

FEI Tecnai F20 S/TEM: EDS system operation  
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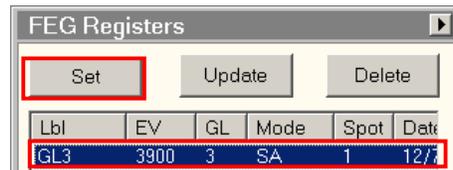
The user should already be familiar with operation of the instrument in STEM mode, use of the Microscope Control interface, and TIA.

## 1. Holder selection and plasma cleaning for EDS

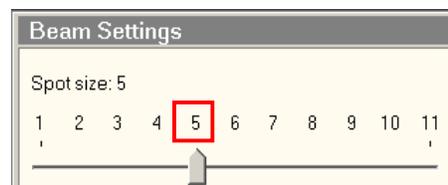
- 1.1. Either the double tilt or single tilt holders may be used for performing EDS. However, if the single tilt holder is used, it must be tilted a specific way to allow for efficient X-ray collection. It is also recommended that plasma cleaning of the specimen be performed (if permissible for the specimen) due to the high beam currents typically used for EDS.

## 2. Instrument settings for EDS

- 2.1. Select and apply the “GL3” FEG register.



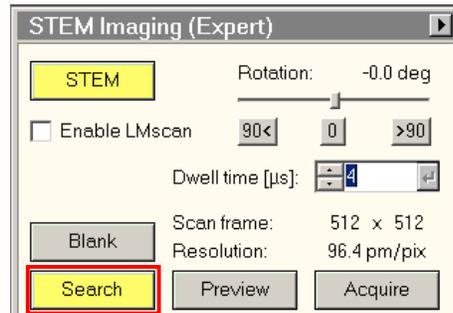
- 2.2. Verify that the objective aperture is retracted (this will prevent overload to the EDS detector).
- 2.3. If using the single tilt holder, set the alpha tilt to  $\alpha = +15^\circ$  and then reset eucentric height.
- 2.4. In Microscope Control, select the **STEM** tab and enter STEM mode; select spot size = 5 (after entering STEM mode); align the probe using the same procedure described for general STEM operation (use the #2 C2 aperture).



- 2.5. Set the camera length as desired (100 mm will produce a HAADF-STEM image) and center the Ronchigram inside the inner rim of the STEM detector.

### 3. Finding an area of interest

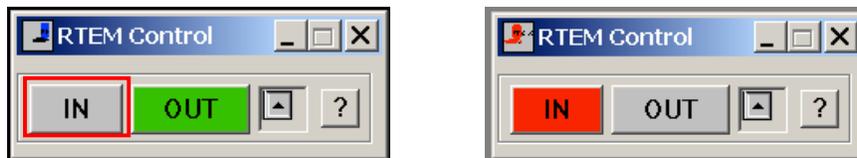
- 3.1. In Microscope Control, select the **EDS** tab; navigate to the “STEM Imaging” control panel and select “Search” to acquire a live image in TIA; adjust the magnification to change the field of view and the trackball to move around as needed.



- 3.2. After locating an area of interest, adjust the scan rotation, focus, and condenser stigmators, if needed.

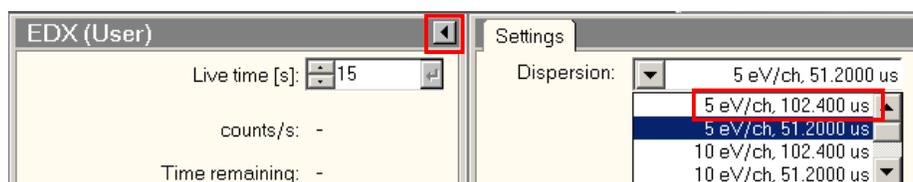
### 4. Inserting the EDS detector

- 4.1. In RTEM Control, select “IN” to insert the EDS detector; the “OUT” button will turn gray and the “IN” button will turn red.



### 5. Acquiring a survey spectrum; EDS detector settings

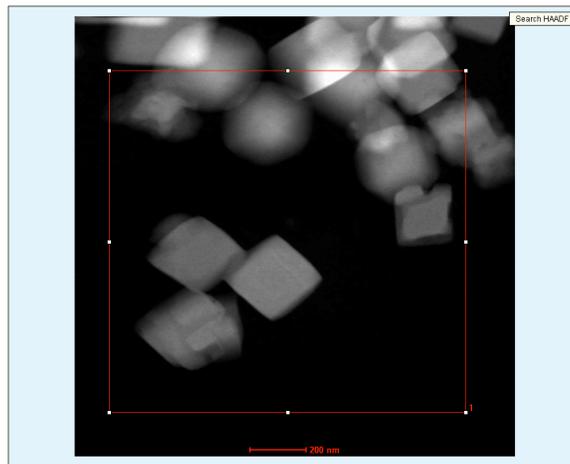
- 5.1. Make sure you are acquiring a live image using “Search”; otherwise, you cannot collect a survey spectrum.
- 5.2. In Microscope Control, select the **EDS** tab; navigate to the “EDX” control panel and select the flap out arrow  to expand the panel. Select “Dispersion” and select from the list of available combinations of eV/ch and process times; these values will also remain selected for any further analysis; when finished, select the flap out arrow  to collapse the panel.



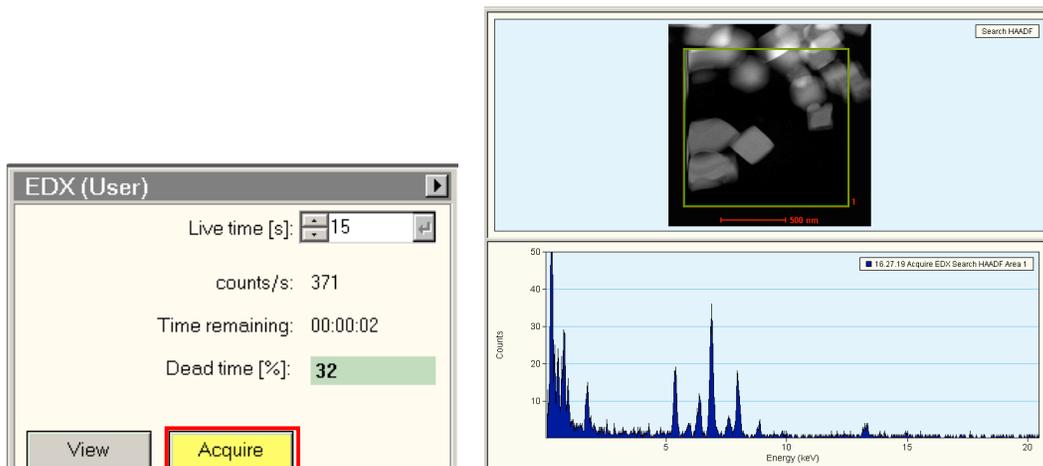
- 5.3. In the “EDX” control panel, enter in the desired value for “Live time” (the time the detector is actively collecting X-rays); a value of 15 – 60 s is usually sufficient for acquiring a survey spectrum



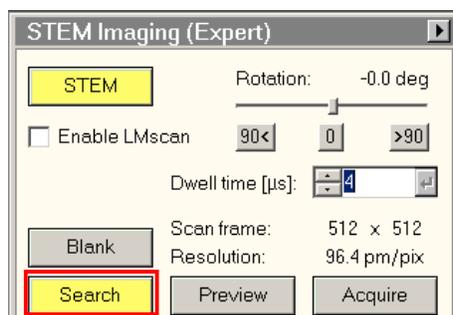
- 5.4. In TIA, select the image selection tool  and draw a box around the region in the image where a survey specimen is to be collected; the box may be repositioned and resized as needed.



- 5.5. In Microscope Control, navigate to the  tab; under the “EDX” control panel, select “Acquire” to collect a survey spectrum from the defined area (outline will turn green); the survey spectrum will appear below the STEM image in TIA and automatically update until complete.

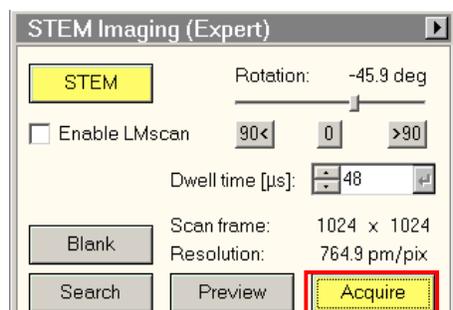


- 5.6. In TIA, select save  to save the display (STEM image and spectrum); then select close  to close the display.
- 5.7. The beam will freeze once the acquisition is complete. To restart the live image, go to Microscope Control and select the **EDS** tab; navigate to the “STEM Imaging” control panel and select “Search” to start acquiring a new live STEM image.

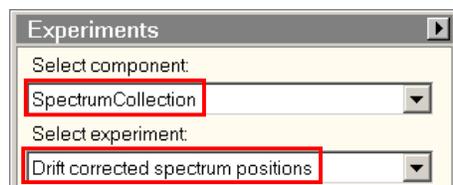


## 6. Point analysis

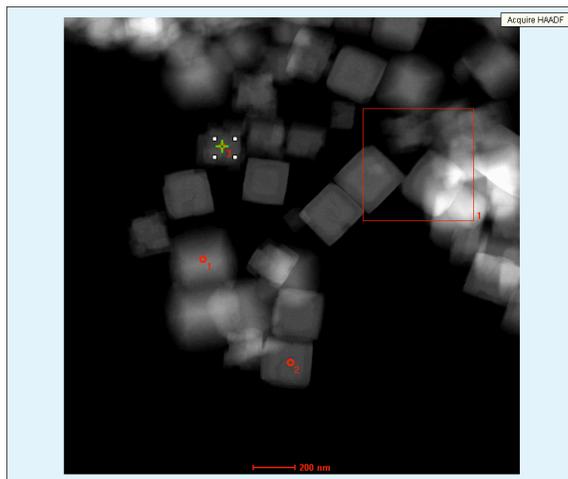
- 6.1. In Microscope Control, select the **EDS** tab; navigate to the “STEM Imaging” control panel and select “Acquire” to acquire a STEM image (you must use “Acquire” to collect the image or point analysis won’t work).



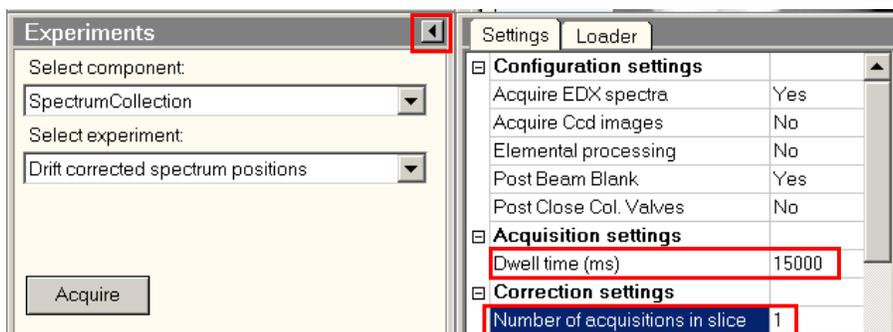
- 6.2. Navigate to the “Experiments” control panel; under “Select component”, select “SpectrumCollection” and under “Select experiment”, select “Drift corrected spectrum positions”.



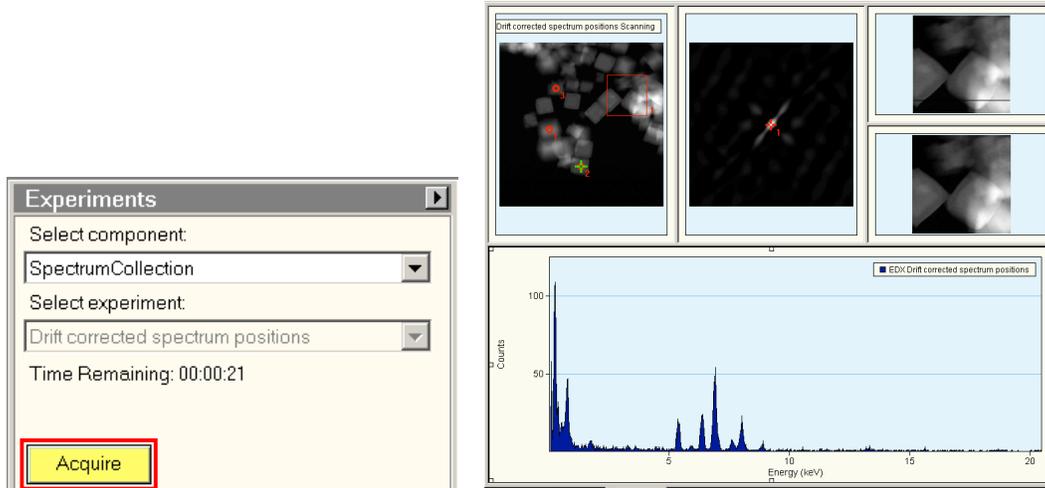
- 6.3. In TIA, a box and two beam position markers will appear on the image; position the markers where the spectra are to be collected; if you want to add another point, select the beam position marker tool  and click on the image to add another point (repeat for any additional points); then position the box around a distinct region in the image, not containing any of the markers (this will be used as the reference for drift correction).



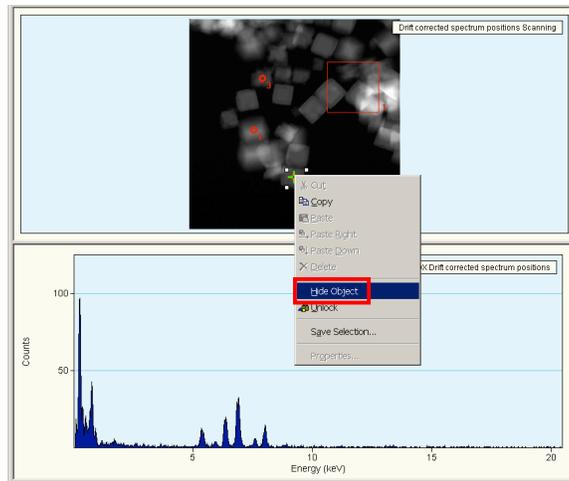
- 6.4. In Microscope Control, navigate to the “Experiments” control panel and select the flap out arrow  to expand the panel; under “Acquisition settings”, enter the dwell time to be used for each point; under “Correction settings”, make sure “Number of acquisitions in slice” is set to 1; when finished, select the flap out arrow  to collapse the panel.



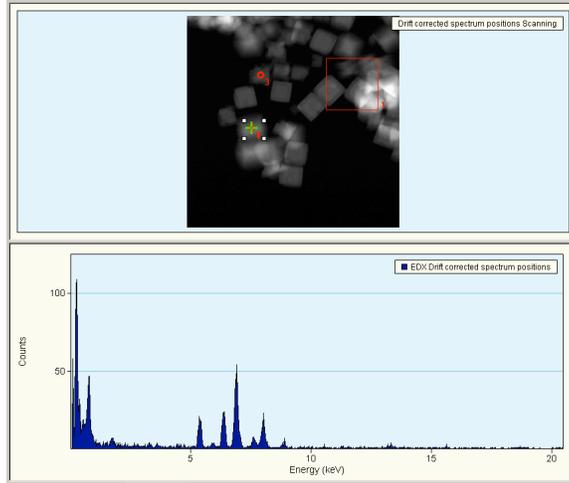
- 6.5. In the “Experiments” control panel, select “Acquire” to start collecting the spectra at the defined positions; after each spectrum is collected, it will be displayed below the STEM image in TIA (the final spectrum shown will be the one collected from the last point).



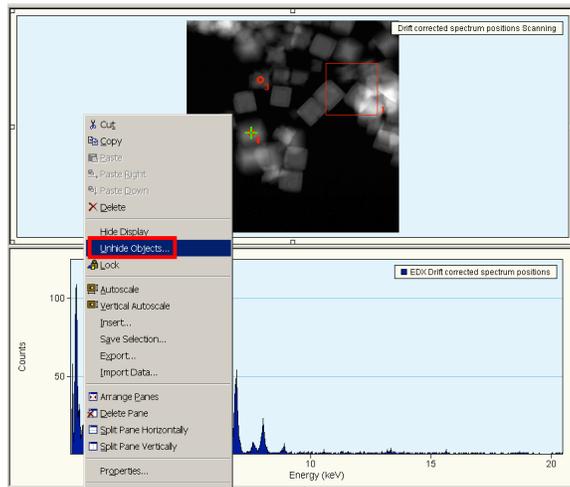
- 6.6. Once acquisition is complete and you want to view the individual spectra collected at the different points, you must first hide the beam position markers. To do this, simply right click on a beam position marker and select “Hide Object” and repeat for all remaining beam markers.



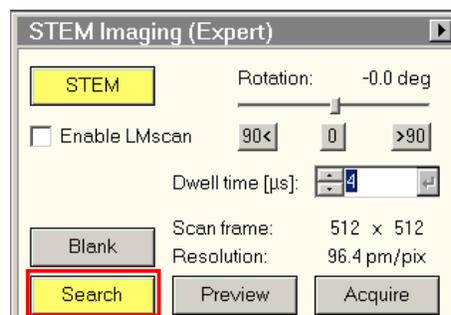
- 6.7. In TIA, select the image position marker tool  and click anywhere on the image; then hold down the “Alt” key and click and drag it onto the spectrum below the image (the actual dark blue portion, not the background); the marker will now snap to a beam position marker and the spectrum for that point will appear. You can now simply click and drag the image position marker and it will snap to all the beam position markers and show the spectrum for each position.



- 6.8. To reveal the beam position markers again, right click on the light blue background in the panel containing the image and select “Unhide Objects”.

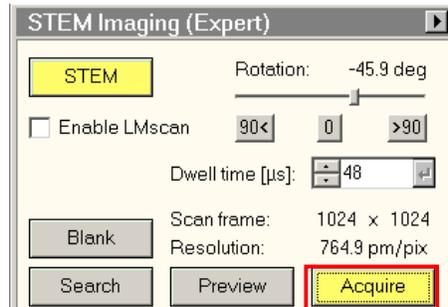


- 6.9. To restart the live image after acquisition, navigate to the “STEM Imaging” control panel and select “Search” to start acquiring a new live STEM image.

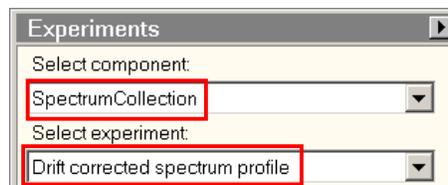


## 7. Spectrum profiles (line scans)

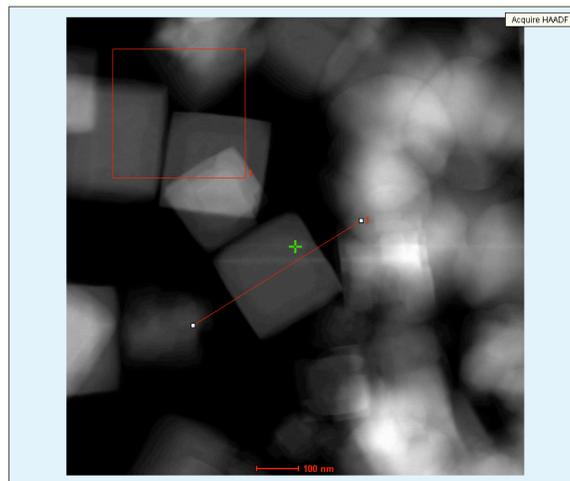
- 7.1. In Microscope Control, select the **EDS** tab; navigate to the “STEM Imaging” control panel and select “Acquire” to acquire a STEM image (you must use acquire to collect the image or spectrum profiles cannot be collected).



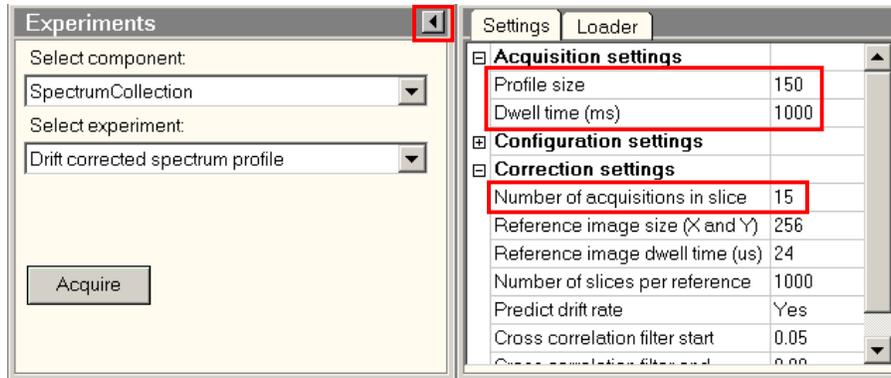
- 7.2. Navigate to the “Experiments” control panel; under “Select component”, select “SpectrumCollection” and under “Select experiment”, select “Drift corrected spectrum profile”.



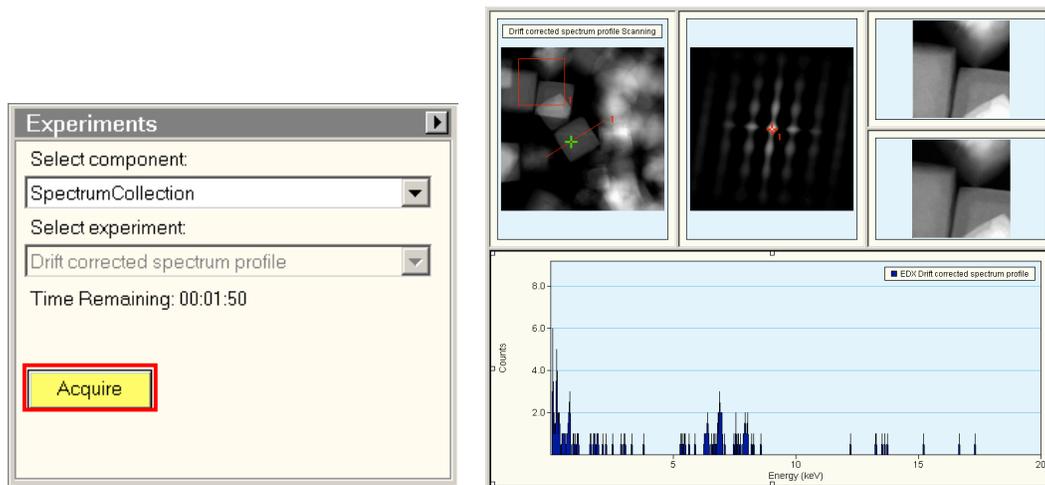
- 7.3. In TIA, a box and line marker will appear on the image; position/resize the line to where you want to collect the spectrum profile; then position the box around a distinct region in the image, not containing line marker (this will be used as the reference for drift correction).



- 7.4. In Microscope Control, navigate to the “Experiments” control panel and select the flap out arrow  to expand the panel; under “Acquisition settings”, enter the desired profile size (number of points in the spectrum) and the dwell time (how long the probe collects a spectrum at each point); under “Correction settings”, set “Number of acquisitions in slice” to 10% of the number of points in the profile (e.g. if there 150 points in the profile, set Number of acquisitions in slice = 15); when finished, select the flap out arrow  to collapse the panel.

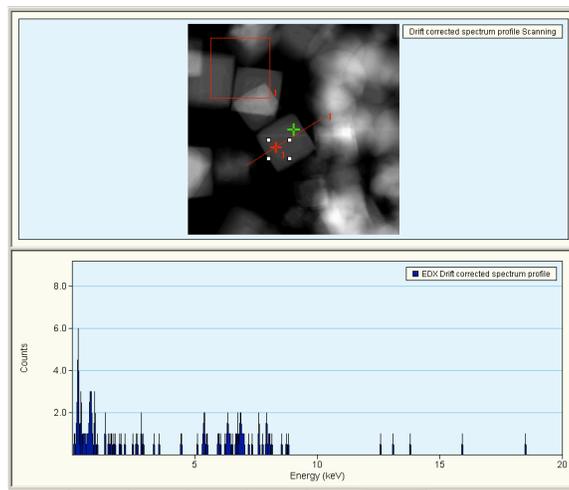


- 7.5. In the “Experiments” control panel, select “Acquire” to start collecting the spectrum profile; after each spectrum at each point is collected, it will be displayed below the STEM image in TIA (the final spectrum shown will be the one collected from the last point in the profile).



- 7.6. Once acquisition is complete you can view the spectrum collected at any point on the profile; select the image position marker tool  and click anywhere on the image; then hold down the “Alt” key and click and drag it onto the spectrum below the image (the actual dark blue portion, not the background); the marker will now snap to the line marker and the spectrum for that point will appear. You can now simply click and drag the image position marker along the line and it will snap to all the beam position markers and show the spectrum for each position.

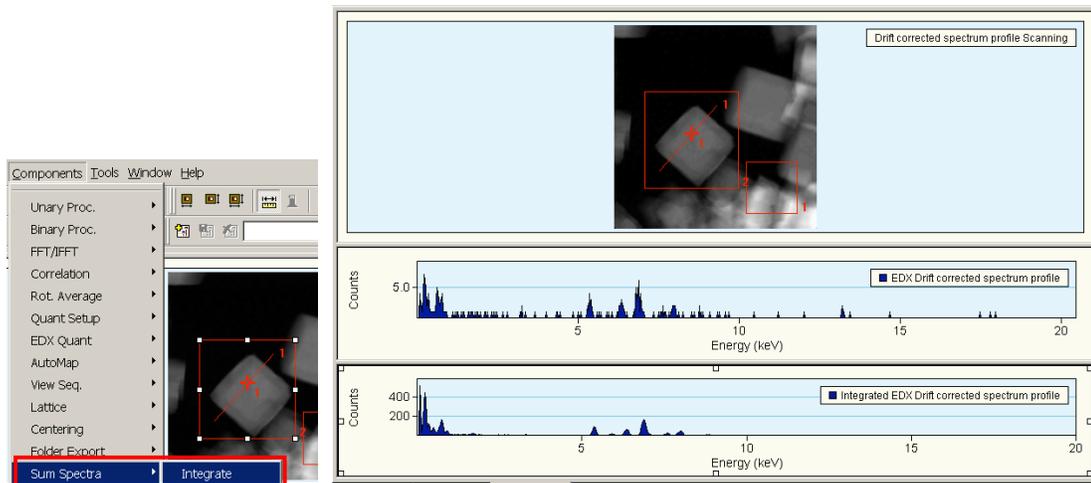
**Do not delete the pane containing the spectrum corresponding to the image position marker;** otherwise, the information needed to generate the profiles will be lost.



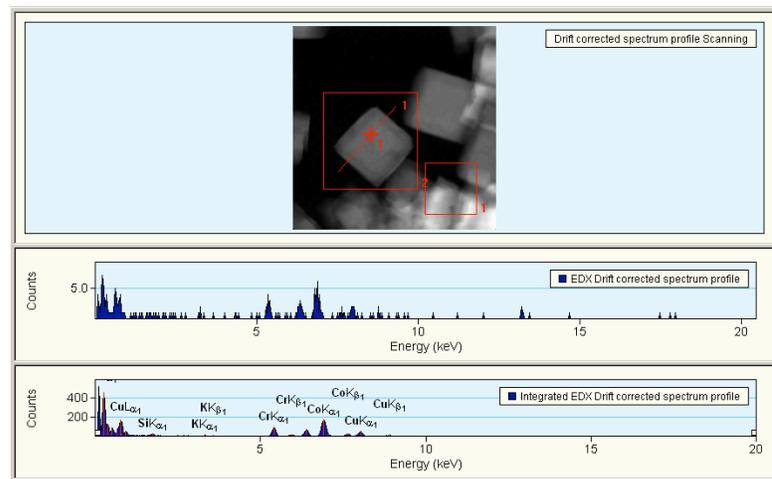
- 7.7. To generate the profiles for the elements of interest, the data must first be saved  and TIA must then be expanded to analysis mode; select  from the lower information panel; the button will change to  and TIA will be in analysis mode.



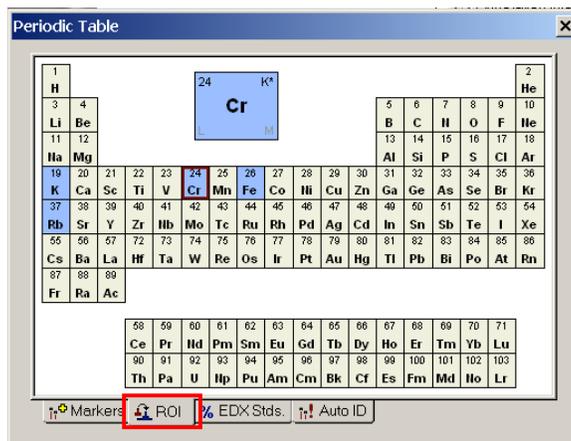
7.8. It will be difficult to identify individual peaks in a spectrum from a single point, so the sum spectrum from all the points should be used for this. To generate the sum spectrum, select the image selection tool  and draw a box around the entire line on the STEM image; then select “Components” from the pull-down menu, then “Sum Spectra” and then “Integrate”; the sum spectrum will be generated in a new pane below the pane containing the spectrum from a single point.



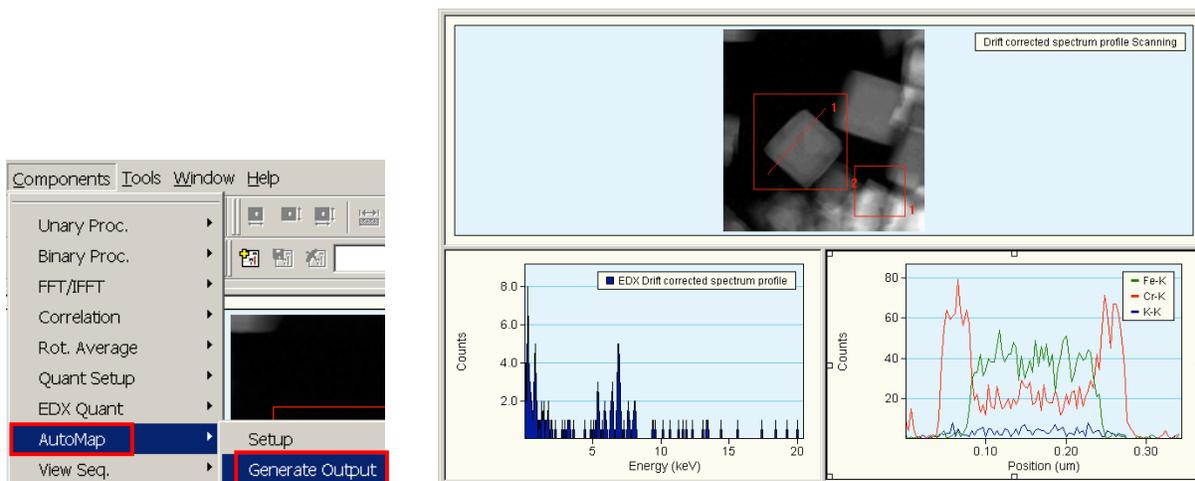
7.9. Select the sum spectrum and then select auto ID  to automatically identify the different peaks (this is usually sufficiently accurate) in the sum spectrum; you can also select KLM labels button  to further characterize the peaks by transition.



- 7.10. Select the periodic table button ; a periodic table dialogue box will pop up. Select the “ROI” tab  to choose which elements will or will not be used to generate the profiles. Each time you select an element, you can select which X-ray peak(s) to use for mapping by selecting from the **K L M** buttons



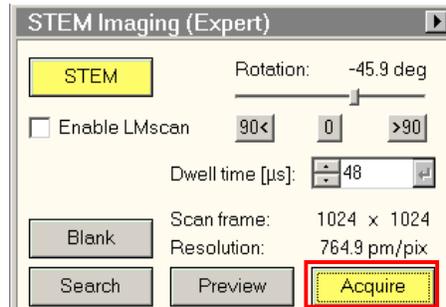
- 7.11. Once the ROIs for the different elements of interest have been selected (Cr K, Fe K, etc.), select “Components” from the pull down menu, then “AutoMap”, then “Generate Output”; a new pane will be generated containing the profiles for each of the selected X-ray peaks. After doing this, it is a good idea to save the data 



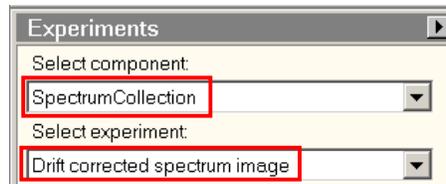
- 7.12. When finished, select  to switch from analysis mode back to acquisition mode .

## 8. Spectrum imaging (mapping)

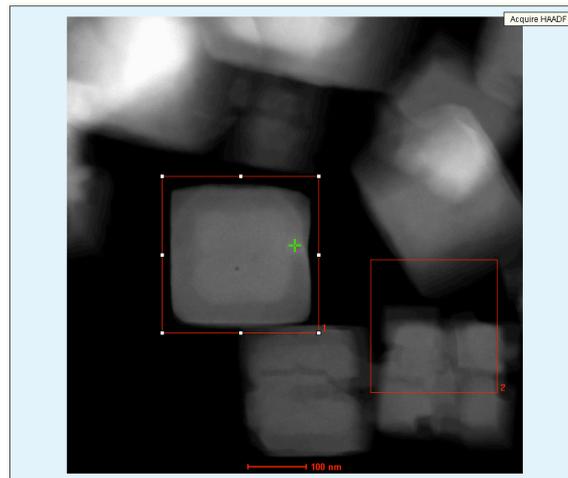
- 8.1. In Microscope Control, select the **EDS** tab; navigate to the “STEM Imaging” control panel and select “Acquire” to acquire a STEM image (you must use acquire to collect the image or spectrum imaging cannot be performed).



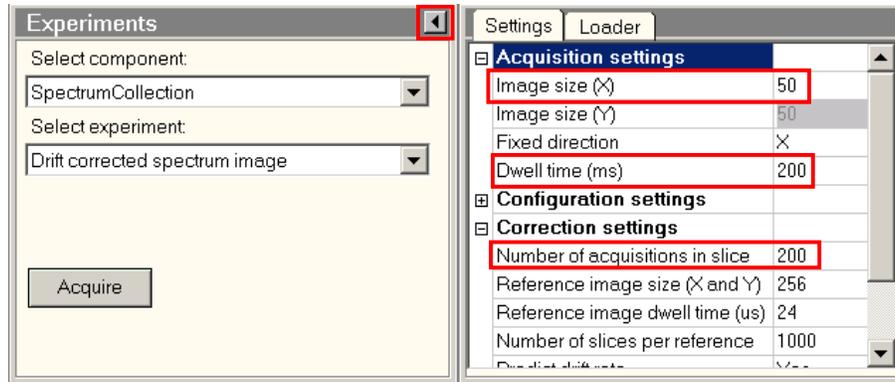
- 8.2. Navigate to the “Experiments” control panel; under “Select component”, select “SpectrumCollection” and under “Select experiment”, select “Drift corrected spectrum image”.



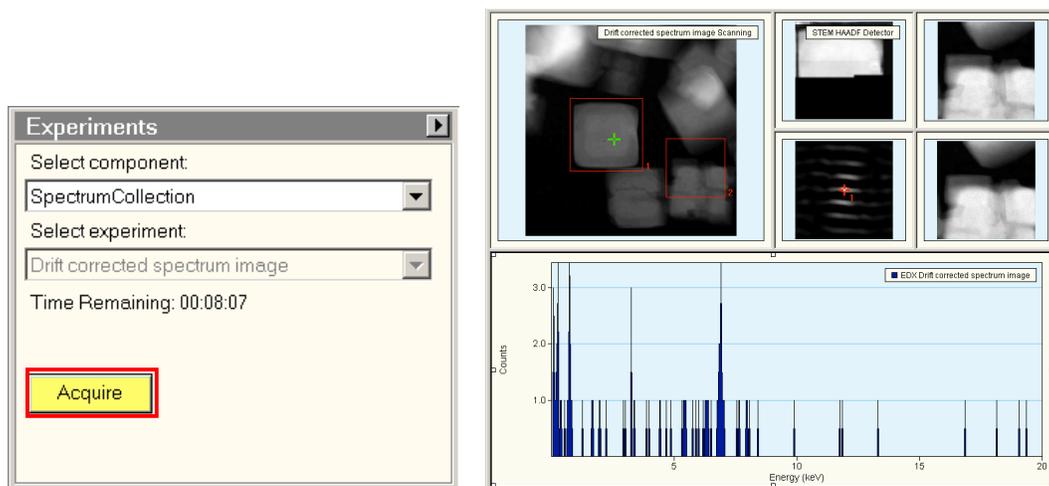
- 8.3. In TIA, 2 boxes will appear on the image; box 1 is the area where spectrum imaging will be performed and box 2 is the area that will be used as the reference for drift correction (again, box 2 should not overlap with box 1 and should ideally contain distinct features); resize and reposition the boxes as necessary.



- 8.4. In Microscope Control, navigate to the “Experiments” control panel and select the flap out arrow  to expand the panel; under “Acquisition settings”, enter the desired image size (how wide the imaged area is in terms of points) and the dwell time (how long the probe collects a spectrum at each point); under “Correction settings”, set “Number of acquisitions in slice” to 10% of total number of points in the image (e.g. if Image size = 50, there will be 2500 total points, so set Number of acquisitions in slice = 250); when finished, select the flap out arrow  to collapse the panel.

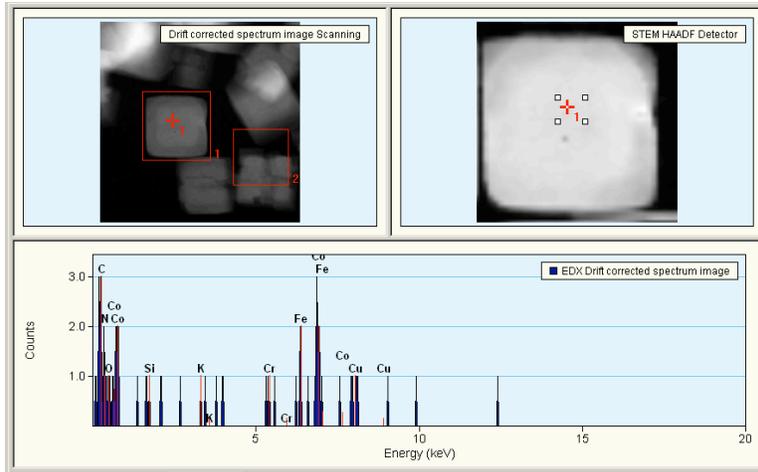


- 8.5. In the “Experiments” control panel, select “Acquire” to start collecting the spectrum image; after each spectrum at each point is collected, it will be displayed below the STEM image in TIA (the final spectrum shown will be the one collected from the last point in the defining region).



- 8.6. Once acquisition is complete you can view the spectrum collected at any point in the area used for spectrum imaging; simply click and drag on the image position marker tool  in the area (you do not need to add a new marker) and the spectrum collected at that position will be shown below.

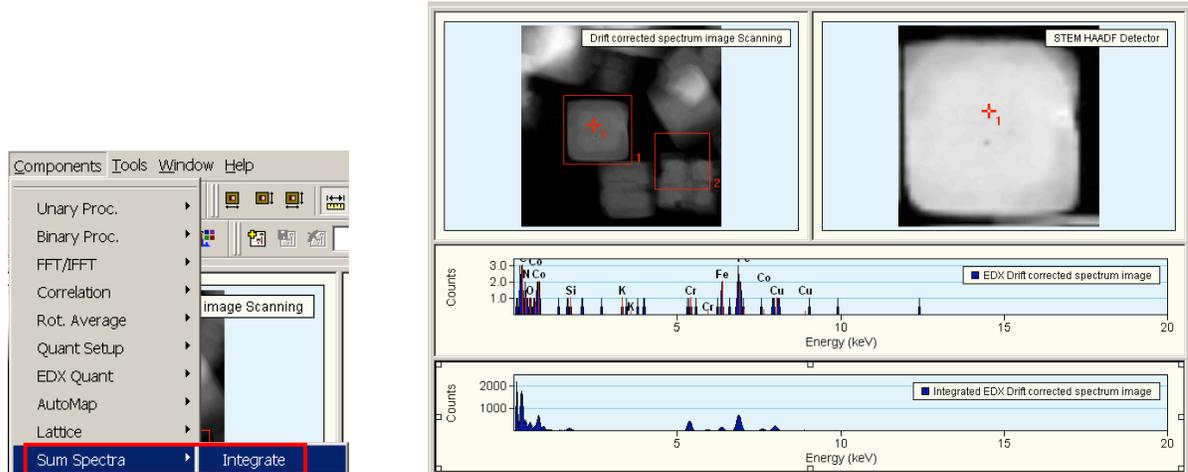
**Do not delete the pane containing the spectrum corresponding to the image position marker;** otherwise, the information needed to generate the maps will be lost.



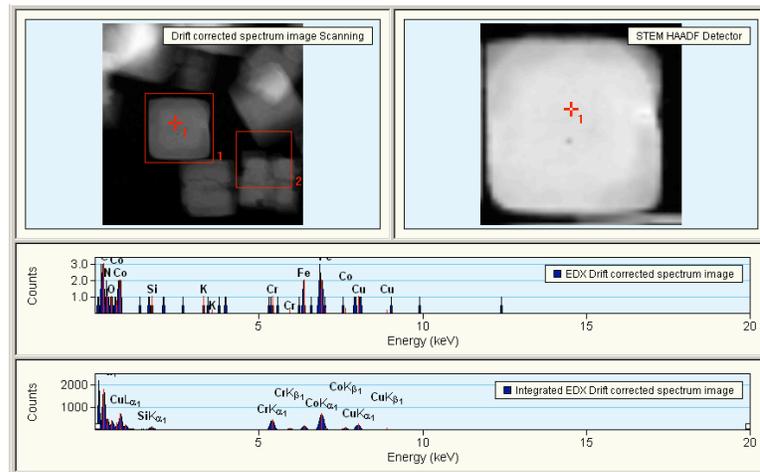
- 8.7. To generate the spectrum images for the elements of interest, the data must first be saved  and TIA must then be expanded to analysis mode; select  from the lower information panel; the button will change to  and TIA will be in analysis mode.



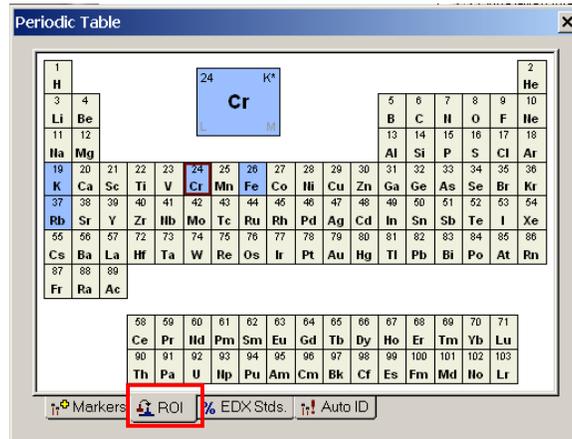
8.8. It will be difficult to identify individual peaks in a spectrum from a single point, so the sum spectrum from all the points should be used for this. To generate the sum spectrum, select the red square used to define the area for spectrum imaging; then select “Components” from the pull-down menu, then “Sum Spectra” and then “Integrate”; the sum spectrum will be generated in a new pane below the pane containing the spectrum from a single point.



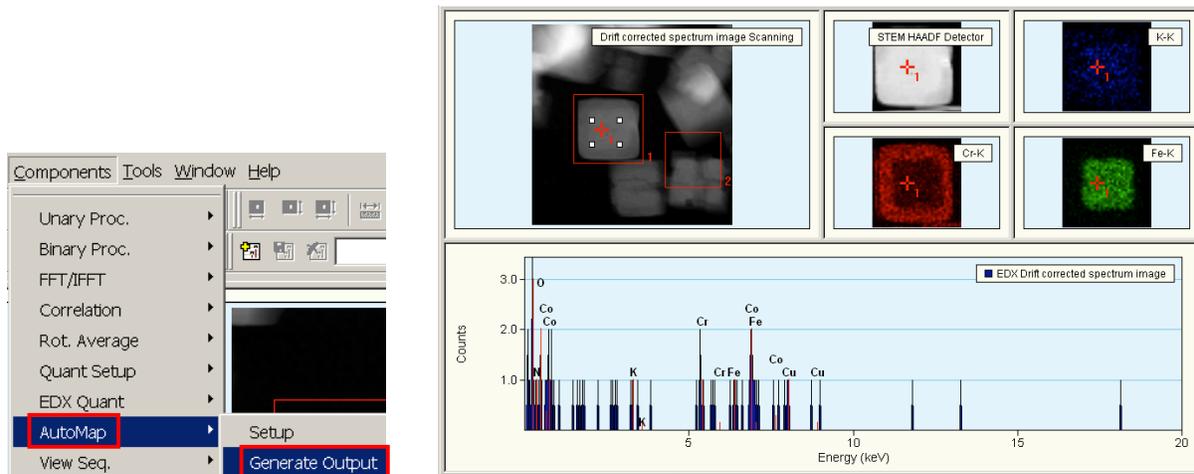
8.9. Select the sum spectrum and then select auto ID  to automatically identify the different peaks (this is usually sufficiently accurate) in the sum spectrum; you can also select KLM labels button  to further characterize the peaks by transition.



- 8.10. Select the periodic table button ; a periodic table dialogue box will pop up. Select the “ROI” tab  to choose which elements will or will not be used to generate the spectrum images. Each time you select an element, you can select which X-ray peak(s) to use for mapping by selecting from the **K L M** buttons.



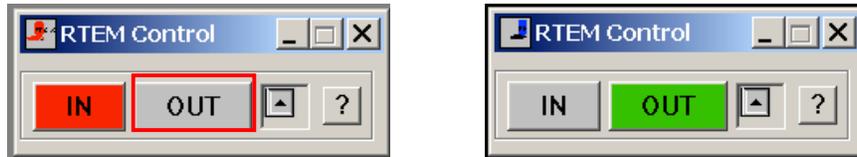
- 8.11. Once the ROIs for the different elements of interest have been selected (Cr K, Fe K, etc.), select “Components” from the pull down menu, then “AutoMap”, then “Generate Output”; for each of the selected X-ray peaks, a new pane will be generated showing the X-ray map. After doing this, it is a good idea to save the data .



- 8.12. When finished, select  to switch from analysis mode back to acquisition mode .

## 9. Finishing

- 9.1. In RTEM Control, select “OUT” to retract the EDS detector; the “OUT” button will turn green and the “IN” button will turn gray. Do not leave the EDS detector inserted unless it is actively in use.



- 9.2. In Microscope Control, select the STEM tab and exit STEM mode.