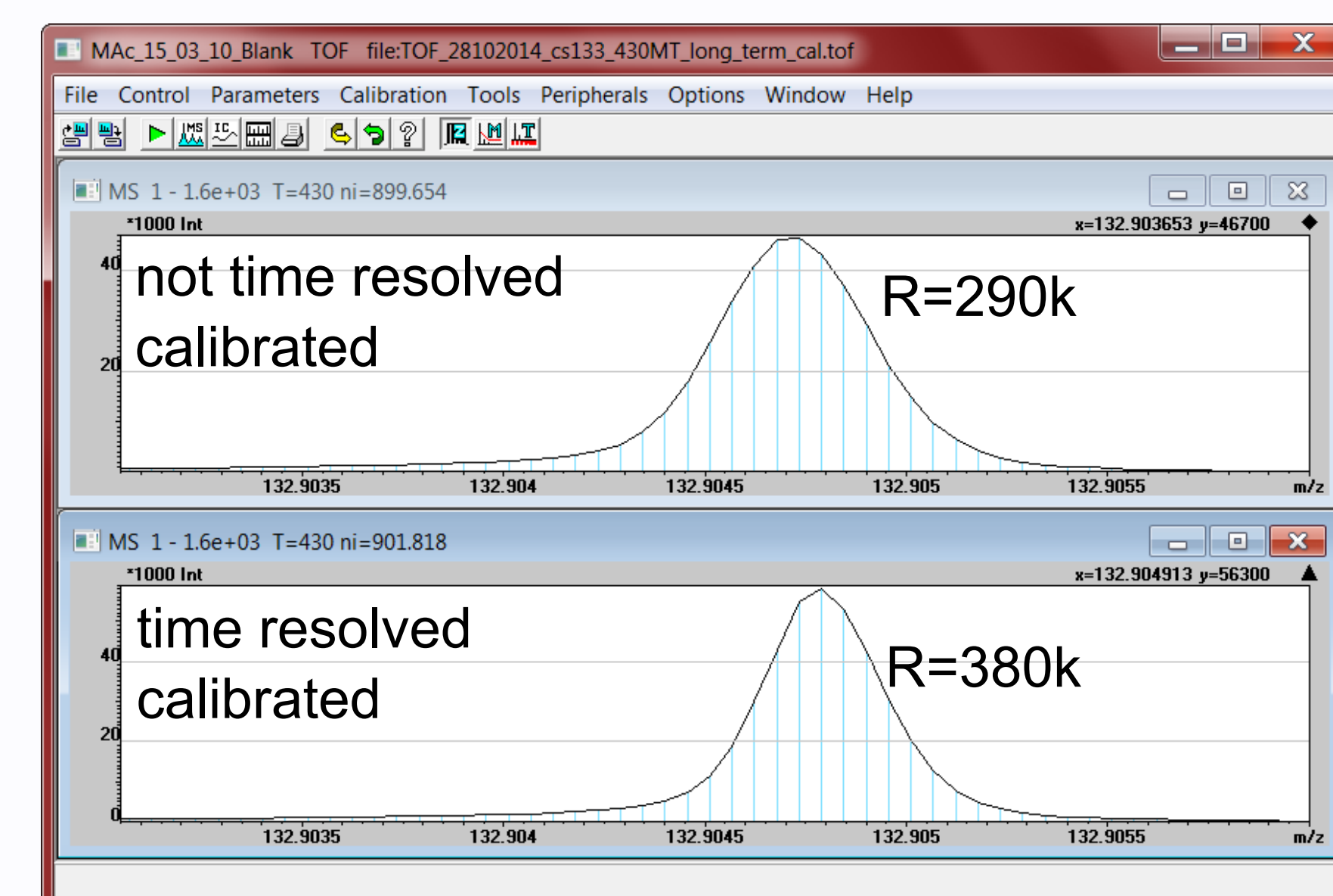
- voltages can vary during measurement
- thus flight length varies during turns
- mass lines shift in time

### Solution:

- + each spectra gets calibrated → calibrant stays fixed in mass

### MAC:

- + automatic detection of calibrant
- + correction of 2000 spectra in under 1 second
- + recalibrated data can be displayed and used inside MAC
- + ideally suited for long term or high resolution measurements



## Poster Information

Julian Bergmann  
julian.bergmann@physik.uni-giessen.de  
Justus Liebig Universität Gießen