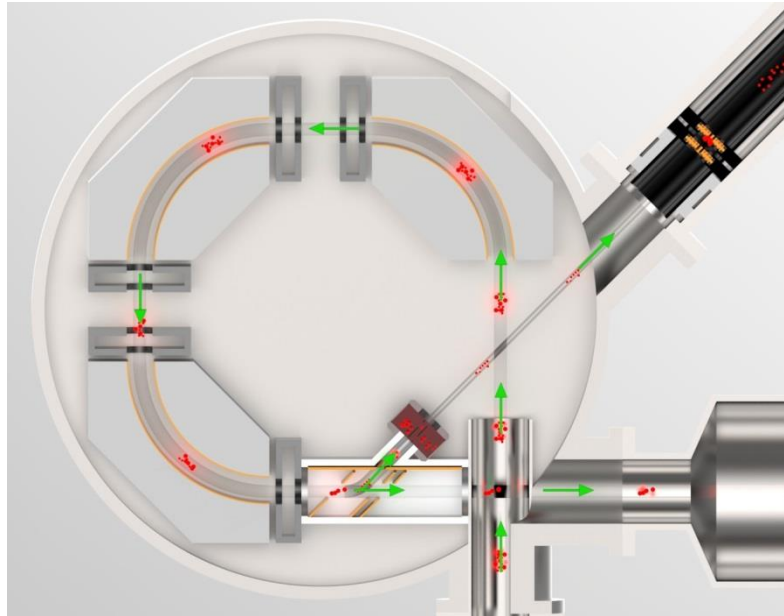


Where do Drug Molecules go Inside of Cells?

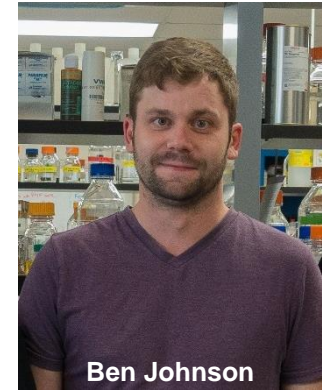
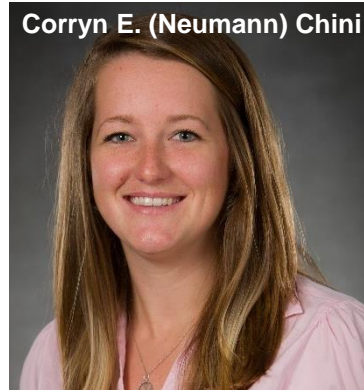
A New Method to Probe the Composition of Cellular Organelles



Ashley Ellsworth
Physical Electronics, USA.

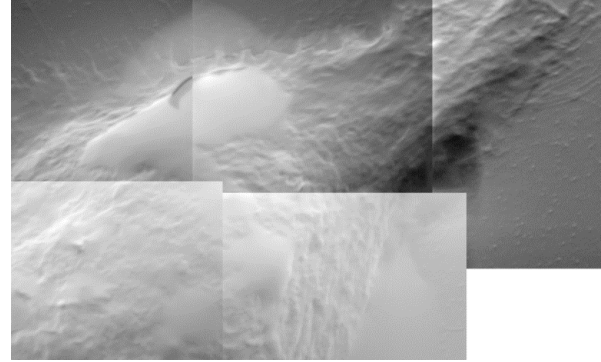
Collaborators

*Physical Electronics,
University of Illinois at Urbana-Champaign,
& Colorado State University*

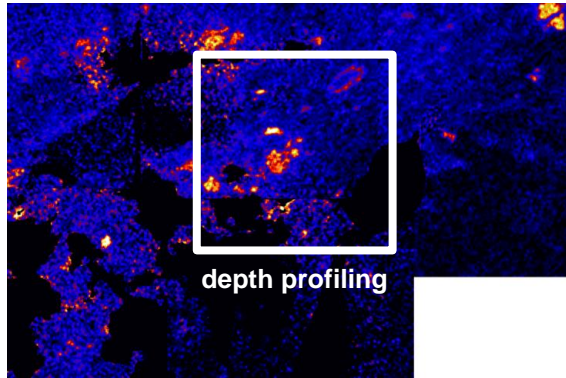


MS 2D Mapping of Lipids & Sterols on MDCK Cell

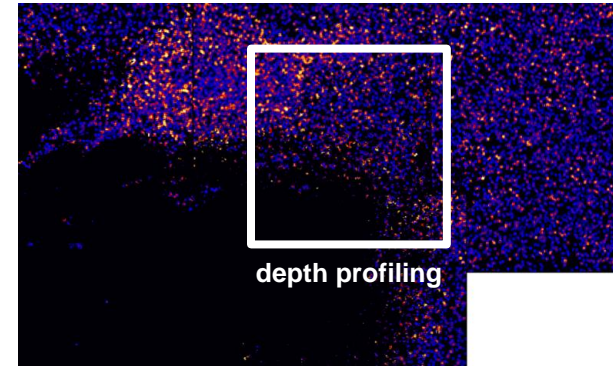
Ion-Induced SE Image of MDCK Cell



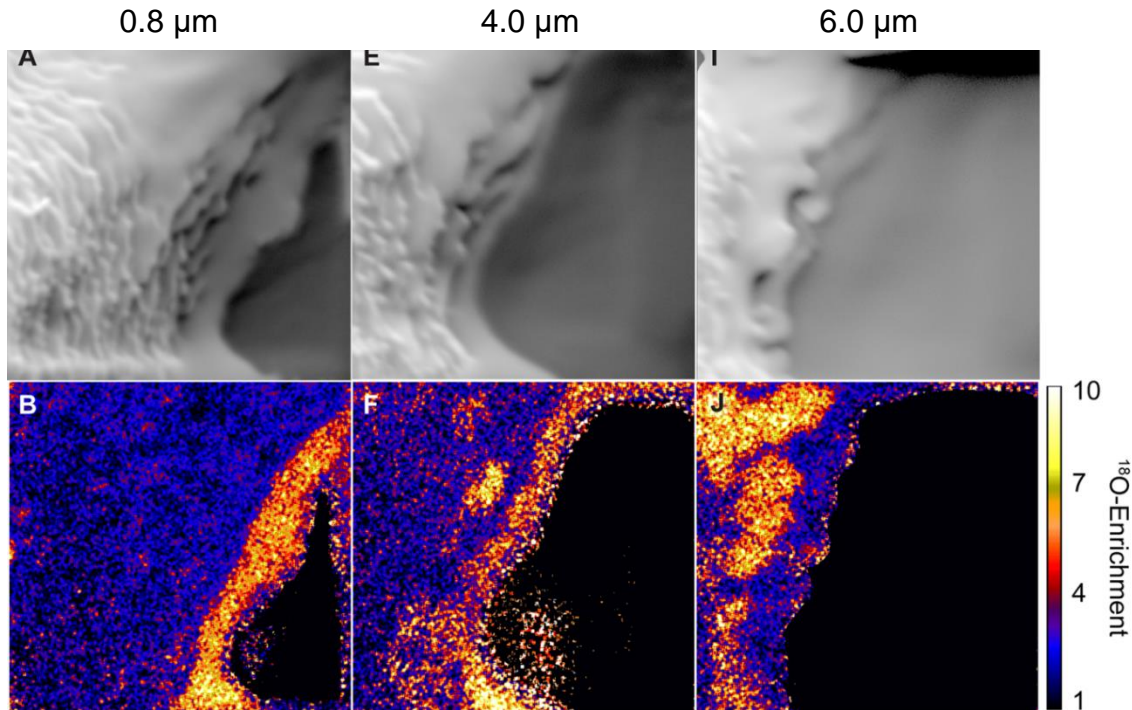
^{15}N -sphingolipids



^{18}O -cholesterol



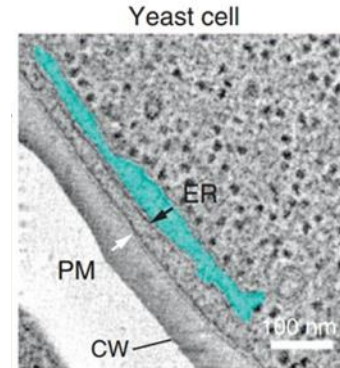
MS 3D Mapping of Lipids & Sterols in MDCK Cell



Are the cholesterol-rich regions part of the endoplasmic reticulum, or an organelle involved in cholesterol synthesis and transport??

Identifying specific cellular compartments or structures by MS imaging isn't easy!!

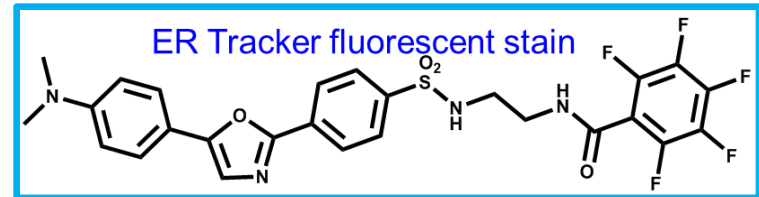
Organelle Labeling: Transfected & Stained HEK Cells



yeast cells



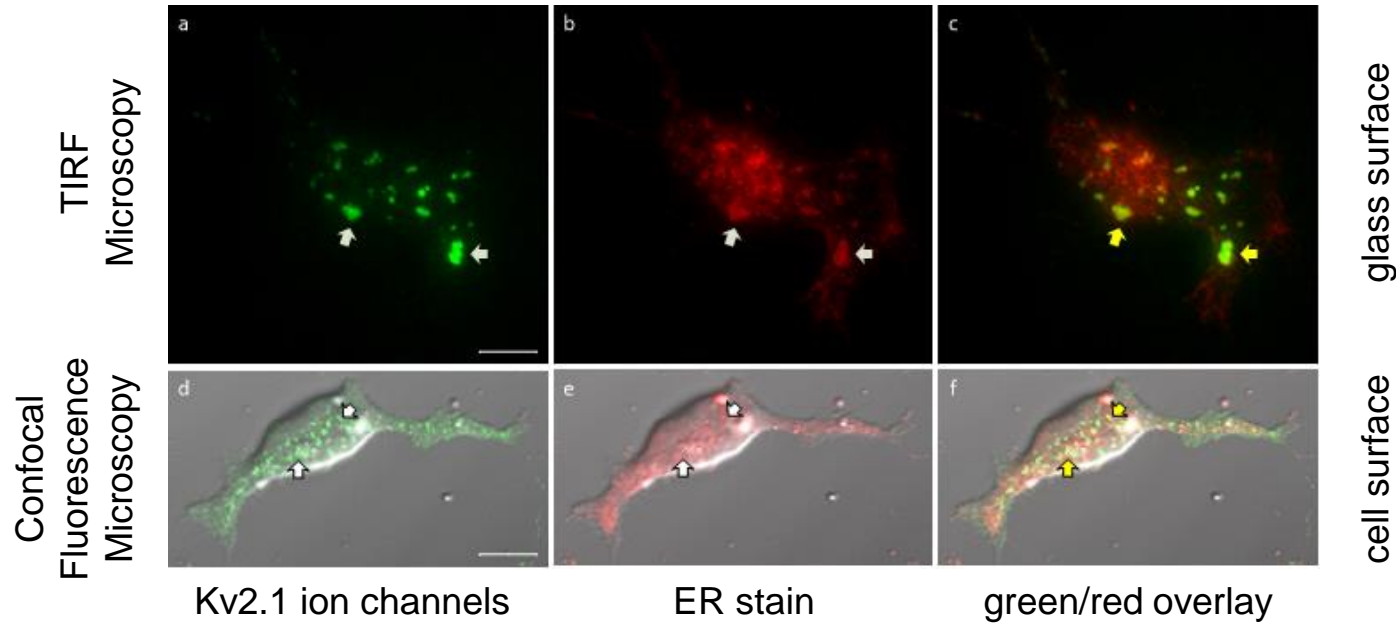
Can we repurpose the ER-Tracker stain for imaging of the endoplasmic reticulum by MSI?



HEK-293 cells were transfected and then stained using the ER-Tracker, and then fixed for analysis.

TIRF and Confocal Fluorescence Microscopy of HEK Cells

The arrows indicate coincidence of the Kv2.1 ion channels and the ER tubules in the stained HEK cells.



The ER tubules measure approximately 0.3 to 2.0 microns in diameter.

MS imaging & tandem MS imaging of HEK cells was performed using a
PHI *nanoTOF* II TOF-SIMS Parallel Imaging MS/MS instrument



3D MS¹ / MS² Imaging

analysis beam

30 keV Bi₃⁺, unbunched
3.0 nA, 1.16 x10¹³ ions/cm²
80 nm beam, 137 nm/pixel

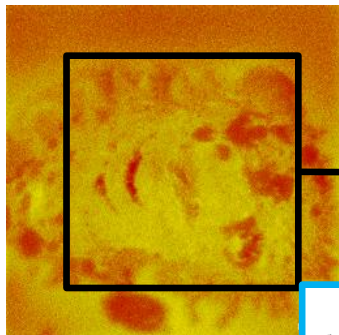
dc sputter beam

5 keV Ar_{2,500}⁺, 6.0 nA,
8.06 x10¹⁴ ions/cm² (total)

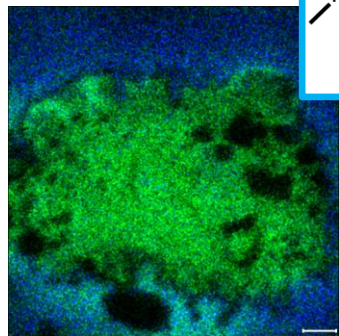
Potential MS¹ Observation of ER Tubules

No molecular ion signals (i.e. $\pm m/z$ 580) of the ER-Tracker stain were observed.

100 μm ; 5 min.



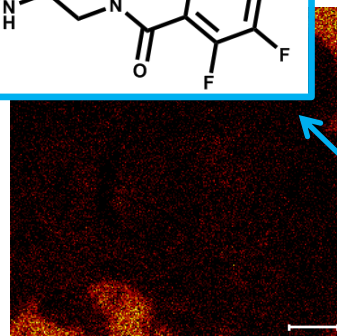
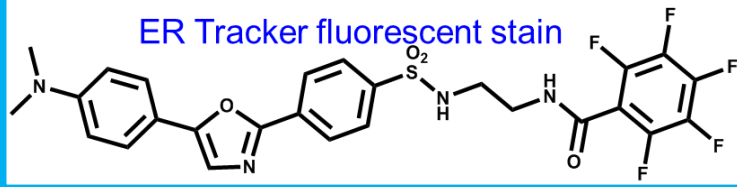
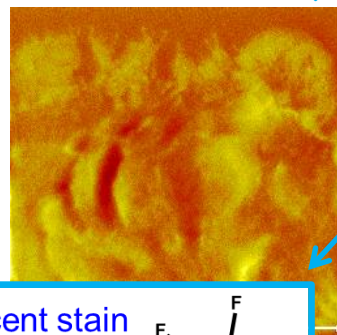
TIC (+SIMS)



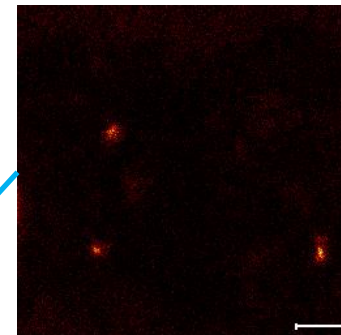
Si^+ (m/z 28); $\text{C}_5\text{H}_{15}\text{NPO}_4^+$ (m/z 184)

?

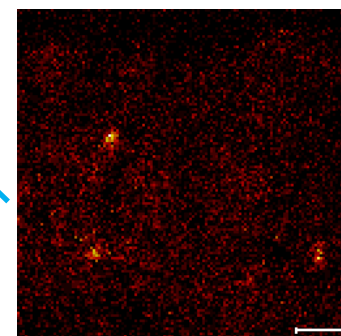
70 μm ; 8 min.



SiO_3H^- (m/z 77)




F^- (m/z 19)



$-m/z$ 167

PHI *nanoTOF* II Standard TOF-SIMS Mode



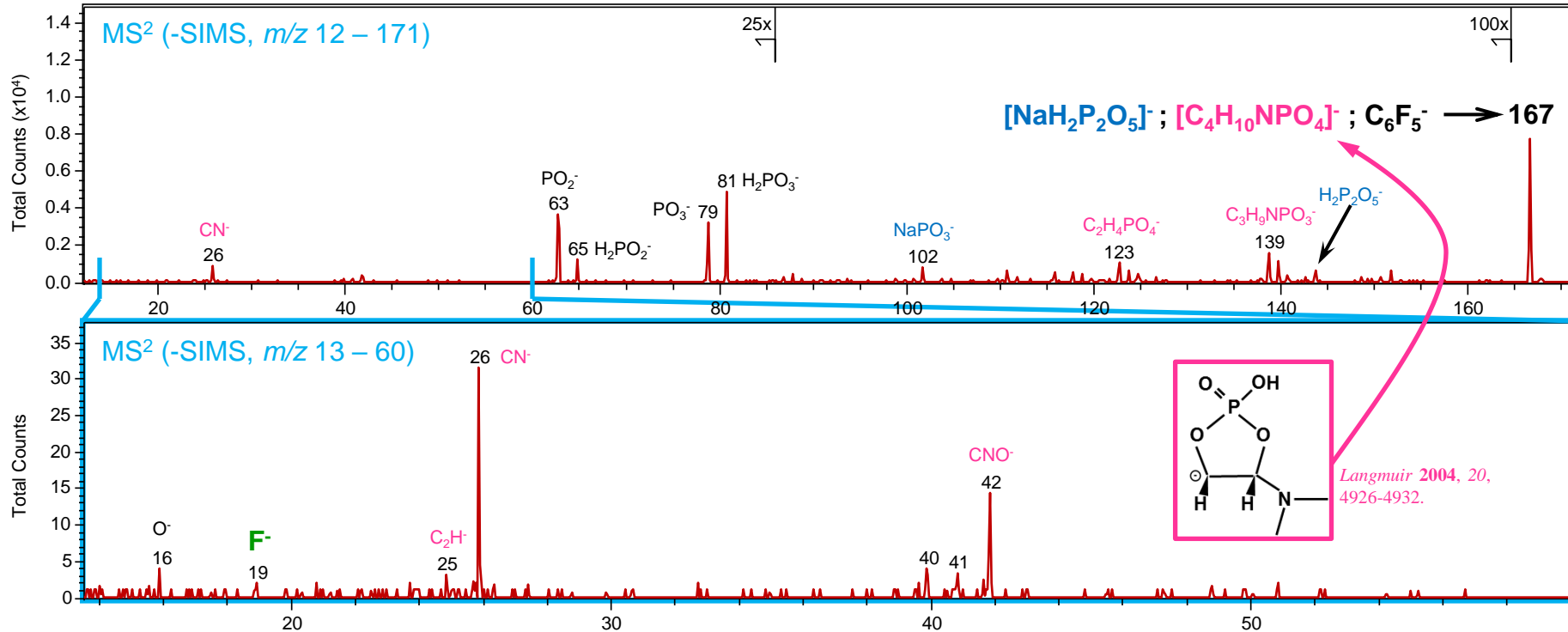
In the standard TOF-SIMS operation mode, all ions are collected at the MS1 detector.

Parallel TOF-SIMS and MS/MS

In MS/MS mode, the precursor ion is deflected into the collision induced dissociation (CID) cell. The resulting fragment ion spectrum is collected with the MS2 detector while the rest of the ions are collected as usual with the MS1 detector.

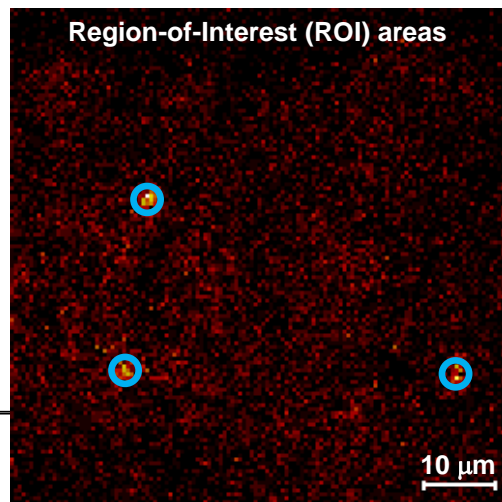
Total Area Tandem MS in Transfected & Stained Cells

MS² spectra reveal exogenous F⁻ in the ER-Tracker-stained cells, potentially from C₆F₅⁻ ions.

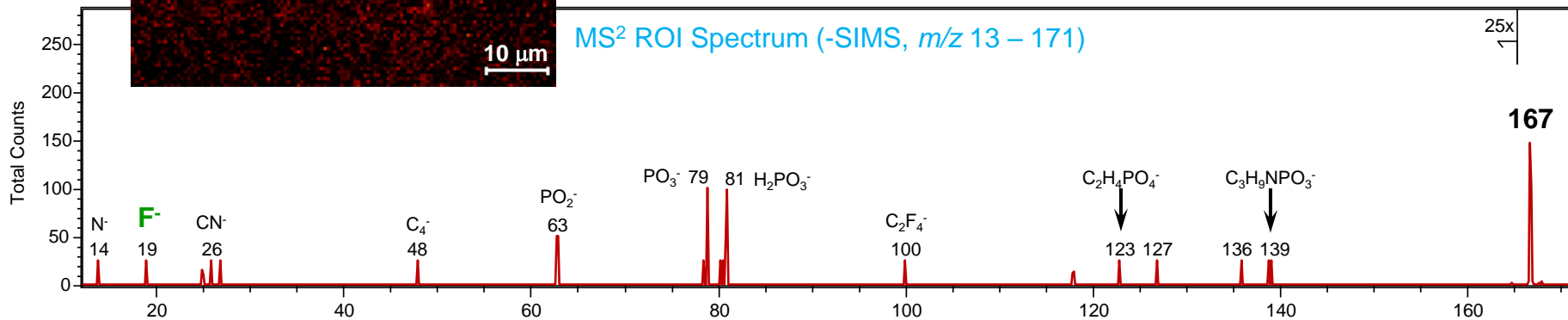


Tandem MS Region-of-Interest (ROI) Spectrum

The F⁻ in the MS² ROI spectrum indicates localization of a fragment of the ER-Tracker.

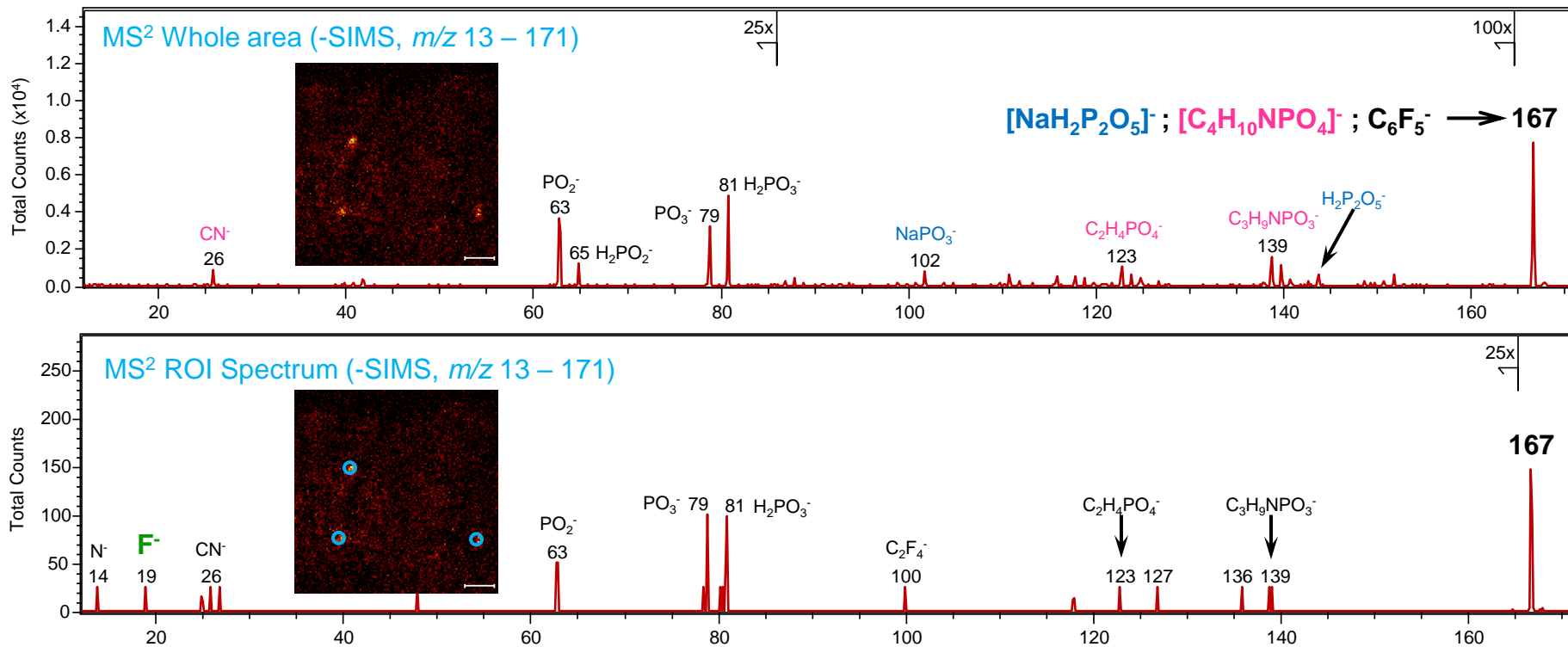


The relative counts of F⁻ ions are elevated in the ROI product ion (MS²) spectrum; *potentially, the F⁻ ions arise from a pentafluorophenyl anion.*



Tandem MS Spectra Comparison

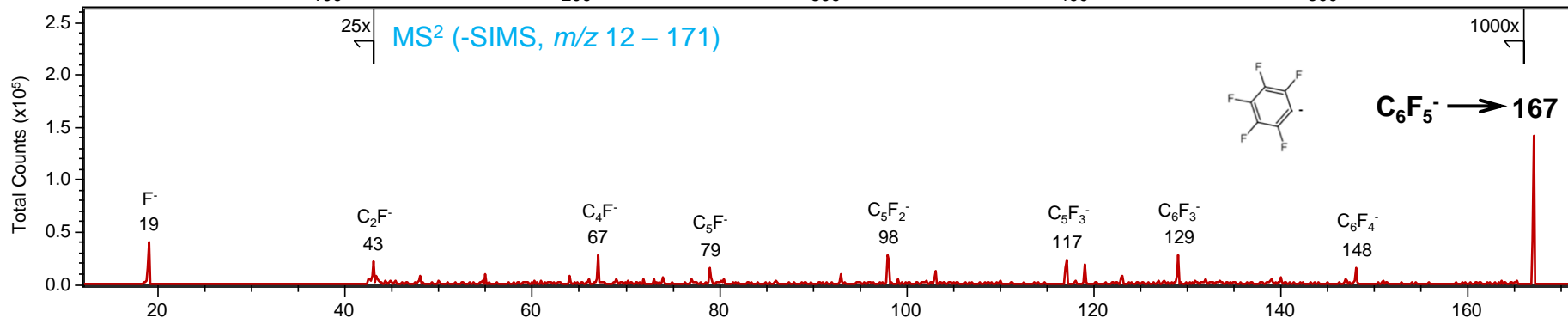
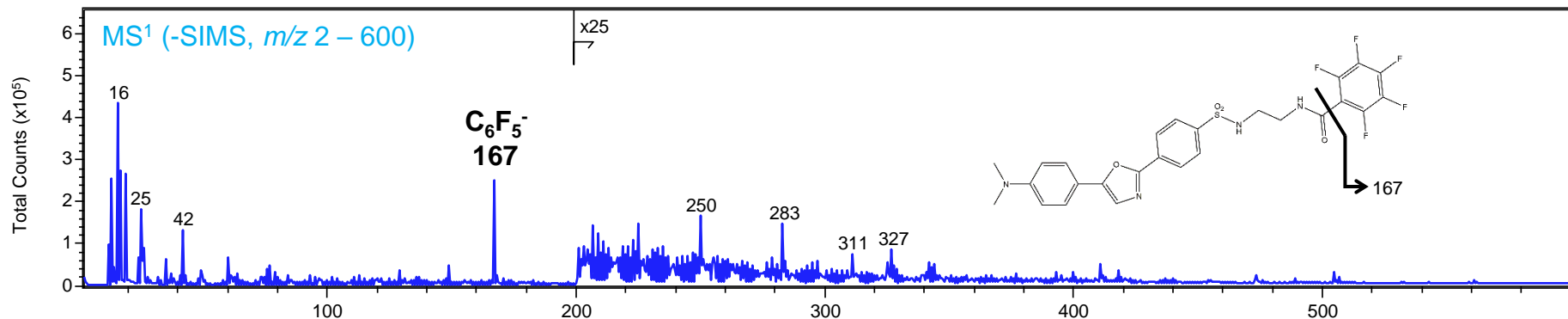
The F^- in the MS^2 ROI spectrum indicates localization of a fragment of the ER-Tracker.



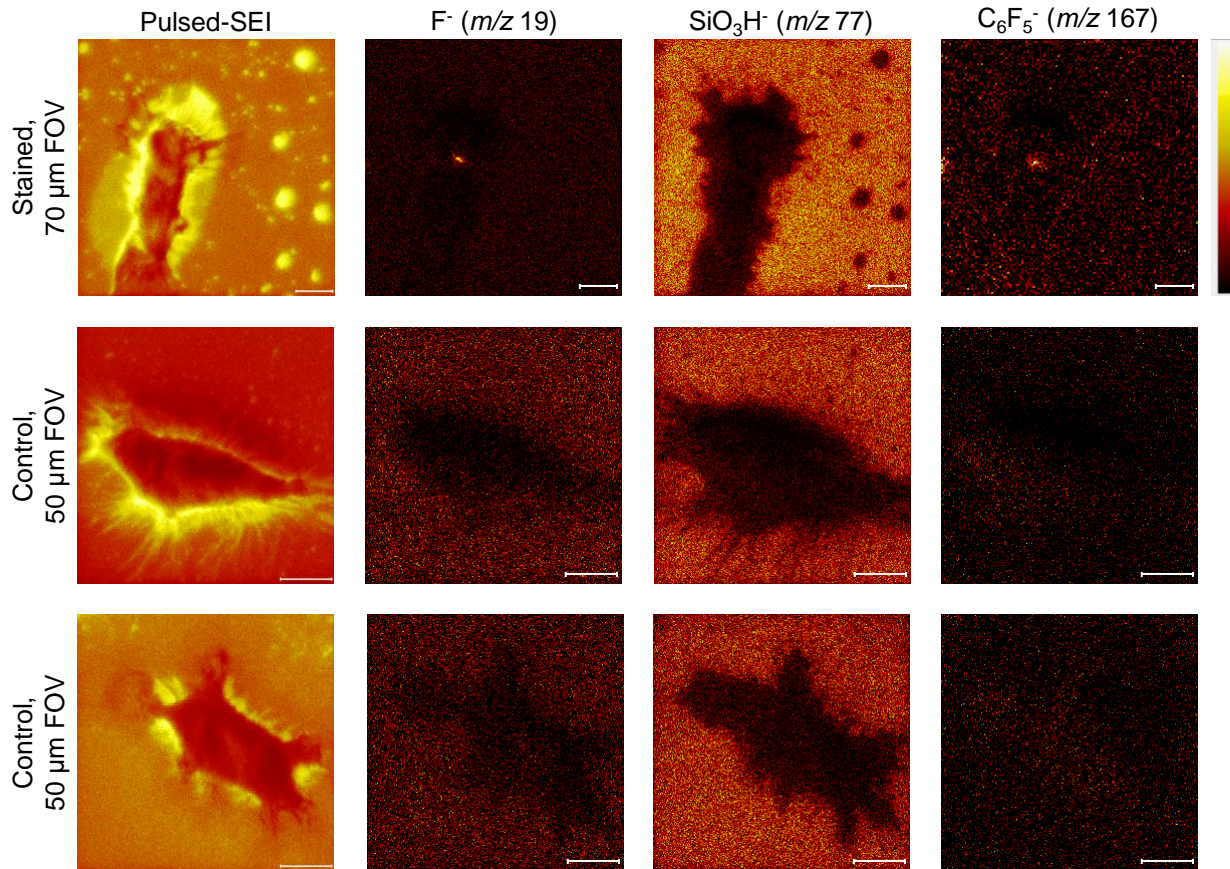
The relative counts of F^- ions are elevated in the ROI product ion (MS^2) spectrum; *potentially, the F^- ions arise from a pentafluorophenyl anion.*

Spectra of the ER-Tracker Reference

No molecular ion observed; MS² of -*m/z* 167 confirms a C₆F₅⁻ composition.



Are ER Features Observed in Unstained Control Cells?

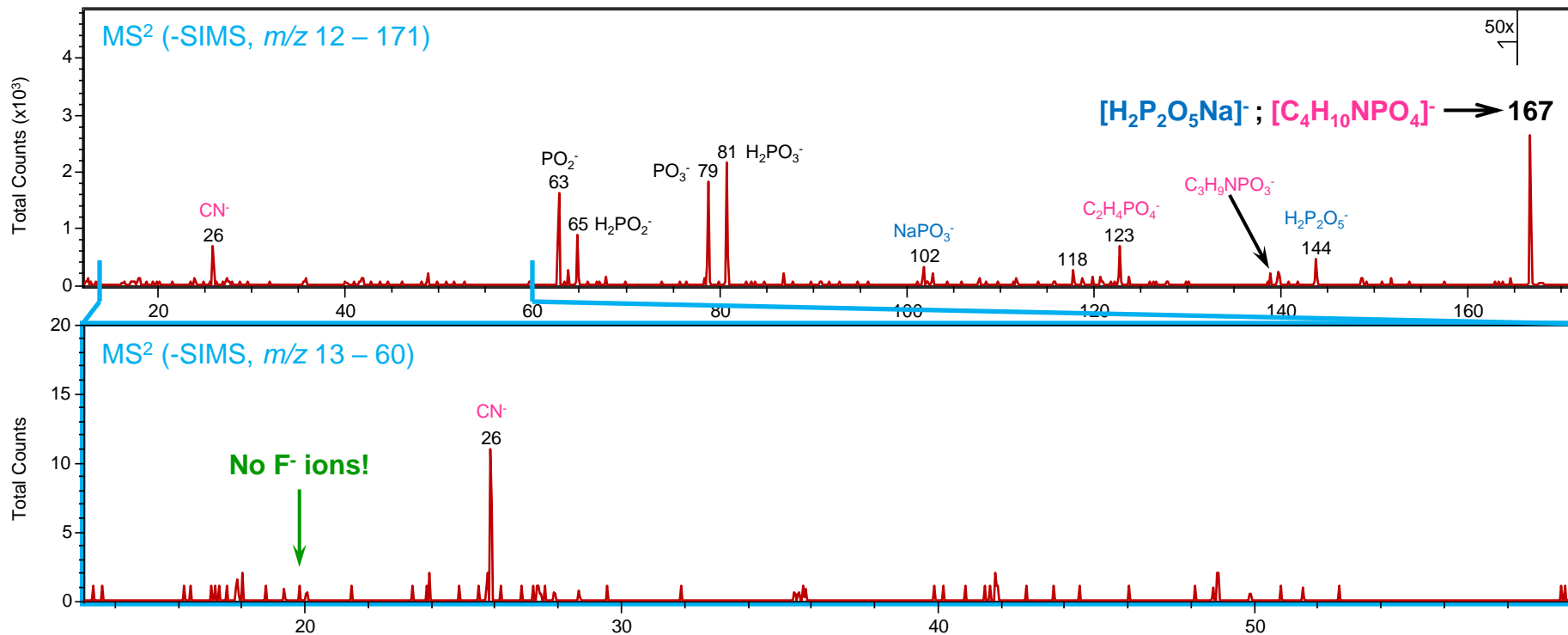


Possible ER tubules and ER-PM junctions are observed in stained cells as noted by localization and correspondence of the F^- and $C_6F_5^-$ ions.

The F^- and $C_6F_5^-$ chemical signatures are not observed in the control cells.

MS² of $-m/z$ 167 in Control Cell

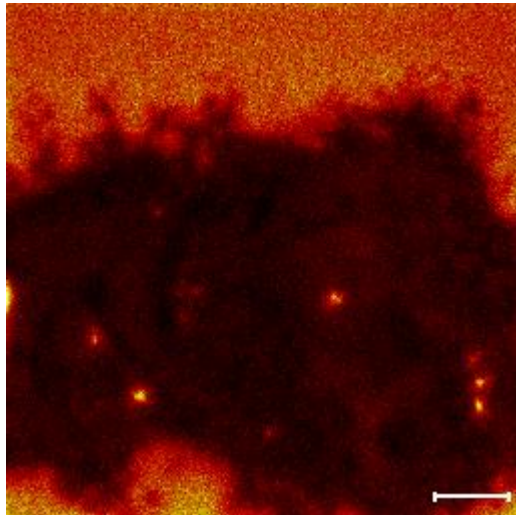
The product ion spectra show no evidence of the pentafluorophenyl anion.



ER Tubules Observed in Both Ion Polarities

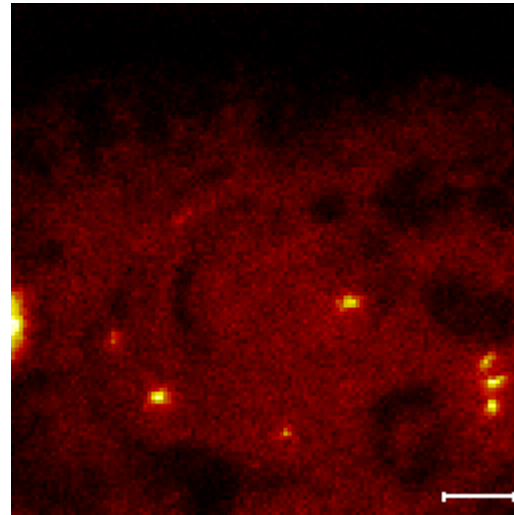
What is the source of the $C_5H_5^+$ ions?

-SIMS; 70 μm ; 8 min.



F^- (m/z 19)

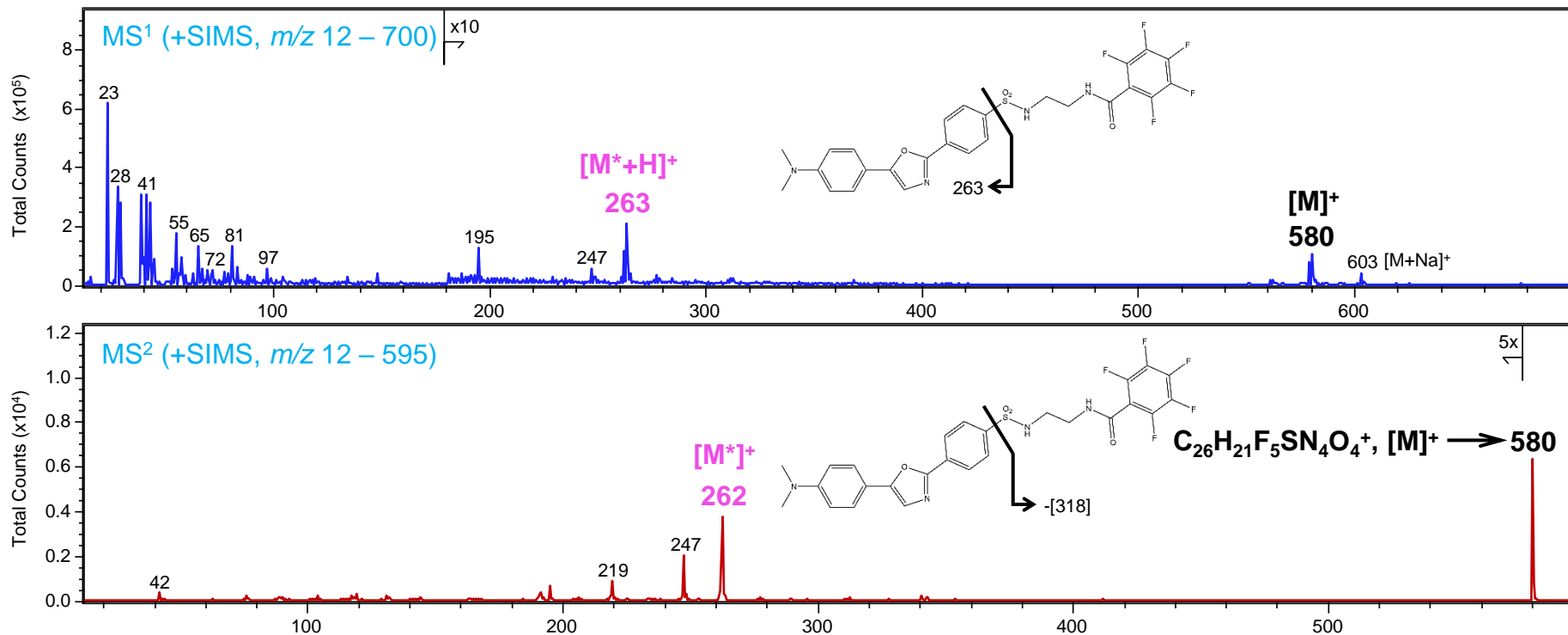
+SIMS; 70 μm ; 16 min.



$C_5H_5^+$ (m/z 65)
binned

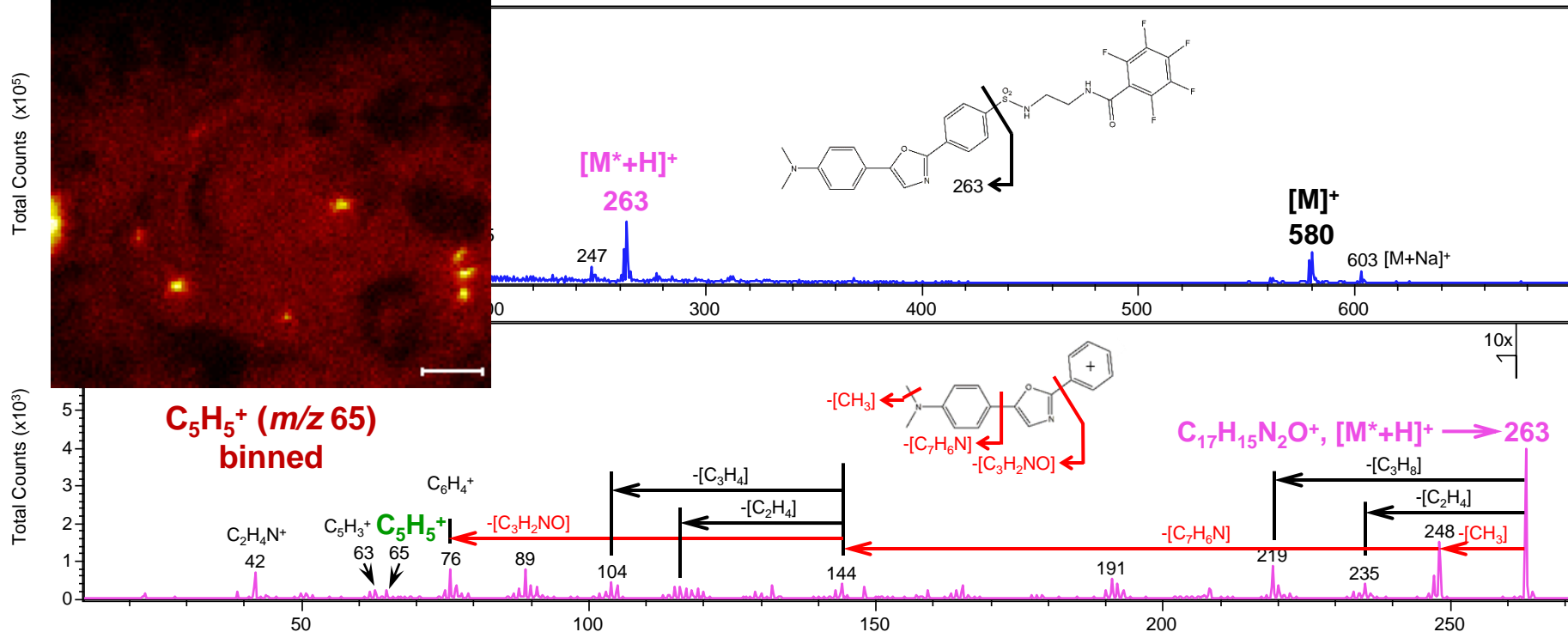
Spectra of the ER-Tracker Reference

MS¹ reveals the [M]⁺ and a fragment at *m/z* 263; MS² of [M]⁺ reveals a product ion at *m/z* 262.



Spectra of the ER-Tracker Reference

MS² of $[M^*+H]^+$ of the dapoxyl dye fragment and gives rise to $C_5H_5^+$ ions.



3D Tandem MS Imaging of Transfected/Stained HEK Cell

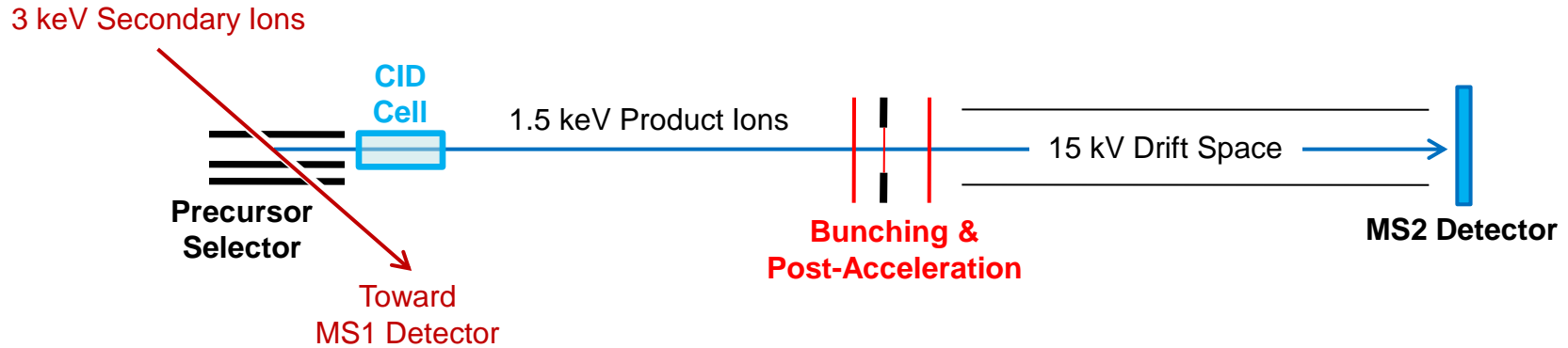
A 50 μm x 30 μm x 40 nm volume revealing the ER tubules and ER-PM junctions in one cell.



Summary & Outlook

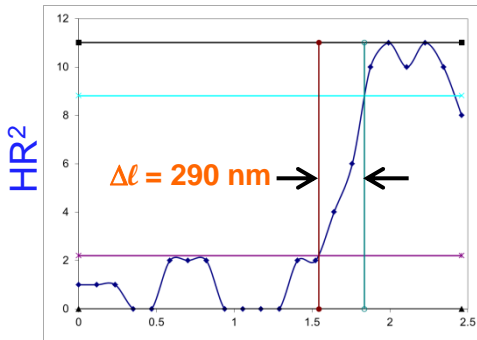
- Organelles must be “labeled” in order to achieve definitive localization of chemistry by MSI
- We employed a combination of staining (ER-Tracker) and tandem MS imaging to directly observe ER tubules and ER-PM junctions in transfected HEK cells
 - high resolving power TOF-SIMS (MS^1) imaging & tandem MS (MS^2) imaging at $\Delta\ell \approx 100$ nm
 - simultaneous collection of MS^1 and MS^2 data from each pixel enhances molecular ID and imaging
 - the same method can be applied to MS imaging of other organelles and cellular structures
- Full 3D MS^1 / MS^2 imaging of the ER in the entire cell volume is underway
- We further aim to identify organelle-specific lipids and differentiate them from e.g. PM lipids
- By extension, it should be possible to identify the location of drugs and metabolites in cells

Tandem MS (MS²) Spectrometer Schematic

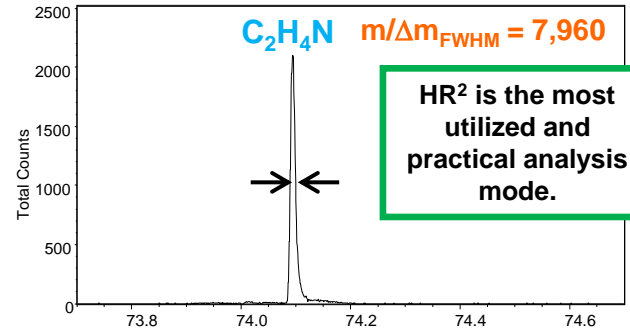
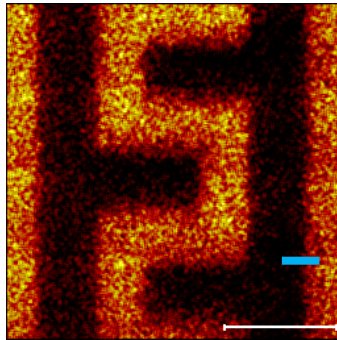


HR² Imaging versus Unbunched Imaging

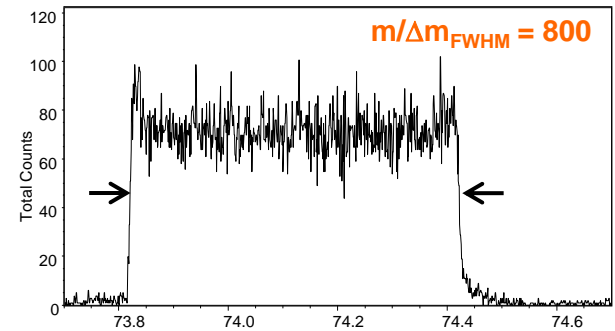
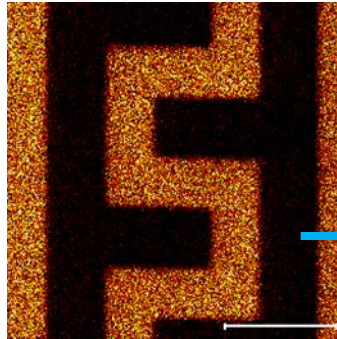
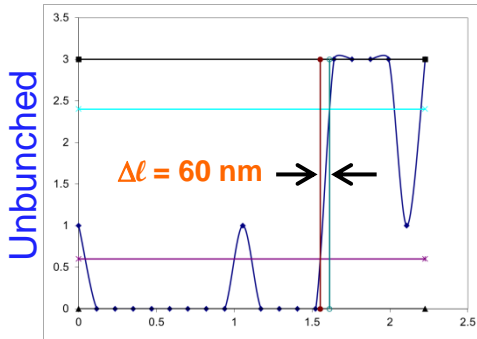
Chemical identification and high lateral resolution together.



Si⁺ (*m/z* 28); 30 μm FOV

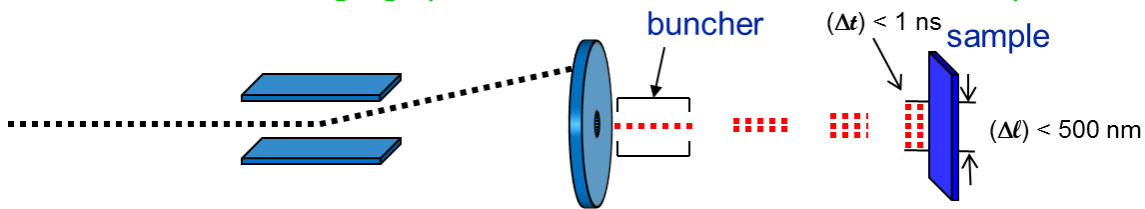


Ultimate lateral resolution but with little or no chemical information.

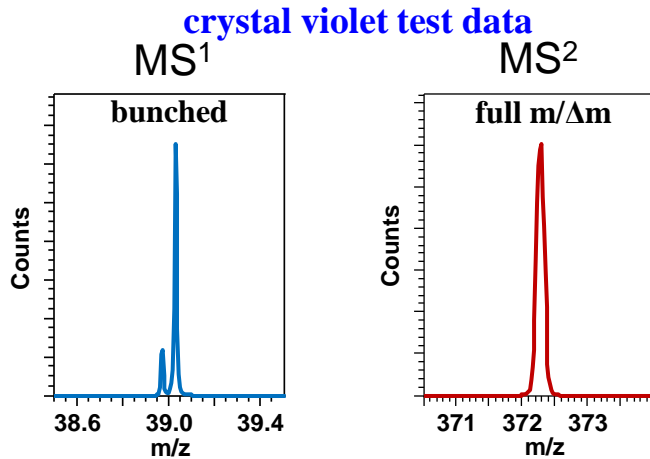


High Resolving Power & Full Tandem MS Resolution

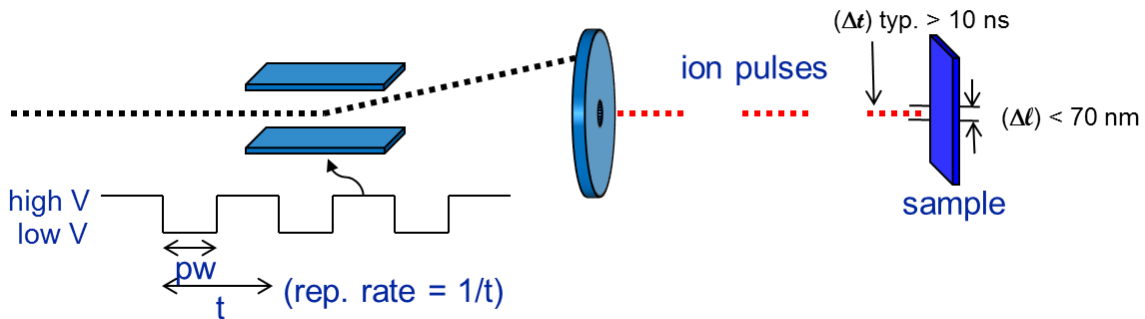
HR² Mode Imaging: (best mass resolution, $m/\Delta m > 10,000$)



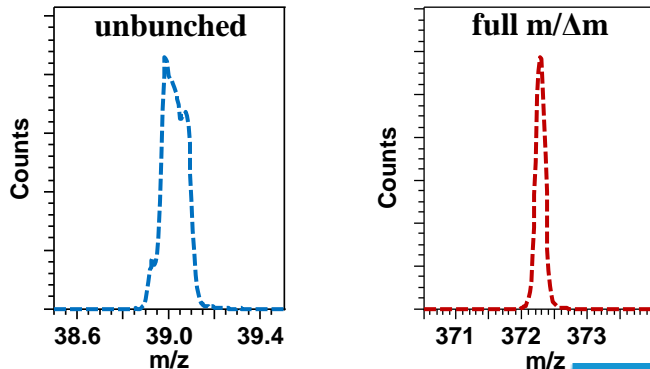
$\Delta \ell < 0.5 \mu\text{m}$



Unbunched Mode Imaging: (best lateral resolution, $\Delta \ell < 70 \text{ nm}$)



$\Delta \ell < 70 \text{ nm}$



What is the Composition at $-m/z$ 167 ?

Peak compositions are not confidently identified from the TOF-SIMS (MS^1) data alone.

