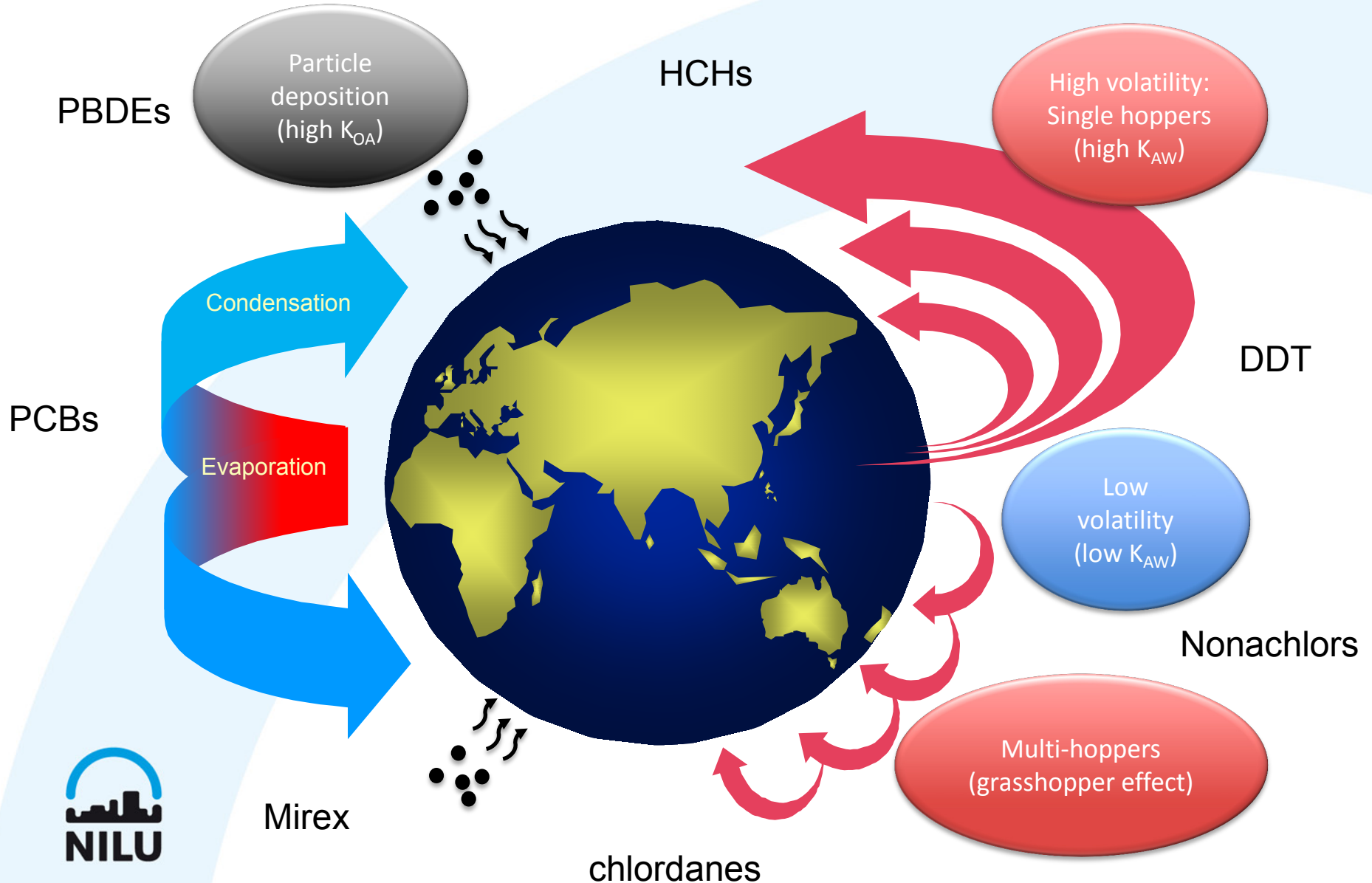


# Versatility of GC-Orbitrap mass spectrometry for the ultra-trace detection of persistent organic pollutants in penguin blood from Antarctic



**Dr. Nicholas A. Warner**  
*NILU-Norwegian Institute for Air Research*  
*Fram Centre, Tromsø, Norway*

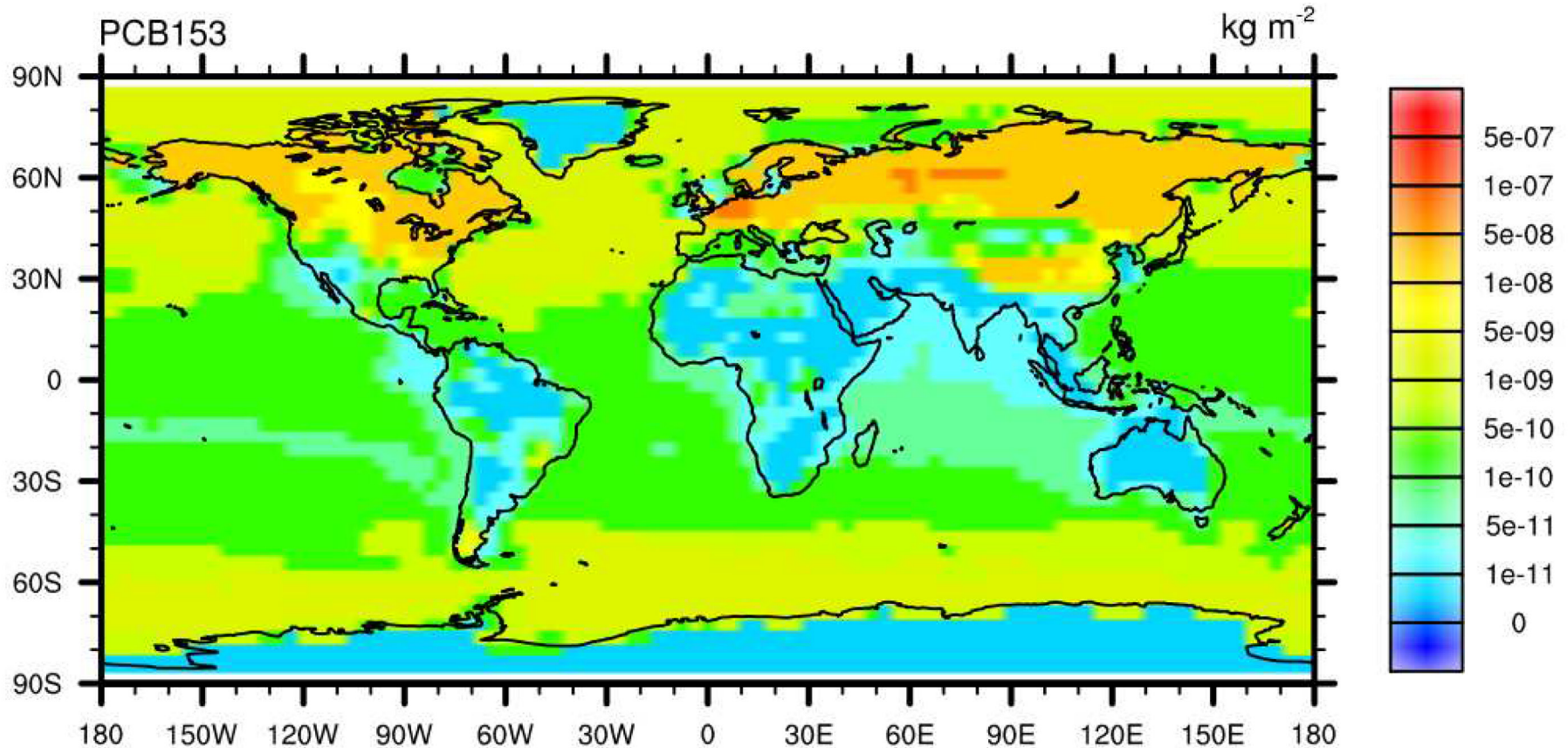
# Contaminants in Polar Regions



# Arctic vs. Antarctic

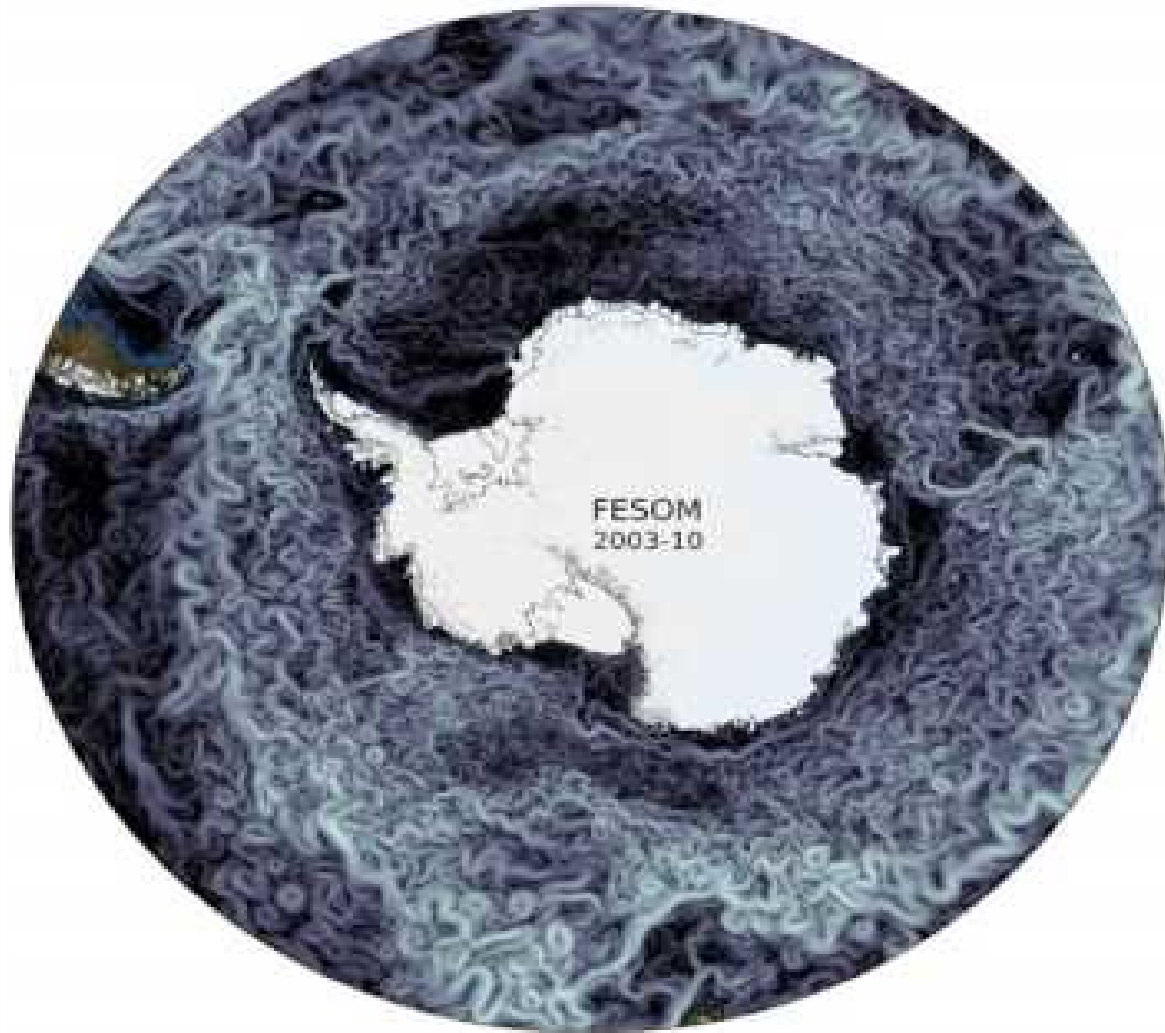
## Predicted global distribution of total PCB 153 environmental burden (2010)

(Lammel and Stemmler. Atmos. Chem. Phys. 2012)



# Arctic vs. Antarctic

## Antarctic Circumpolar Current

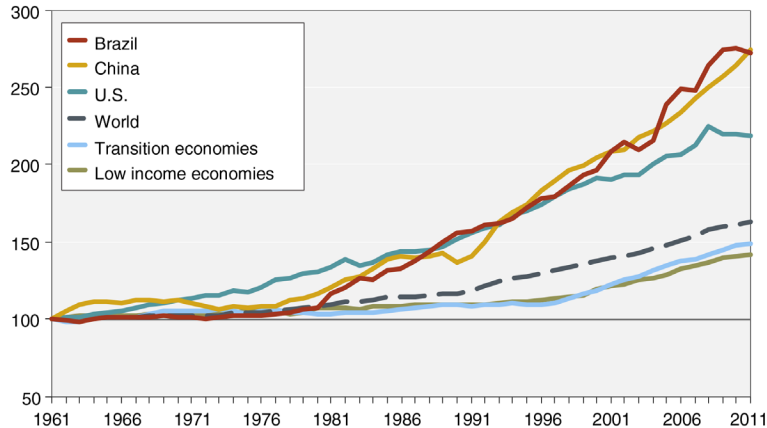




# Exposure risks to Antarctica?

Agricultural productivity (measured as total factor productivity) growth for selected countries/country groups, 1961-2011

Index (1961 = 100)



Note: Total factory productivity (TFP) takes into account all of the land, labor, capital, and material resources employed in farm production and compares them with the total amount of crop/livestock output. If total output is growing faster than total inputs, we call this an improvement in TFP.

Source: USDA, Economic Research Service, International Agricultural Productivity data product.

## Export of e-waste



Source: Greenpeace, Basel Action Network

activities

OC compounds  
detected in:  
*air, soil, biota*

**Contaminant monitoring needed to assess impacts of increased emission sources in Southern Hemisphere on Antarctic environment**



# POLAR-ECOTOX project

***«To quantify temporal changes in contaminants to identify climatic, ecological, and anthropogenic drivers affecting exposure in Antarctic wildlife»***

- King penguin blood (2000-2017)

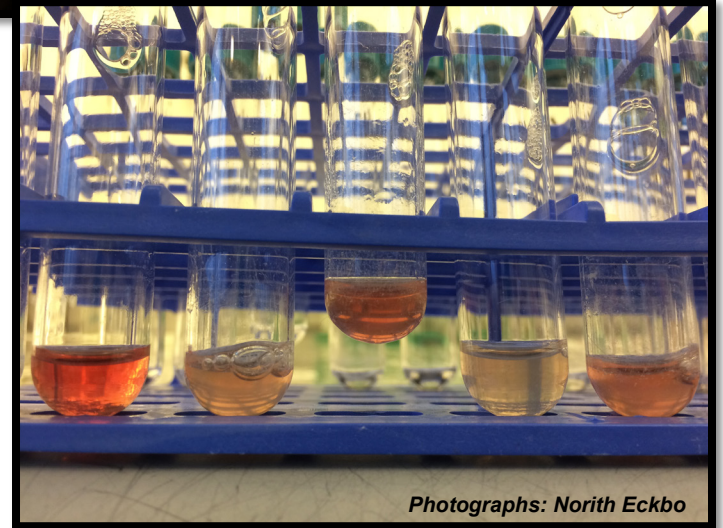


- Reside within and rely on the circumpolar current
- Provide a baseline to assess temporal changes in OC contaminant exposure



# Challenges

- Sample material
  - non-destructive sampling
  - Limited material
- Low concentration levels
- Sample processing minimized to avoid analyte losses



*Photographs: Norith Eckbo*

# Sample preparation and analysis

Liquid-liquid extraction  
50% EtOH( $\text{NH}_4$ ) $_2$ SO $_4$ :Hexane

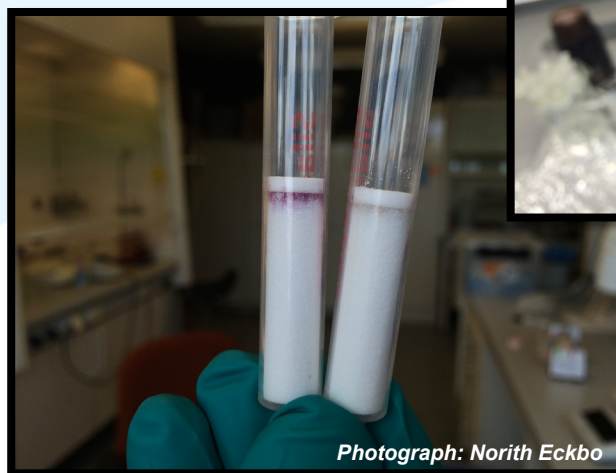
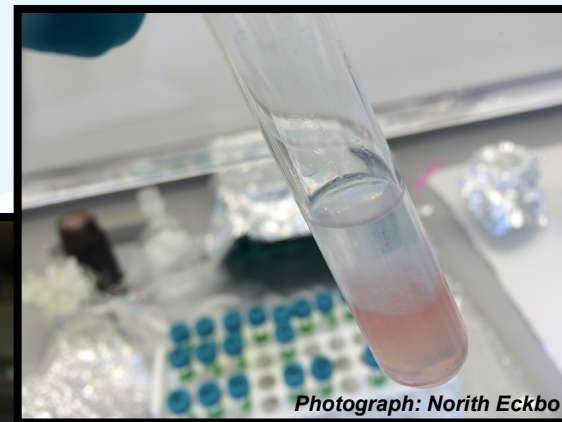


Automated SPE clean-up  
using activated Florisil



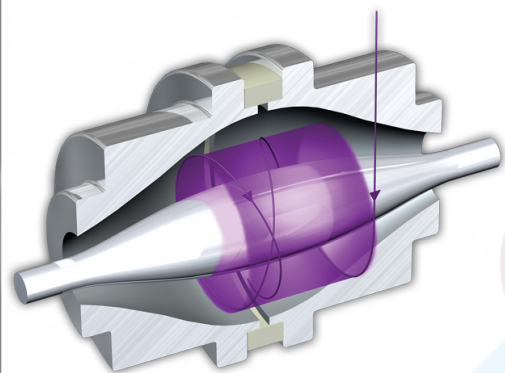
Concentrated extract (50  $\mu\text{L}$ )  
analyzed using GC-Orbitrap  
HRAM

- Full scan
- Targeted SIM





# GC-Q-Exactive



HCD Cell

C-TRAP

AQT  
Quadrupole

Bent  
Flatapole

Orbitrap  
Mass  
Analyzer

ExtractaBrite™  
Ion Source

120K mass resolution  
FWHM (routine at 60K)

Resolving power useful to  
remove co-extracted  
matrix/chemical interferences

Detection sensitivity at ppt  
levels using full scan



# PCB standard (7 pg/ $\mu$ L)

## Method details

1  $\mu$ L injection (SSL injector, 250°C)

Splitless time: 1.5 min

Column: Thermo TG-5Sil MS

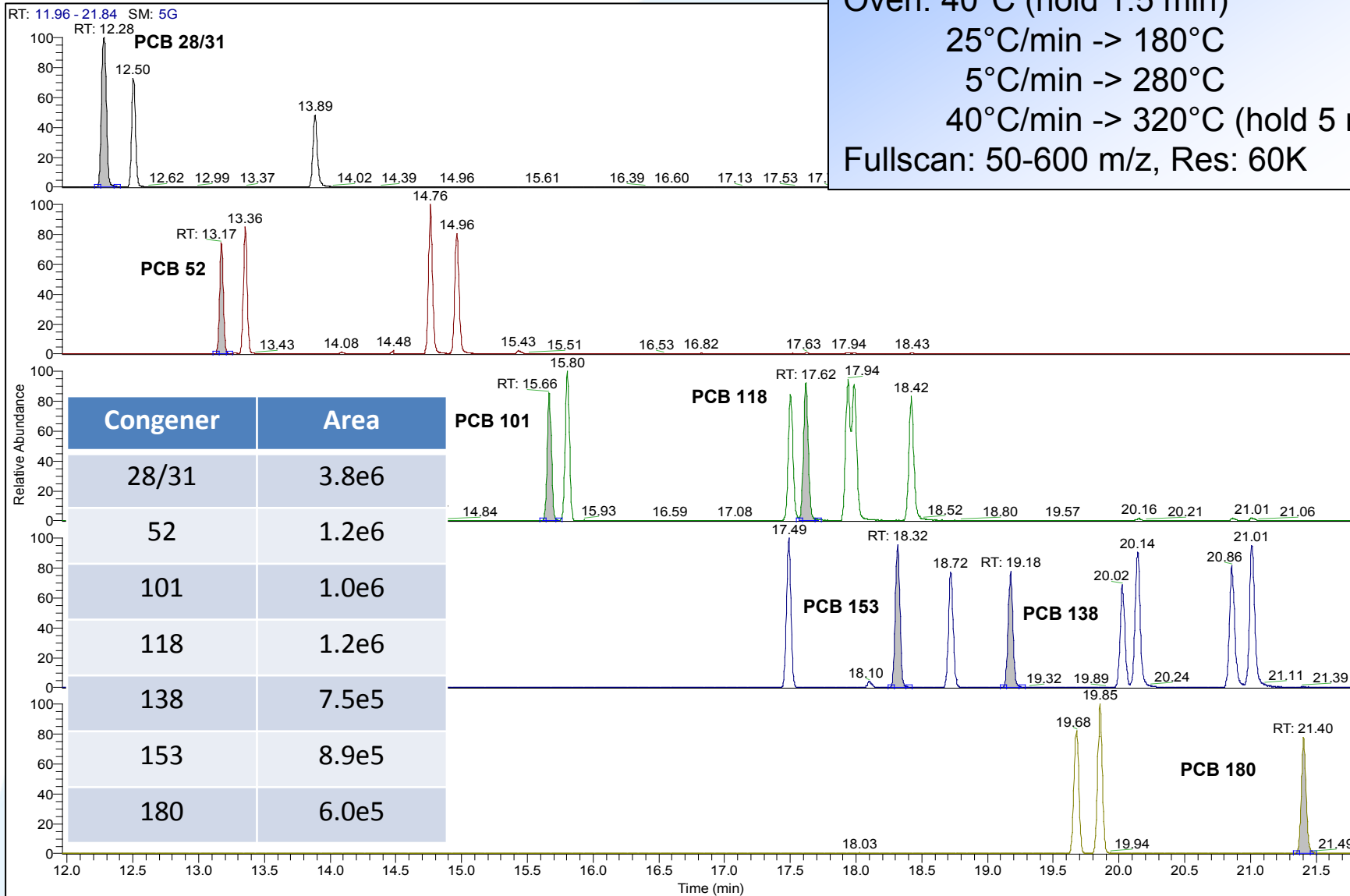
Oven: 40°C (hold 1.5 min)

25°C/min  $\rightarrow$  180°C

5°C/min  $\rightarrow$  280°C

40°C/min  $\rightarrow$  320°C (hold 5 min)

Fullscan: 50-600 m/z, Res: 60K



# Pesticide standard

## Method details

1  $\mu$ L injection (SSL injector, 250°C)

Splitless time: 1.5 min

Column: Thermo TG-5Sil MS

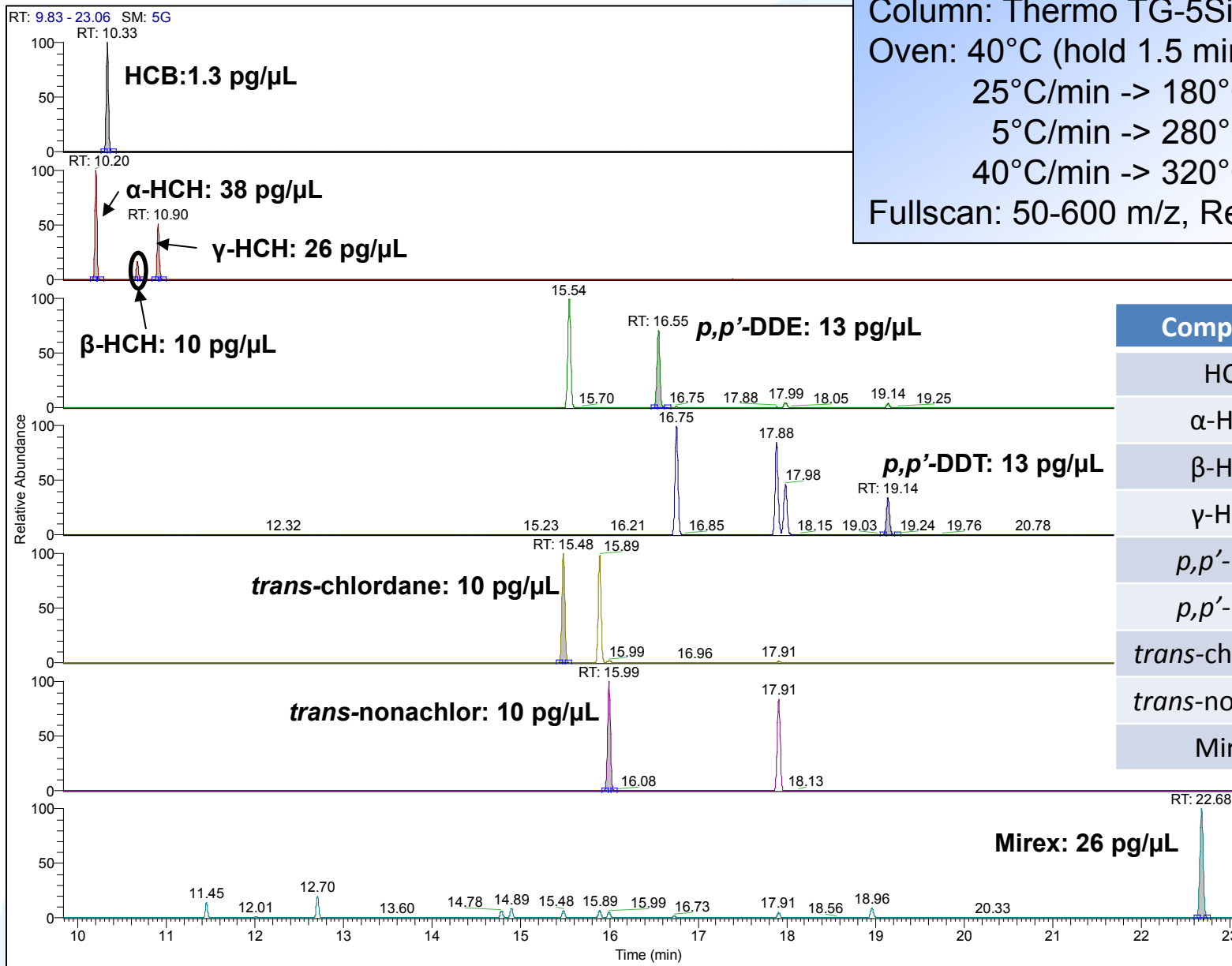
Oven: 40°C (hold 1.5 min)

25°C/min  $\rightarrow$  180°C

5°C/min  $\rightarrow$  280°C

40°C/min  $\rightarrow$  320°C (hold 5 min)

Fullscan: 50-600 m/z, Res: 60K



| Compound                | Area  |
|-------------------------|-------|
| HCB                     | 4.8e5 |
| $\alpha$ -HCH           | 1.0e7 |
| $\beta$ -HCH            | 1.8e6 |
| $\gamma$ -HCH           | 5.5e6 |
| <i>p,p'</i> -DDE        | 4.3e6 |
| <i>p,p'</i> -DDT        | 1.7e6 |
| <i>trans</i> -chlordane | 1.3e6 |
| <i>trans</i> -nonachlor | 1.4e6 |
| Mirex                   | 7.4e6 |

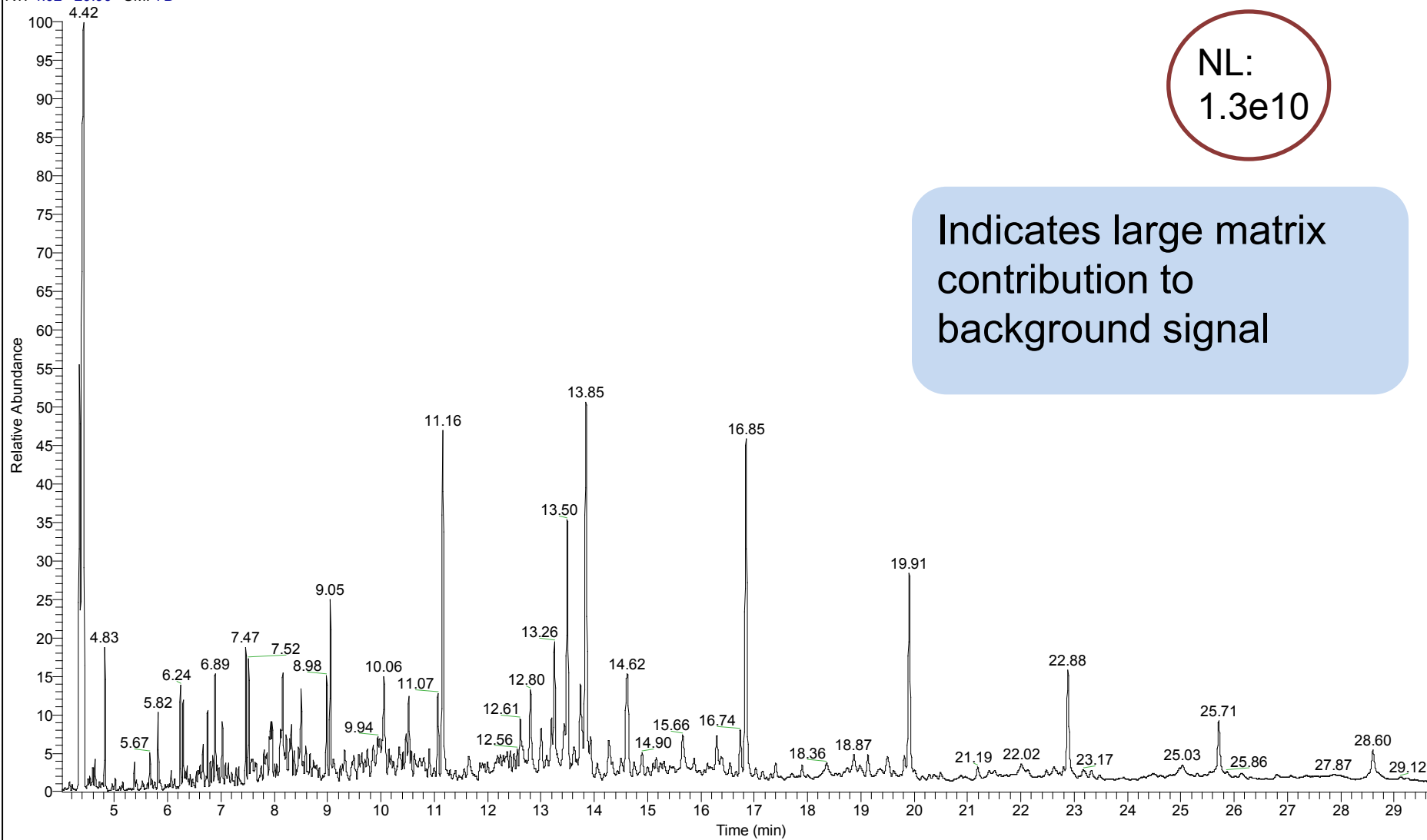


# Antarctic penguin blood sample (full scan)

RT: 4.02 - 29.90 SM: 7B

NL:  
1.3e10

Indicates large matrix  
contribution to  
background signal



# Antarctic penguin blood sample (PCBs)

## Method details

1  $\mu$ L injection (SSL injector, 250°C)

Splitless time: 1.5 min

Column: Thermo TG-5Sil MS

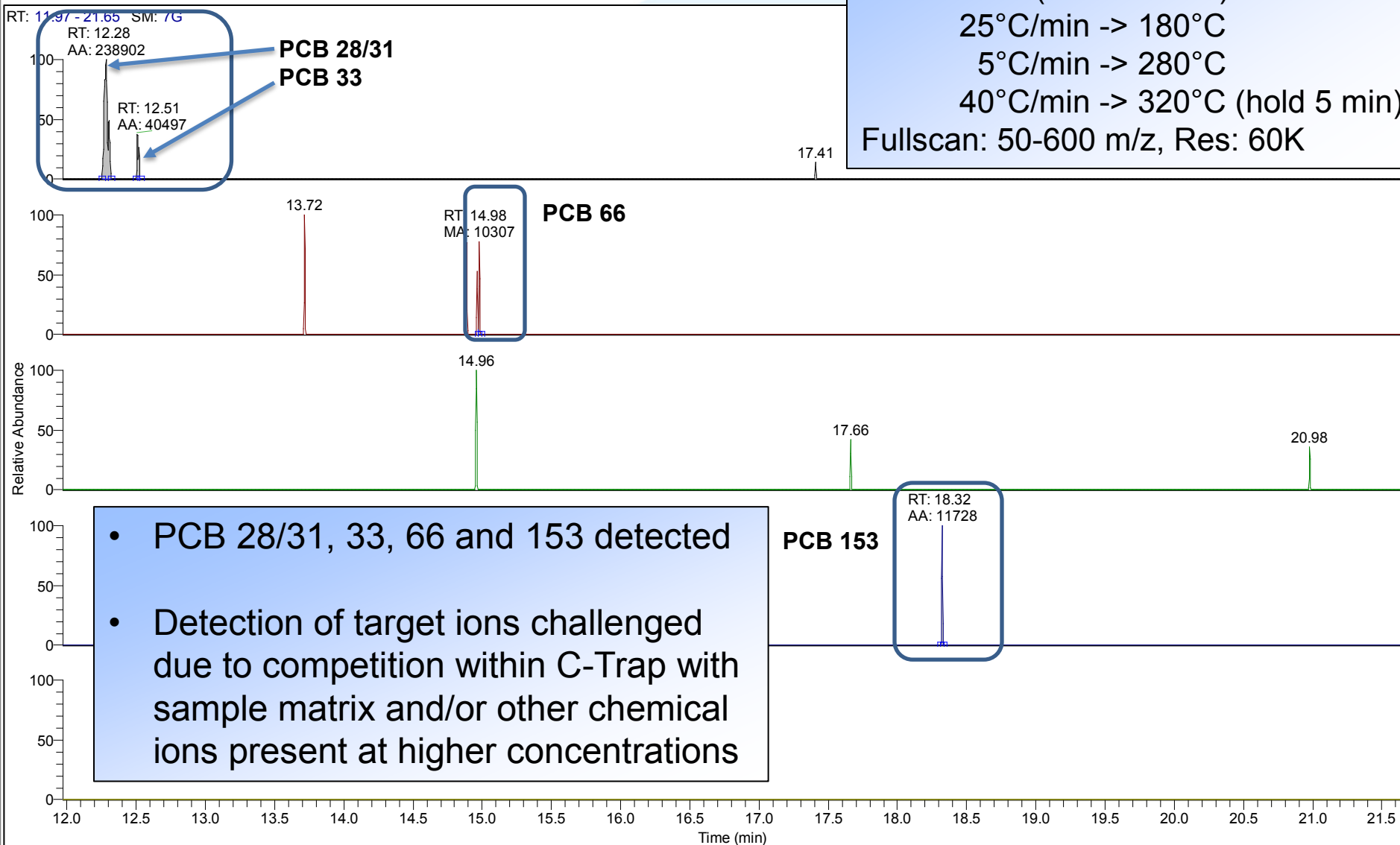
Oven: 40°C (hold 1.5 min)

25°C/min  $\rightarrow$  180°C

5°C/min  $\rightarrow$  280°C

40°C/min  $\rightarrow$  320°C (hold 5 min)

Fullscan: 50-600 m/z, Res: 60K



# Antarctic penguin blood sample (pesticides)

## Method details

1  $\mu$ L injection (SSL injector, 250°C)

Splitless time: 1.5 min

Column: Thermo TG-5Sil MS

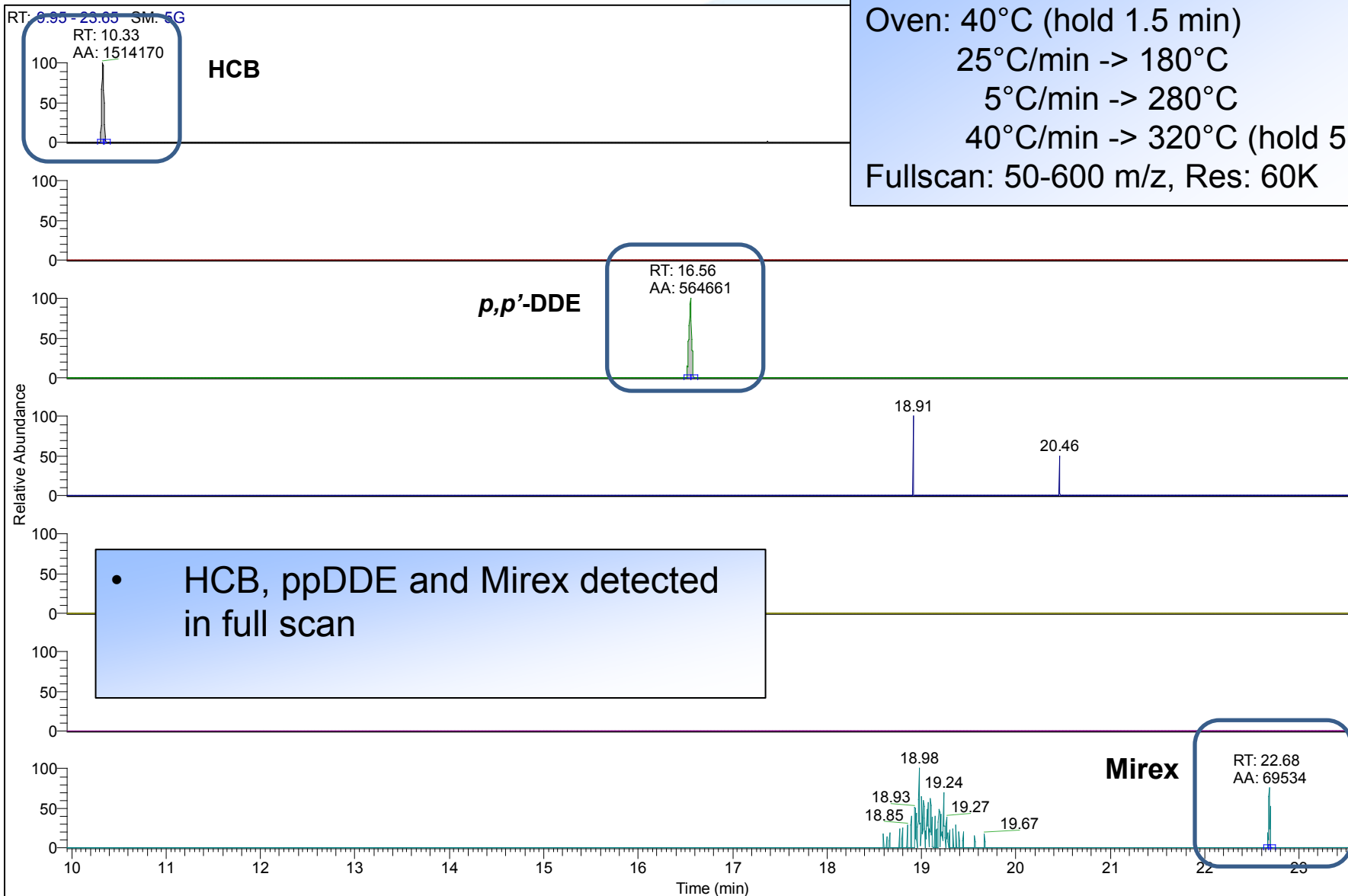
Oven: 40°C (hold 1.5 min)

25°C/min  $\rightarrow$  180°C

5°C/min  $\rightarrow$  280°C

40°C/min  $\rightarrow$  320°C (hold 5 min)

Fullscan: 50-600 m/z, Res: 60K

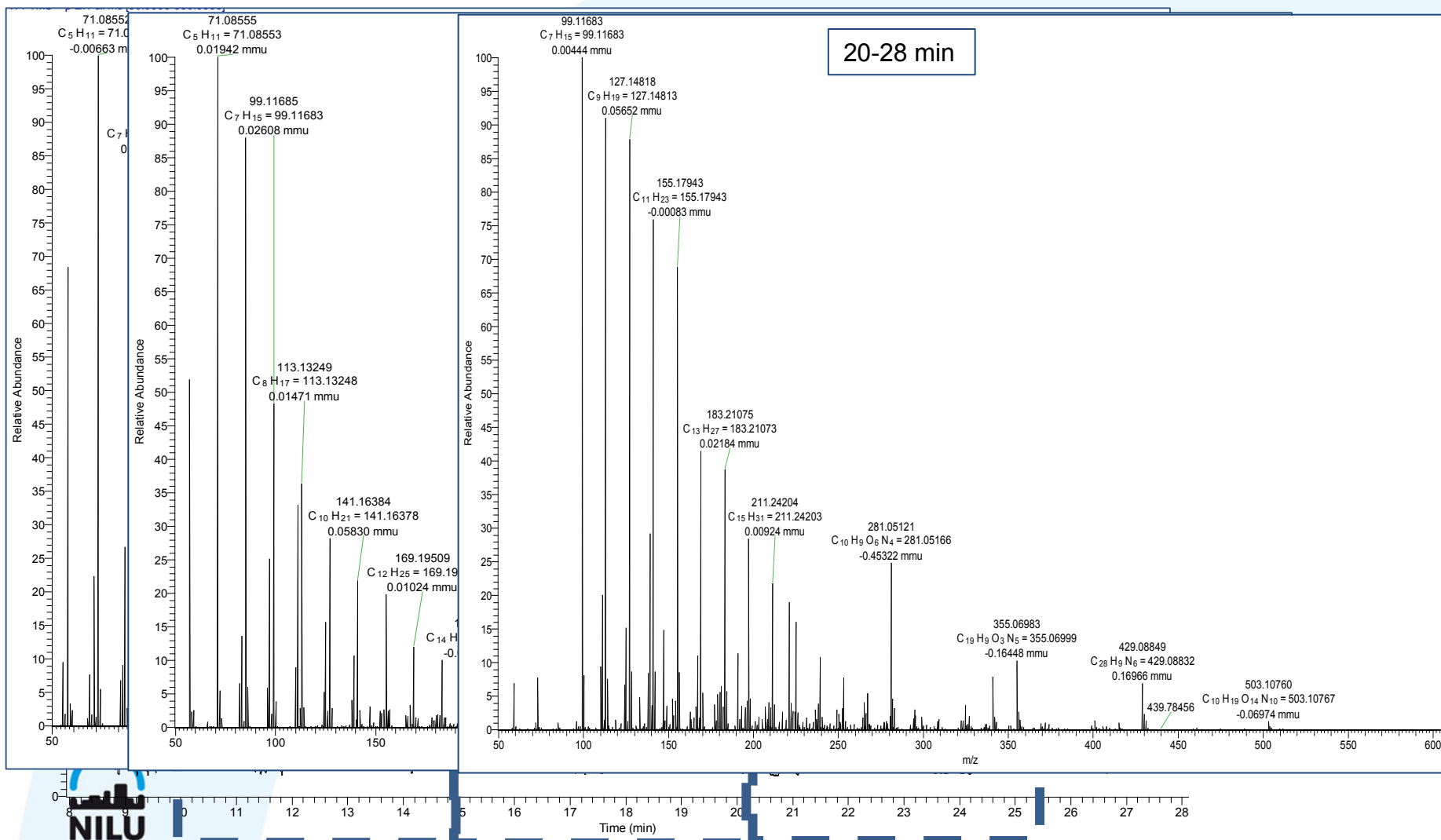


# Antarctic penguin blood sample

10-15 min

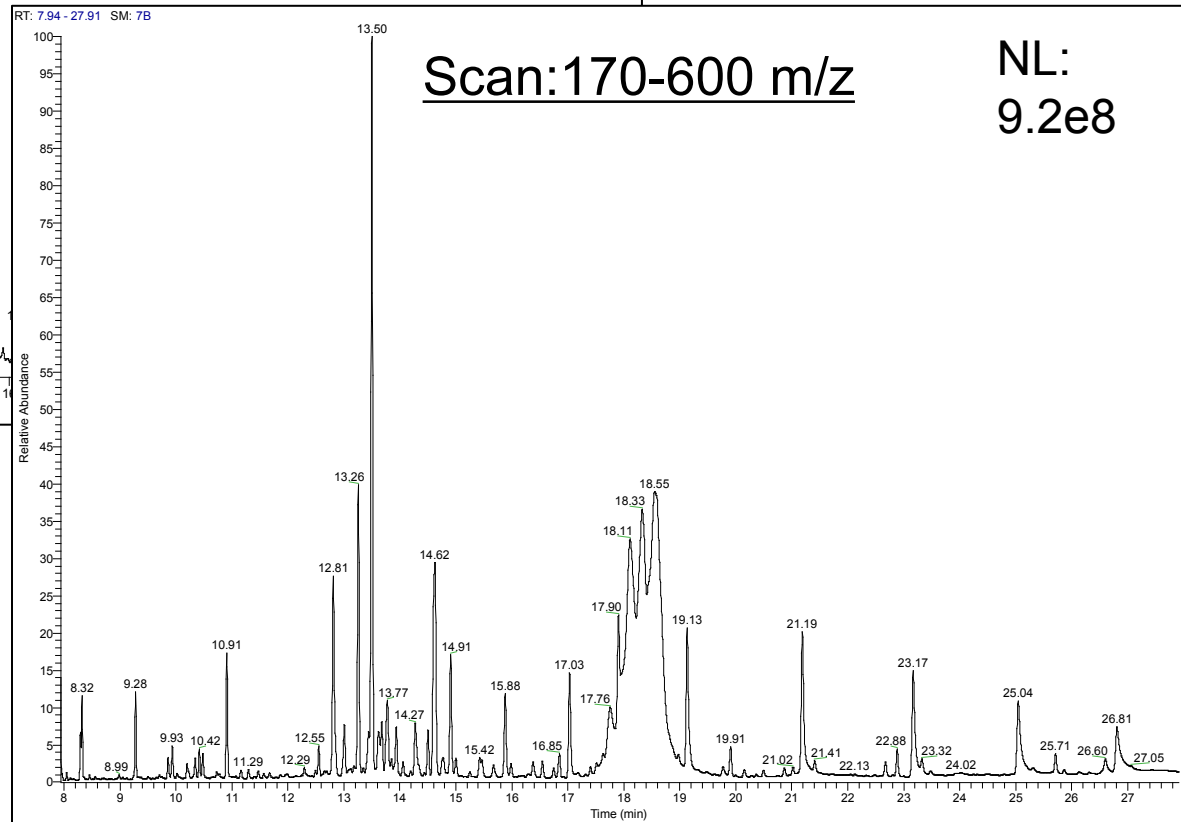
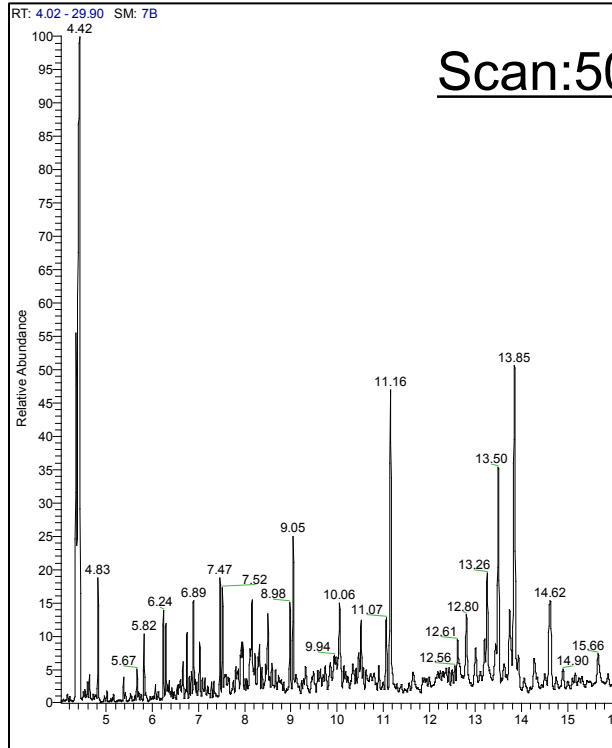
15-20 min

20-28 min

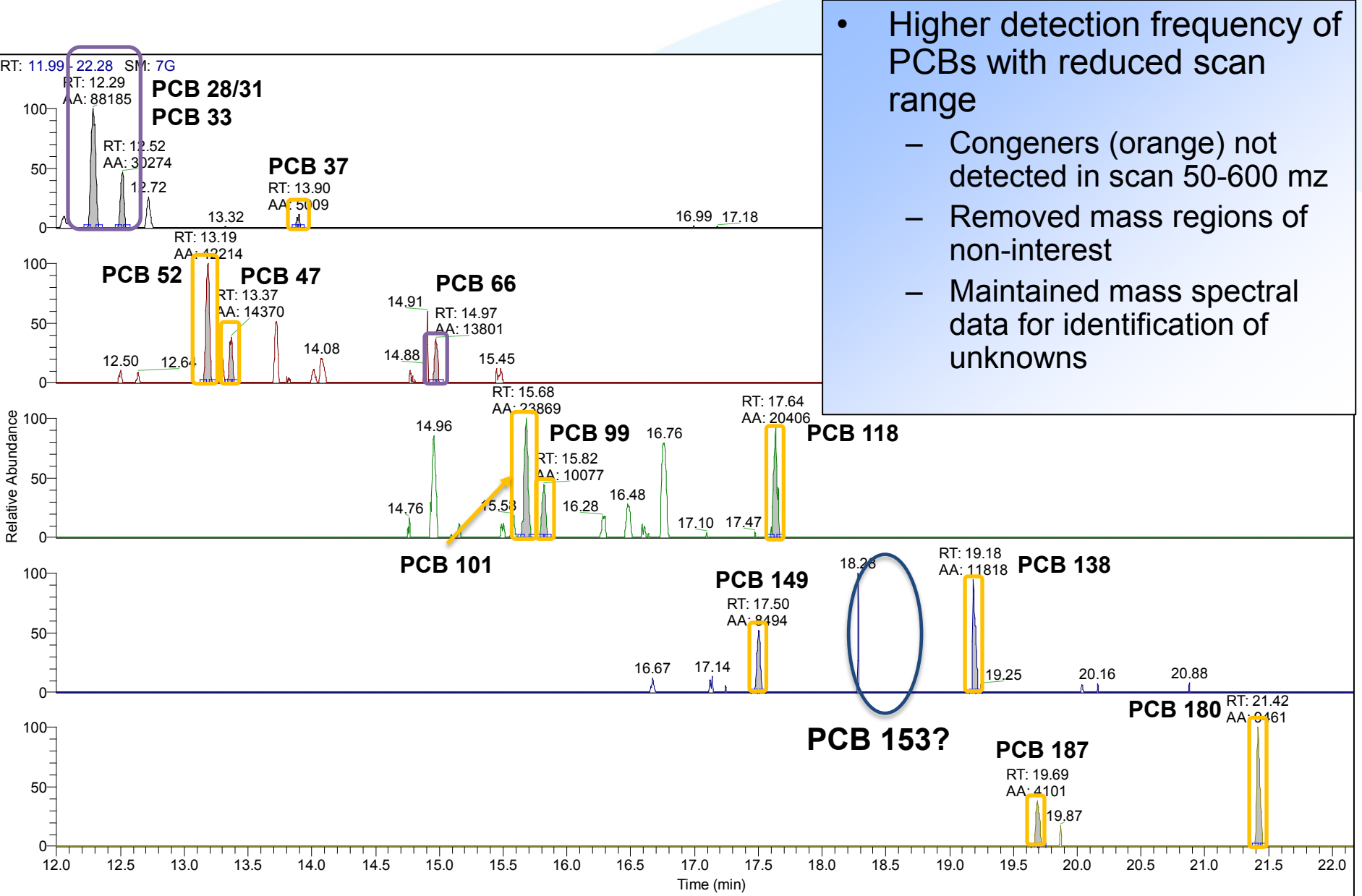




# Full scan vs. reduced full scan

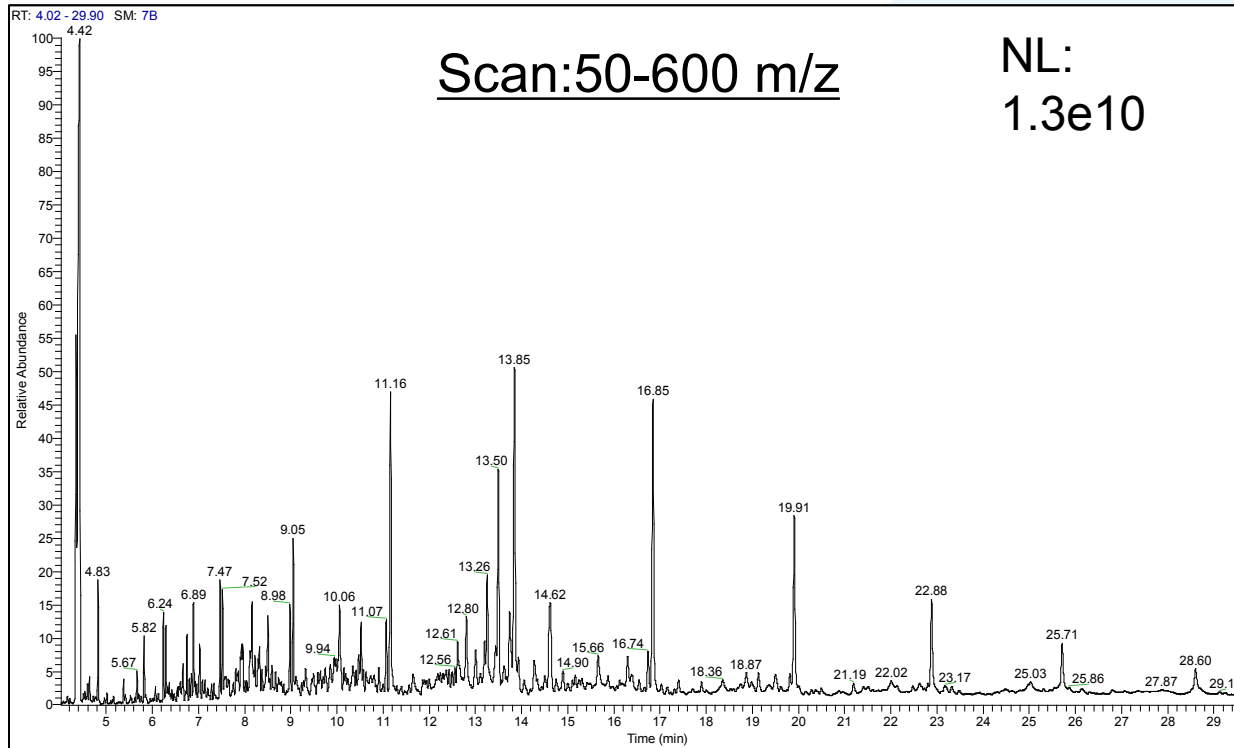


# Antarctic penguin blood sample (PCBs, 170-600 m/z)

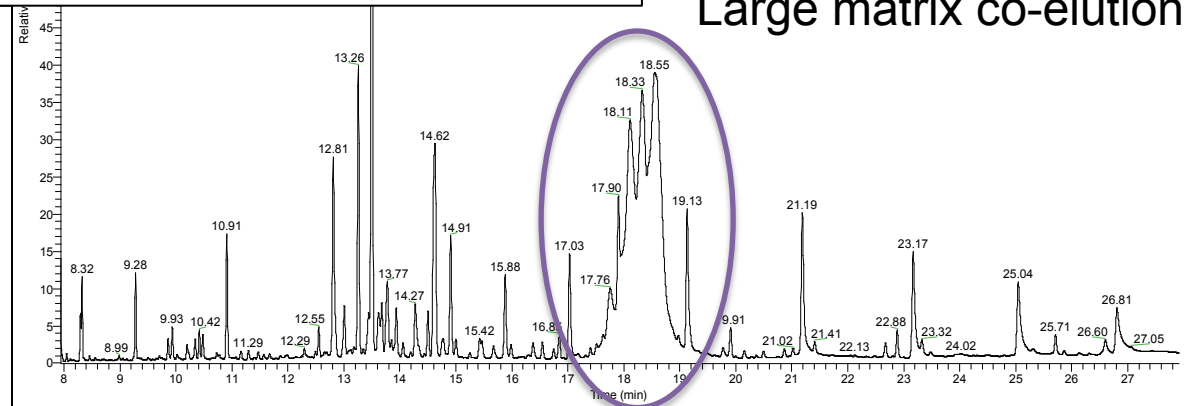


- Higher detection frequency of PCBs with reduced scan range
  - Congeners (orange) not detected in scan 50-600 m/z
  - Removed mass regions of non-interest
  - Maintained mass spectral data for identification of unknowns

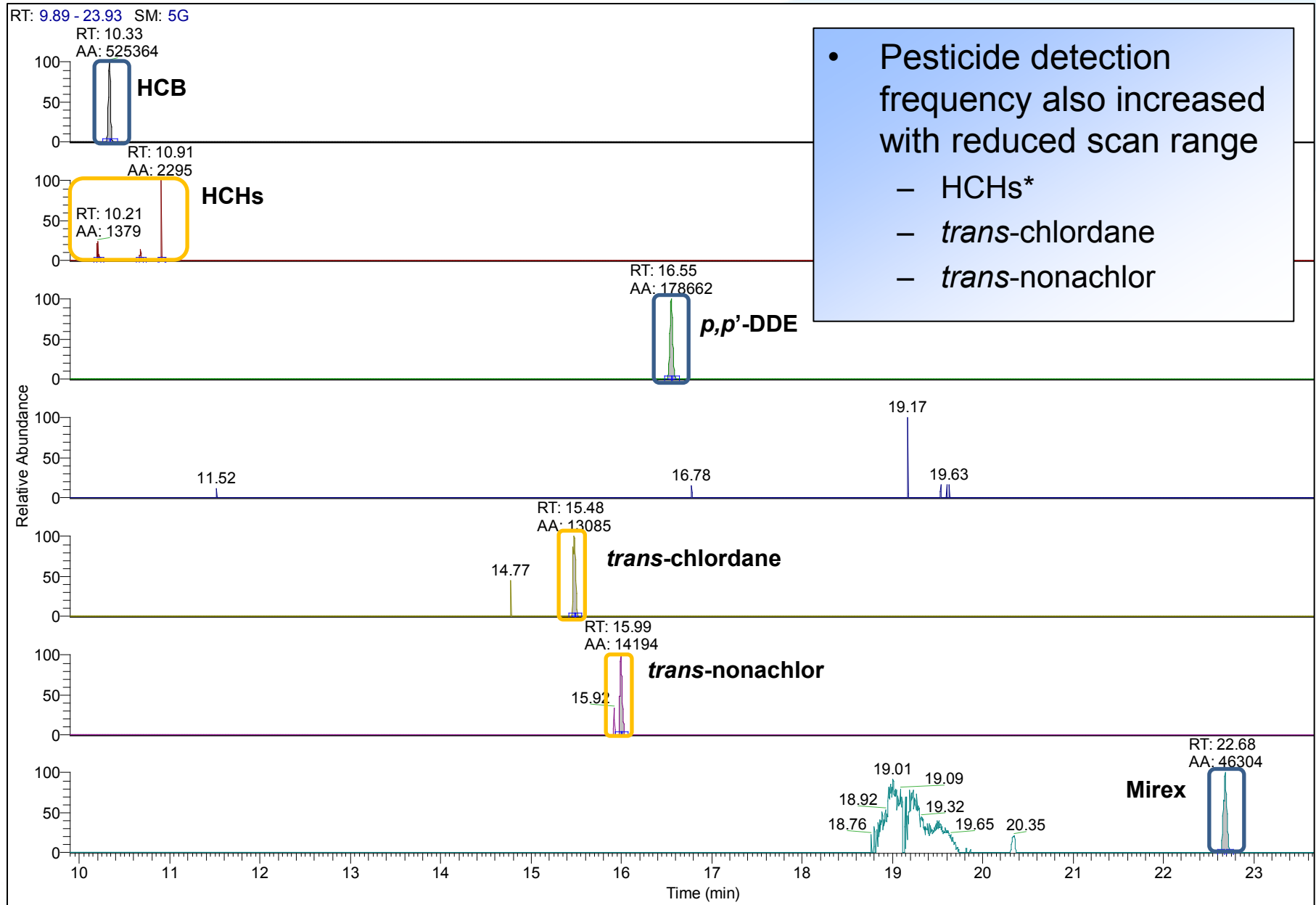
# Full scan vs. reduced full scan



-600 m/z NL: 9.2e8



# Antarctic penguin blood sample (pesticides, 170-600 m/z)





# Antarctic penguin blood sample (Targeted SIM (TSIM))

## Method details

1  $\mu$ L injection (SSL injector, 250°C)

Splitless time: 1.5 min

Column: Thermo TG-5Sil MS

Oven: 40°C (hold 1.5 min)

25°C/min  $\rightarrow$  180°C

5°C/min  $\rightarrow$  280°C

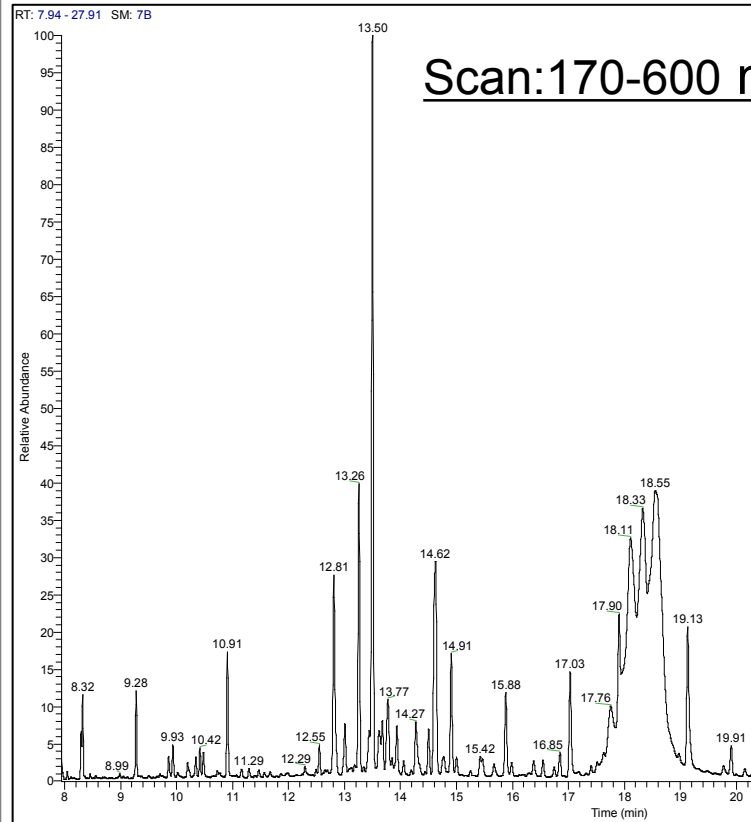
40°C/min  $\rightarrow$  320°C (hold 5 min)

Targeted SIM: 8 m/z scan window,

Resolution: 60K

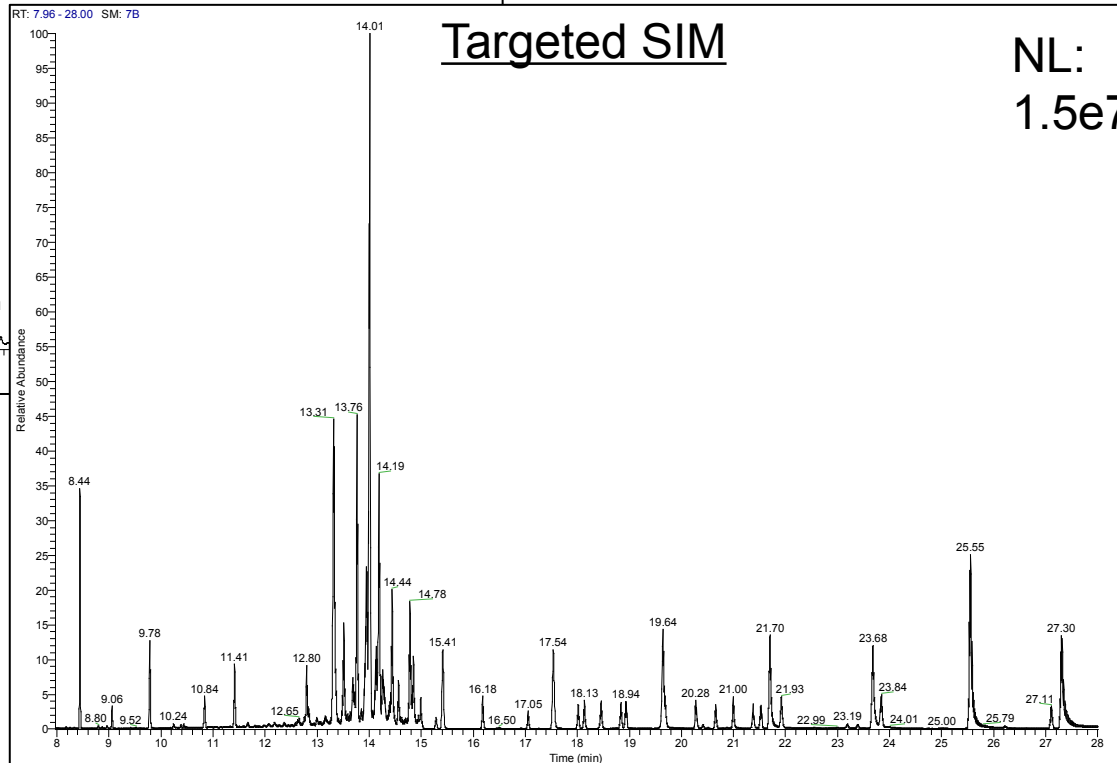
Scan:170-600 m/z

NL:  
9.2e8

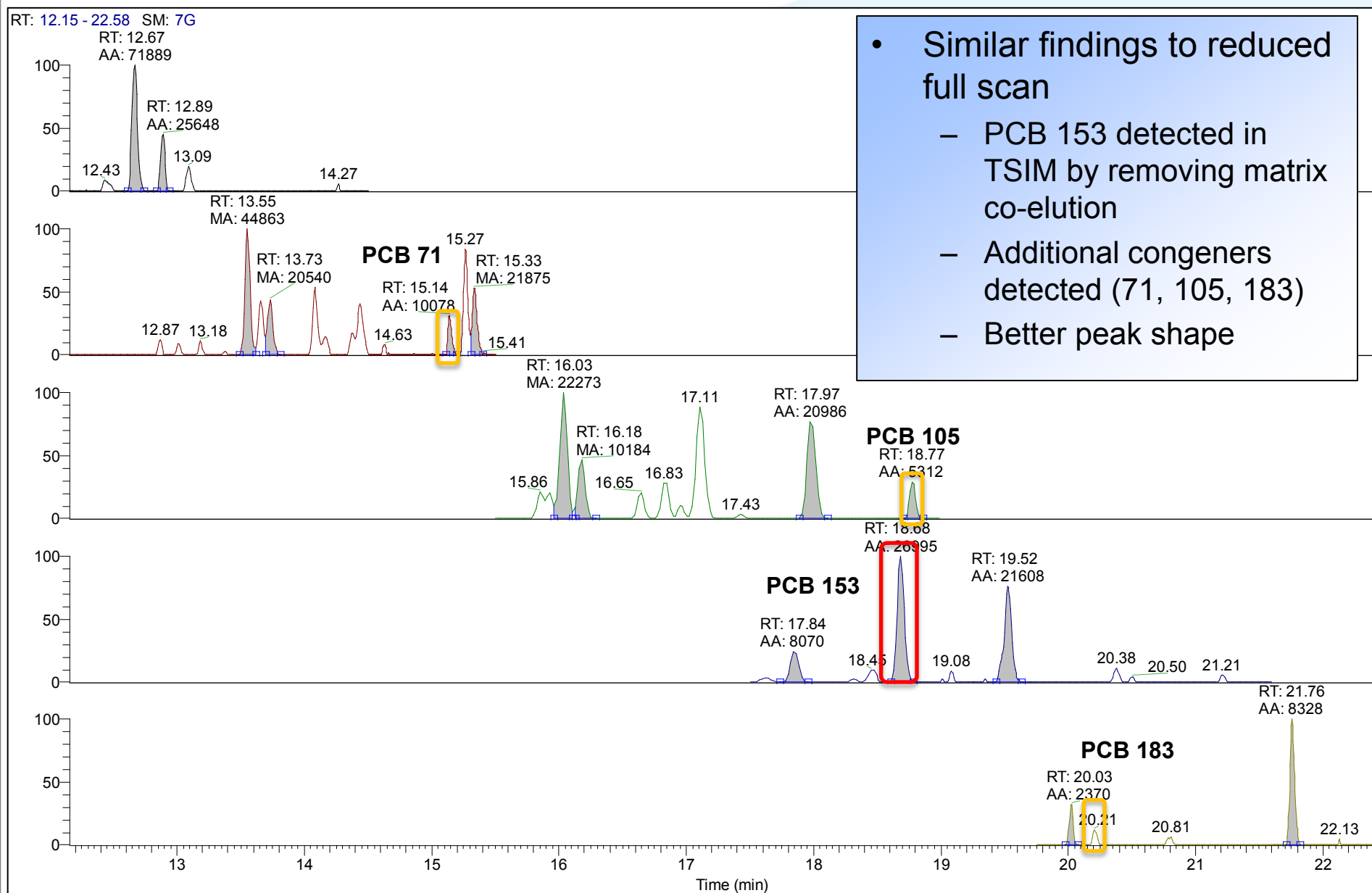


Targeted SIM

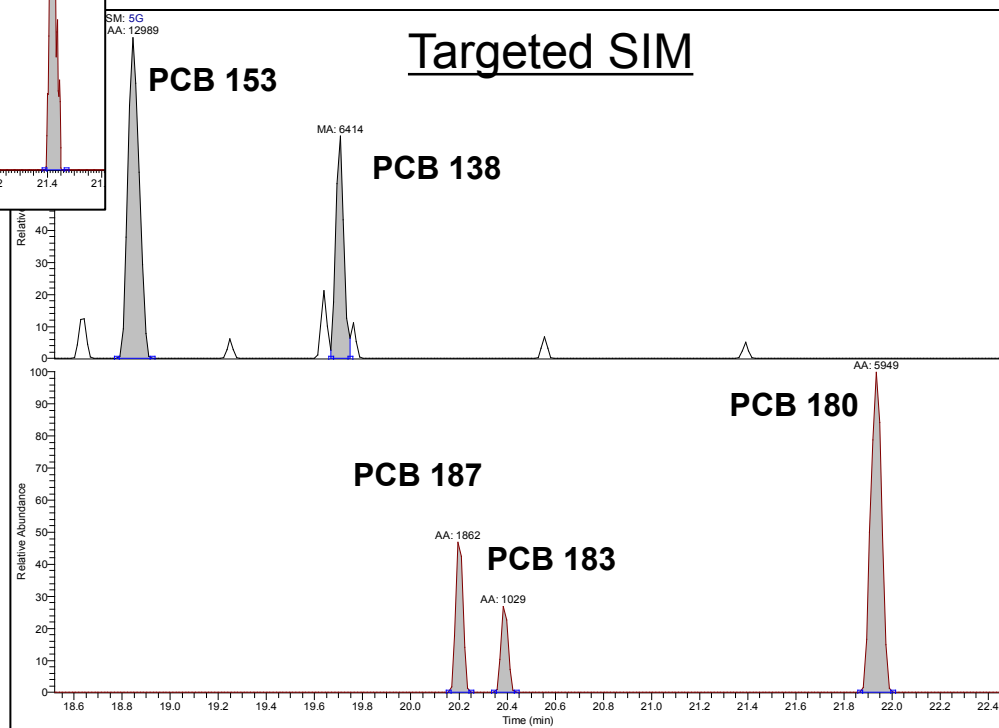
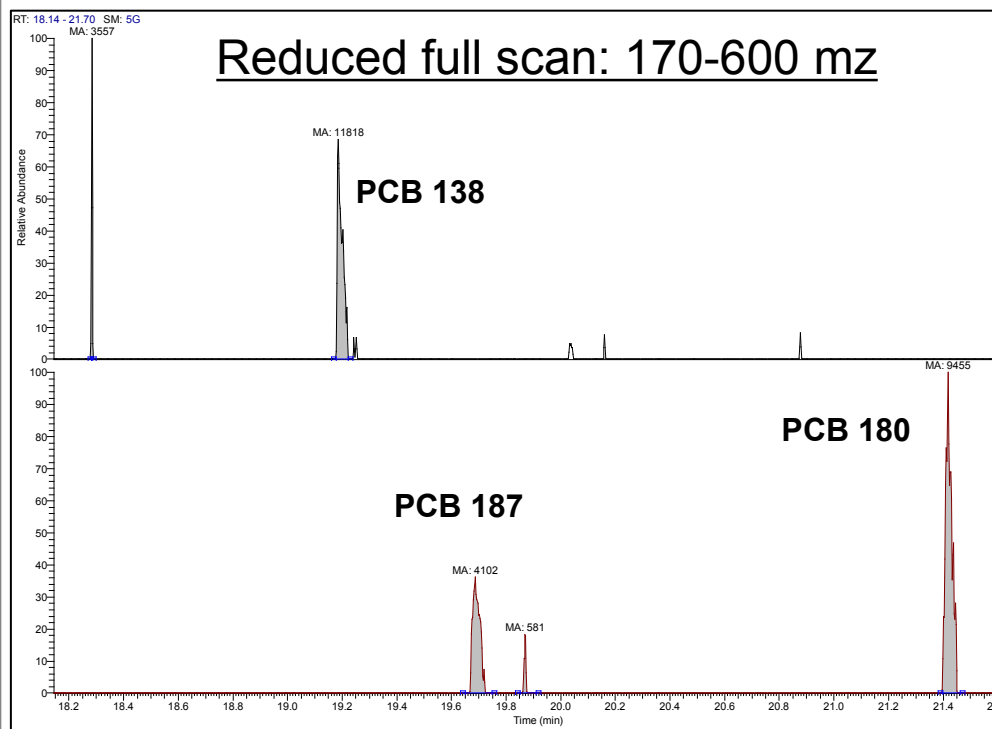
NL:  
1.5e7



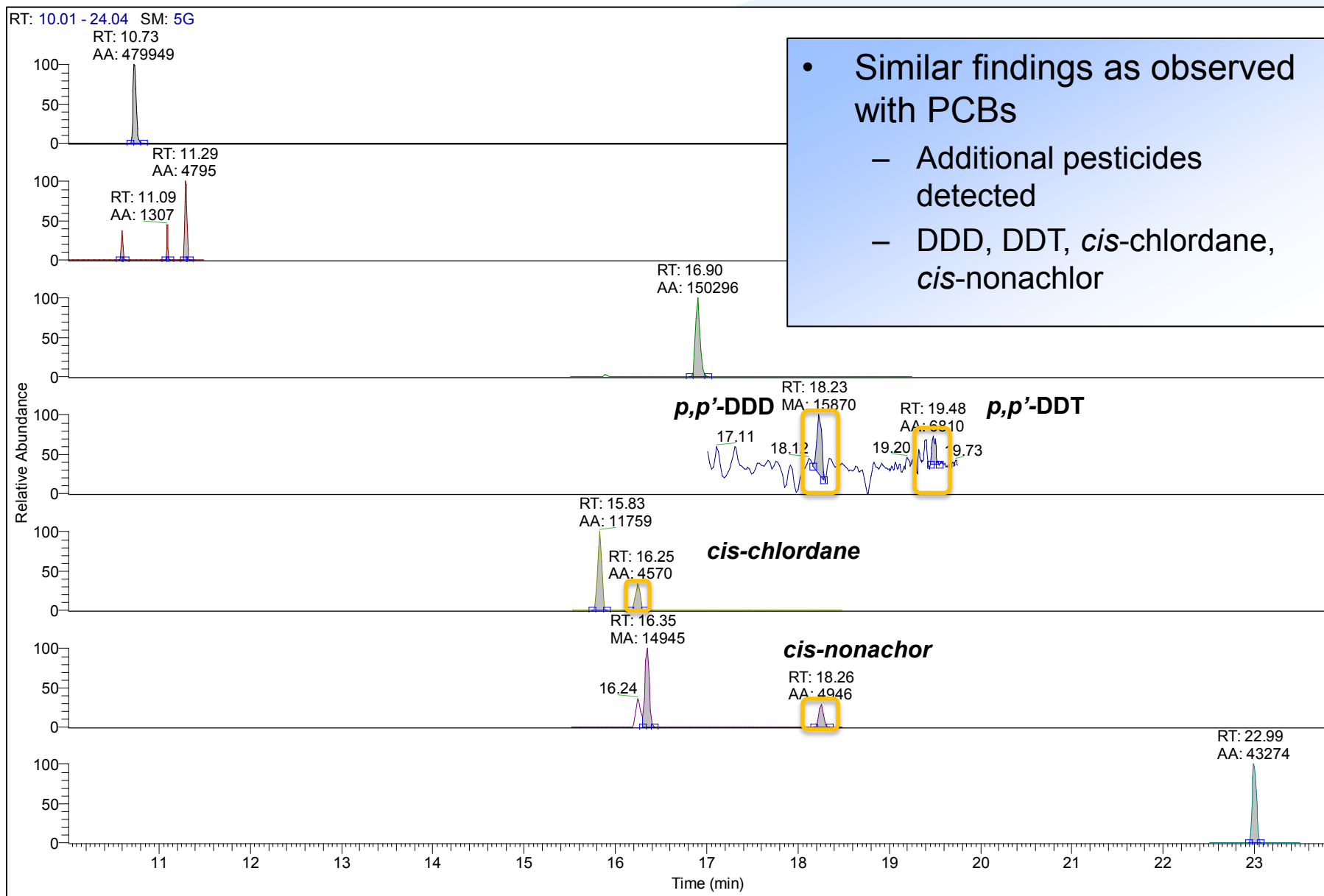
# Antarctic penguin blood sample (PCBs, Targeted SIM)



# Peak shape comparison



# Antarctic penguin blood sample (pesticides, Targeted SIM)



# Simultaneous full scan and targeted SIM

NewNILUPOP\_EL\_SSL\_TSIM30K\_Fullscan60K.meth - Thermo Xcalibur Instrument Setup

File Help

Global Lists

- Lock Masses
- Inclusion
- Exclusion
- Tag Masses

Tune Files

C:\Xcalibur\methods\EL\_70EV\_March\_26\_2018.mstune

Scan Groups

Targeted-SIM

Full MS - SIM

time (min)

0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30

time (min)

0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30

Experiments

General

- Full MS - SIM
- Full MS / dd-MS<sup>2</sup> (TopN)
- Targeted-SIM
- PRM
- Targeted-SIM / dd-MS<sup>2</sup>

Targeted SIM acquisition at 30K mass resolution

Full scan acquisition at 60K mass resolution

Can obtain targeted analysis specificity without loss of full scan mass spectral data for identification of unknown compounds

Properties of the method

Global Settings

- User Role: Standard
- Use lock masses: best

Time

- Method duration: 30.00 min

EI/CI Source

- Filament on delay: 7.00 min
- MS transfer line: 280 °C
- Ion source temp: 250 °C
- Ionization mode: EI
- CI gas type: None
- CI gas flow: 0.00 mL/min

Method duration

Duration of the method

Properties of Tunefiles

General

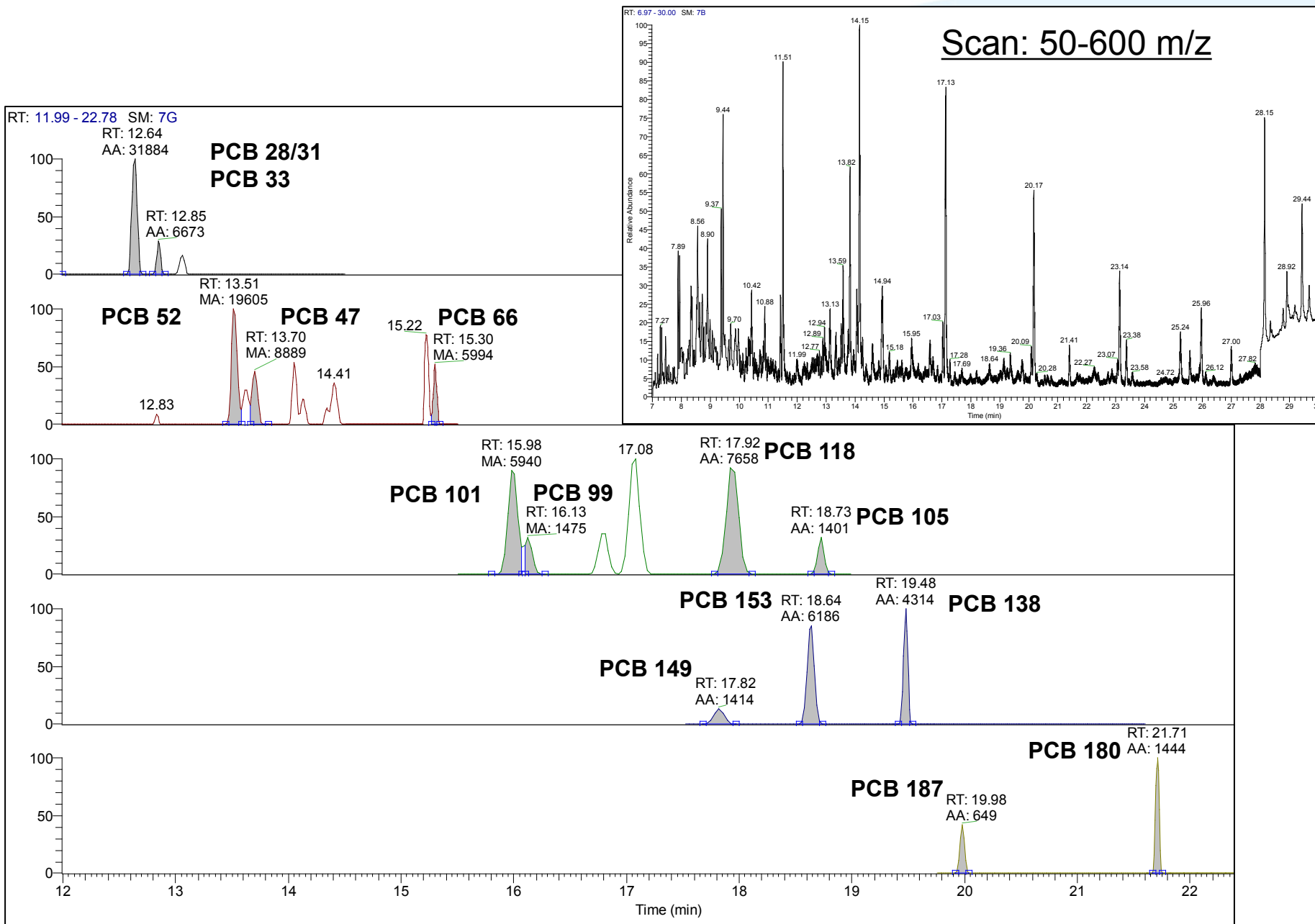
- Switch Count: 0
- Base Tunefile: C:\Xcalibur\methods\EL

Switch Count

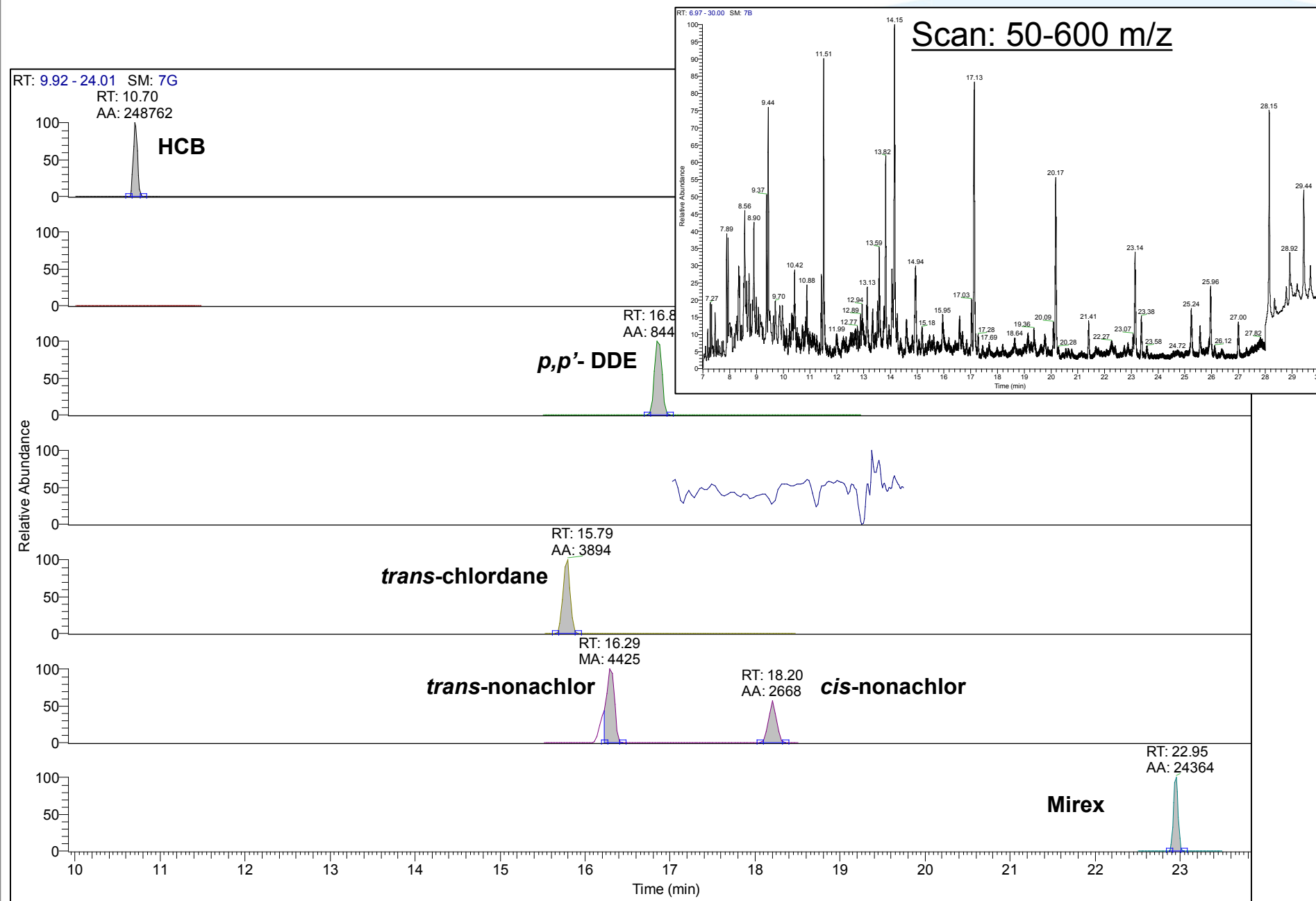
Select the number of tune file switches. Range 0-50



# Antarctic penguin blood (Target SIM/Full scan)



# Antarctic penguin blood (Target SIM/Full scan)



# OC concentrations in penguin blood sample

| PCB congener | Concentration (pg/g ww) |
|--------------|-------------------------|
| 52           | 11.5                    |
| 101          | 16.3                    |
| 118          | 6.4                     |
| 138          | 13.2                    |
| 149          | 3.8                     |
| 153          | 11.6                    |
| 180          | 5.3                     |
| 183          | 1.4                     |
| 187          | 2.0                     |

| Pesticide               | Concentration (pg/g ww) |
|-------------------------|-------------------------|
| HCB                     | 290                     |
| aHCH                    | 3.9                     |
| <i>p,p'</i> -DDT        | 1.38                    |
| <i>p,p'</i> -DDD        | 4.72                    |
| <i>p,p'</i> -DDE        | 101                     |
| <i>trans</i> -chlordane | 11.7                    |
| <i>cis</i> -chlordane   | 3.5                     |
| <i>trans</i> -nonachlor | 11.7                    |
| <i>cis</i> -nonachlor   | 0.9                     |
| Mirex                   | 38                      |

- Blank subtraction performed on sample results
- LOQ: 0.1 – 14.2 pg/g ww (compound dependent)

GC-Q-Exactive provides  
excellent sensitivity for  
OC analysis in penguin  
blood

Sample matrix can  
hinder detection at  
ppt concentrations

PCBs and OC  
pesticides  
detected at ppt  
concentrations in  
full scan and TSIM

Targeted SIM can  
reduce influence of  
matrix ions in C-Trap

HCB, ppDDE,  
Mirex dominate in  
penguin blood

Full scan data acquired  
simultaneously without  
compromising targeted  
analysis





A wide-angle photograph of a vast, icy Antarctic landscape. In the foreground, a large, white ice formation slopes down towards a body of dark blue water. A small group of penguins is visible on the ice slope. The background shows more ice formations under a clear sky.

# Versatility of GC-Orbitrap mass spectrometry for the ultra-trace detection of persistent organic pollutants in penguin blood from Antarctica

## Authors

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Cristian I. Cojocariu<sup>2</sup>

<sup>1</sup>NILU-Norwegian Institute for Air  
Research, Fram Centre, Tromsø,  
Norway, <sup>2</sup>Thermo Fisher Scientific,  
Runcorn, United Kingdom

## Goal

In this study, the performance of the Thermo Scientific™ Q Exactive™ GC Orbitrap™ mass spectrometer was evaluated for routine analysis of POPs within King penguin blood from Antarctica.

## Introduction

Persistent organic pollutants (POPs) have been studied extensively over the past five decades due to their toxicity and environmental persistence

# Acknowledgements

**Cristian Cojocariu (Thermo Scientific)**

**Strategic Institute Funding grant (CleanArctic, NRC)**

**POLAR-ECOTOX project (NRC)**



**Thank you for your attention**