Ultra-55

Zeiss Ultra-55 Field Emission Scanning Electron Microscope (FESEM)

CNS Harvard

- Prepare your samples in a puck and mark the tape by cutting one corner.
- Make a map of your samples
- Give a description of which elements you expect to find in each sample.



- 1. Open the web browser
- 2. Click on CNS CLEAN
- 3. Select Ultra55 from Room B15 i
- 4. User type \rightarrow User

- 5. Select "Have Reservation"
- 6. Choose your name
- 7. Enter password (cns2tafti)
- 8. Login



Before starting EDX, make sure the detector is in the correct position and before removing samples, make sure the detector is in the safe position

Place the puck in the holder and tighten the screw





Magnification knob

XY-stick

Z-stick

+/- increase/decrease scanning rates. Typically, you need slower scans to clarify backscattered images.







From top menu, select Stage\Navigation Adjust a safe Z-height

IR camera (Chamber Scope)



Sample Holder Carousel 9x 6.5mm At the end of the experiment, you need to insert the information here, enter your PW, and log out by re-entering your password

- 1. Double click on "smart sem"
- 2. Input ID and PW (ftafti, cns2tafti)
- 3. Fix samples on the puck
- 4. Click on "Change" to vents the chamber

- 5. Press "OK" to start pumping
- 6. Wait until "vac" is green
- Press "EHT" and select "EHT on" to start high tension



- 1. On top bar, select "stage" \rightarrow "Navigation"
- 2. Double click on the stage number
- 3. Use the Z-stick to bring the stage up while monitoring with the IR camera
- 4. Be careful not to exceed the safe Z limit

- 5. Double click on "WD" and put it to 8.5, then hit enter
- 6. Use the Z-stick to bring the stage to focus
- 7. Double click on "Mag" to adjust



To save image, File \rightarrow save image \rightarrow choose directory and save



To start EDX, put the detector in the correct position \rightarrow **turn off ccd** (make sure the checkbox is off, no need to close the window) \rightarrow double click on "EDX-Genesis"



Image tab \rightarrow collect e image \rightarrow select region \rightarrow red icon cleans the existing spectrum and stopwatch collects new spectrum



Peak ID \rightarrow click on the drop-down icon and select element, you may need to delete O Select area for or Al or C for better quantitative results, Pt scanning and Pd coating always appear CODAX Spectrum Image Maps/Line Particles File Edit View Collect Display Process MultiField Setup Window Help 🛩 🖆 🖬 🏐 T 🔎 🖤 🗋 🖽 🔠 🔙 🐯 😵 🏥 😐 + 👾 🖬 \downarrow 🔖 🗸 Preset 💌 Amp Time 3.20 uS 💌 👥 🕺 Analyzer Det 1 Scan selected area Untitled kV: 15.0 Mag: 1000 Reso:1024x800 Collect e Strips:1 Conti 4095 16 6E1 1 0 bel Det Smin Smax Reads Matrix: 1024×800 Signal: SE1 Data: ADC Au CutePDF Writer Prin Build Max Spc + = 🏢 Build Spc Range: 0-4095 100.00% Area Frac: **Color Area** 👌 🐹 Untitled: <>> × △ ≚ ④ ▲↓ Q↓ Counts

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- To save spectrum data → Go to spectrum tab → File → Import SCP from → Image/Maps
- Then, Save \rightarrow save as case.csv



After each analysis and saving data, unfreeze the microscope by pressing on "EXT-XY"



Color of each element can be changed after the mapping is finished and individual figures can be saved with a new color.

Panels can be re-arranged

Counts

_4.2k

3.6k

-3.0k

2.4k _____1.8k - 1.2k B



Before logging out, make sure to remove your sample and pump down the chamber. To log out, input the information in the system Log, then input your password and click logout.



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Waiting for clean.rc.fas.harvard.edu...



