Endstation Operation and Data Collection Procedures

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All commands to be typed in the terminal are in italics. **Enter a command without arguments to see usage instructions.**

1 Starting Endstation Software

1.1 Starting SPEC Software to collect data.

- 1. Open a terminal and login to the data acquisition computer by double-clicking runRIXS_ES on the desktop.
 - Enter your **group password**. The beamline staff should have provided login credentials for your group.
- 2. Start spec software on the RIXS ES by typing runRIXS.
 - Report any errors during starting to the beamline staff.
 - The procedure for recovering in case of an accidental terminal closure or computer issue is identical to above.

1.2 Starting SPEC dataGUI to visualize live data.

- 1. Start dataGUI by double-clicking runRIXS_GUI icon.
 - It will take a few seconds to load.

1.3 Starting cameraGUI to for sample alignment.

1. Start the camera software by double-clicking the runRIXS_Cameras icon.

1.4 Starting Jupyter Notebook for Data Analysis.

- 1. Start the analysis software by double-clicking the **runREIXS_Analysis** icon.
 - This will launch a terminal to start juptyer notebook and then launch a web browser.
 - The analysis notebook can only be run on one computer at a time.
 - Documentation can be found here: https://pypi.org/project/reixs/

1.5 Starting EDM screens for additional information.

- 1. Start the EPICS EDM screen by double-clicking the REIXS_Users icon.
 - This will launch window to view some beamline variables within EPICS
 - Launch additional relevant windows or strip-tool using the appropriate buttons.

2 Setup REIXS Beamline

• These steps only need to completed if you are starting for the first time, or if you are changing elemental edges.

2.1 Setup EPU

- 1. Check the current settings of the EPU using statEPU or observe the parameters on the dataGUI.
- 2. Select the EPU polarization using *setEPU polar*. Examples below:
 - Typically one uses horizontal polarization for almost all experiments. Vertical polarization is used to for spectrometer calibration or for specific experiments.
 - *setEPU polar lh* linear horizontal (default)
 - setEPU polar lvn linear vertical negative (use for calibration)
 - setEPU polar lvp linear vertical positive
 - setEPU polar li linear inclined
 - setEPU polar cl circular left
 - setEPU polar cr circular right
- 3. Select the EPU harmonic using *setEPU harmo*. Examples below:
 - Use the correct harmonic to ensure best higher order rejection and optimal photon flux.
 - setEPU harmo 1 set EPU harmonic to "1"
 - * 95 850 eV
 - *setEPU harmo 3* set EPU harmonic to "3"
 - * 850 1500 eV

2.2 Setup Mono

- 1. Check the current settings of the Mono using *statMONO* or observe the parameters on the **dataGUI**.
- 2. Use moveM to select the mirror and moveG to select the grating for the desired energy range.
 - Ensure that you select the mirror coating before selecting the grating.
 - Generally, you can use the following:
 - Carbon mirror and Au LEG when using 100 eV to 250 eV energy range.
 - Nickel mirror and Ni LEG when using 200 eV to 750 eV energy range.
 - Silicon mirror and Au HEG when using 350 eV to 1500 eV energy range.
 - Examples below:
 - Carbon mirror and Au LEG grating
 - * If changing from Nickel/NiLEG:
 - (a) *moveE* 250
 - (b) moveM Carbon
 - (c) moveGAuLEG
 - * If changing from Silcon/AuHEG:
 - (a) *moveE 600*
 - (b) moveM Nickel
 - (c) moveG NiLEG
 - (d) *moveE* 250
 - (e) moveM Carbon
 - (f) moveGAuLEG
 - Nickel mirror and Ni LEG grating
 - * If changing from Carbon/AuLEG:
 - (a) *moveE 250*
 - (b) moveM Nickel
 - (c) moveG NiLEG
 - * If changing from Silcon/AuHEG:
 - (a) moveE 600
 - (b) moveM Nickel

(c) moveG NiLEG

- Silicon mirror and Au HEG grating
 - * If changing from Carbon/AuLEG:
 - (a) moveE 250
 - (b) moveM Nickel
 - (c) moveG NiLEG
 - (d) moveE 600
 - (e) moveM Silicon
 - (f) moveG AuHEG
 - * If changing from Nickel/NiLEG:
 - (a) moveE 600
 - (b) moveM Silicon
 - (c) moveG AuHEG
- 3. Move to the desired energy using moveE
 - Typically the excitation energy is 30-40 eV above the edge. Consult electron binding energy values or consult beamline staff.

Setup Spectrometer 3

- Check the current settings of the Spectrometer using *statXES* or observe the parameters on the **dataGUI**.
 - 1. Select the appropriate grating, use *setXES*.
 - Grating selection is based on a balance between efficiency and resolution.
 - Consult charts on wall for specific performance. In general:
 - * XLEG: 90 200 eV
 - · setXES XLEG
 - * LEG: 200 400 eV
 - · setXES LEG
 - * MEG: 400 700 eV
 - setXES MEG
 - * HEG: 700 1000 eV
 - · setXES HEG
 - 2. Move the spectrometer using moveXES
 - The location of the emission line is generally 10 15 lower than the absorption edge. Consult tables for binding energies or emission energies.
 - * Alternatively ask beamline staff for assistance.
 - The spectrometer requires image correction for optimal resolution and therefore, pre-work is required.
 - Use a location out of the list below (< 5 eV and < 1 deg) to ensure correct image processing. For example:
 - * moveXES 520 -1 to move to 520 eV with -1 deg offset on the LEG.
 - After movement is complete ensure the correct shift file is loaded, otherwise software will report NULL and no image correction will be used.
 - 3. Optimize the position of the spectrometer with 5 eV in later step once your sample is aligned.

SHIFT FILES AVAILABLE

• XLEG 80eV 4deg	• LEG 380eV 0deg	• LEG 575eV -1de
• XLEG 93eV 4deg	• LEG 390eV 0deg	• LEG 645eV -1de
• XLEG 114eV 4deg	• LEG 400eV 0deg	• LEG 675eV -1de
• XLEG 120eV 4deg	• LEG 455eV 0deg	• MEG 276eV 4de
• XLEG 126eV 4deg	• LEG 485eV -1deg	• MEG 380eV 1de
• LEG 117eV 2deg	• LEG 515eV -1deg	• MEG 390eV 0de

• LEG 276eV 0deg • LEG 520eV -1deg

- eg
- eg
- eg
- eg
- eg
- MEG 390eV 0deg
- MEG 400eV 0deg

- MEG 455eV 0deg
- MEG 675eV 0deg
- HEG 785eV 0deg

- MEG 485eV 0deg
- MEG 515eV 0deg
- MEG 520eV 0deg
- MEG 575eV 0deg
- MEG 785eV -1deg • HEG 645eV 2deg

• MEG 710eV 0deg

- HEG 675eV 2deg
- HEG 710eV 1deg

- HEG 860eV 0deg
- HEG 940eV 0deg
- HEG 1020eV 0deg

4 Setup Datafile

• MEG 645eV 0deg

- Before attempting any data collection, it is best to setup the file for saving data. Some tips are:
 - Do not use a new filename for each scan.
 - Do not include spaces or math symbols in the filename.
 - Group scans together either with sample and/or element measured. This will depend on the number of scans, try to keep fewer than 100 scans per file.
 - 1. Set datafile name using newfile
 - For example: *newfile test_file*
 - No extension is needed, all file extensions will be changed automatically.
 - If you wish to resume adding to existing files, just specify the name and it will change files.
 - ENSURE YOU MAKE NOTE OF THE SCAN NUMBER AND FILENAME FOR YOUR LOG BOOK

5 **Aligning Samples**

- While it is tempting to align samples only prior to the measurement, it is best to align all samples, save their positions and then proceed. This will significantly improve throughput.
- The sample alignment is achieved using the camera cross-hairs. The principle is that if the cross-hairs are co-located on the same position, the beam is there and the sample is in focus.
 - 1. Turn on the beam using setBEAM on.
 - 2. Ensure the chamber light is on using *setLAMP*.
 - Can be toggled on or off using setLAMP on and setLAMP off.
 - 3. Roughly change the sample depth to ensure the cross-hairs are properly aligned in that they appear at the same sample position.
 - Use *umvr ssd* to add or subtract sample depth. For example, *umvr ssd* 0.5 to add 0.5 mm of sample thickness.
 - 4. Roughly position the sample in the beam using a combination of *umvr ssv* and *umvr ssh*.
 - *umvr ssv* will move the sample vertical position; positive vertical moves the sample upward.
 - *umvr ssh* will move the sample horizontal position; positive horizontal moves the sample to the right.
 - It is possible to combine movement for faster alignment such as umvr ssv 1 ssh 2
 - All units are in mm.
 - 5. Use a line up scan *lup* to precisely align the sample.
 - For example lup ssh -1 1 10 1 will scan relative -1mm to 1mm with 0.2mm steps counting for 1s each point.
 - ALWAYS scan positive and ensure the step size does not fall for below 50um for ssh and 25um for ssv.
 - Make use of all detectors, tey, sdd and mcp, to assist with determining good sample positions.
 - 6. Save your sample position using setSS
 - For example: setSS Sample_1
 - Same rules for filenames, no spaces or math symbols.
 - If you want to re-save an existing sample, you will be asked to confirm the overwrite.
 - 7. Move samples using moveSS

- For example: *moveSS Sample_1*
- Sample moves can be added to the macro files for automated experiments.
- 8. Save an image of the selected location using captureSS
 - This will save an image with the beamline cameras and cross-hairs indexed by filename, Scan#, sample name and date/time.
 - Images located in ../Screenshots folder.

6 Collecting XAS Spectra

- When collecting XAS spectra, it is important to ensure that all detectors are in an acceptable range.
- Adjusting detectors is easiest done above the edge, non-resonant. However, it may be necessary to collect a quick XAS scan and then move to a resonance to adjust the detectors.
 - 1. Move above the absorption edge using moveE, typically 30 40 eV higher.
 - For K-edges, typically above the edge is 75-80 % of the maximum intensity.
 - For L-edge, you may need to a rough scan and move to the L3 resonance.
 - 2. Count for 1s using *ct* to determine the event rate on detectors.
 - 3. Adjust the count rate on the SDD_A (sdda) using the overall beamline flux, setFLUX
 - Ensure the count rate is < 100,000 cps, typically set to 80,000 cps above resonance.
 - Reduce or increase the flux using *setFLUX* where the argument is % nominal, 100% is largest.
 - For example: setFLUX 100 to set to maximum
 - For example: setFLUX 50 to reduce flux to 50% of maximum.
 - 4. One beamline flux is adjusted, adjust amplifiers using setAMP to optimize dynamic range.
 - Beamline flux meter is Mesh (i0), adjust using setAMP i0 down or setAMP i0 up.
 - Sample current meter is Sample (tey), adjust using setAMP tey down or setAMP tey up.
 - Changes the rate by a factor or 2 or 2.5.
 - Ensure rate does not exceed 1,000,000 cps.
 - Typically, set to 400,000 600,000 cps is ideal.
 - For the sample current (tey), 400,000 600,000 cps is okay above the edge for K-edge. L-edge may need to be lower.
 - Sensitivity can be set explicitly using setAMP tey 10nA/V for example. This is best for macros.
 - 5. Check the count rate once again using ct
 - These settings can be added to a macro file to speed up the experiment.
 - 6. Scan the energy using *Escan*.
 - Typically ones does a quicker rough scan to check beamline calibration and ascertain where there
 are resonances.
 - It is also advisable to always do two XAS scans to check radiation damage, either two normal scans added together to produce the final spectrum, or two quick scans followed by a longer final scan. The latter will aid in detector optimization.
 - For example: *Escan 390 395 25 405 200 415 100 425 50 5* will scan from 390 eV to 425 eV with step regions 0.2 eV, 0.05 eV, 0.1 eV and 0.2 eV with a count time per point of 5 seconds.
 - 7. Do XAS calibration prior to experiment.

6.1 Calibrating XAS Spectra

- Typically, one uses a reference material to calibrate the XAS energy.
 - 1. Select an appropriate reference material, either simple oxide or nitride.
 - 2. Measure an XAS spectrum with the required precision at the resonances used for calibration.
 - 50 meV step size when 100 700 eV
 - 100 meV step size when 700 1200 eV
 - 200 meV step size above 1200 eV
 - 3. Complete a reference XAS scan before an after the experiment per edge.
 - 4. If running multi-day experiment, complete at least once per day.

7 Collecting XES Spectra

- While XES spectra seem to be a separate measurement, they are in fact the same as XAS, in that you acquire an XES spectrum at every point.
- Typically for XES, you simple count longer at specific energy.
 - 1. Ensure that you have not left the flux reduced from previous XAS scans, setFLUX 100.
 - Typically all XES measurements are completed using full beamline flux.
 - 2. Move the beamline energy to the required energy, moveE.
 - Typically, one does at the very least one non-resonant XES, about 30-40 eV above the edge.
 - Additional excitation energies can be selected based on peaks in the corresponding XAS spectra.
 - 3. Ensure sample is in focus using cameras or pinhole.
 - Typically, the camera cross-hairs are sufficient to ensure the sample is in focus, provided the cross-hairs are located at the sample sample position.
 - If you are in doubt or have some challenge viewing your sample, you can use the pinhole to determine the focus location.
 - (a) Set pinhole in place using *setXES pinhole*.
 - (b) Count for 5s using *ct* 5.
 - (c) Toggle image on the lower left plot on the dataGUI.
 - (d) Use *umvr ssx 0.1* or *umvr ssx -0.1* successively to place the image centred vertically at channel 127.
 - (e) Save the sample location using *setSS*.
 - (f) Place the mask back in place using setXES mask.
 - (g) Un-toggle image on the lower left plot on the dataGUI.
 - 4. Determine the appropriate count time for the spectrum using ct.
 - Complete a 10s count *ct 10*.
 - Determine the total time required to achieve the require statistics.
 - Ideally, the best spectrum would have 10,000 peak counts per channel, but 1000 counts would be minimum.
 - For spectra where the point spacing is small < 50 meV, the adjacent channels can be binned to increase the count rate.
 - 5. Collect XES spectra using *Tscan*
 - Collect for a total time with 30s exposures: Tscan 360 counts for a total of 360s.
 - Collect for a total time with custom exposure time: *Tscan 360 120* counts for a total of 360s with 120s exposures.
 - Collect for a total time with custom exposure time and setting the excitation energy: *Tscan 360 120 420* counts for a total of 360s with 120s exposures at 420 eV.
 - If the energy is not specified, it will be acquired at the current beamline energy.
 - 6. Do XES calibration prior to experiment.

7.1 Calibrating XES Spectra

- The calibration of the XES occurs relative to the XAS, with the XAS used as the reference from other beamtimes.
- The calibration consists of 5 elastic peaks and two reference spectra.
 - 1. Collect resonant elastic scattering peaks on a bare material, stainless steel plate is okay, but Indium foil or Gold foil is better.
 - (a) Place beam on bare scattering target using *umvr ssh* and *umvr ssv*
 - (b) Set EPU polarization to linear vert- using setEPU polar lvn
 - (c) Select 5 energy locations distributed throughout the XES energy window. Typically one at the centre and two on either edge.
 - (d) Collect XES such that the peak counts is 1000 for each excitation.
 - (e) Set EPU polarization back to linear horizontal using *setEPU polar lh*
 - (f) Move to the reference compound and collect a non-resonant XES spectrum.
 - (g) Move excitation to the first peak and collect a resonant XES spectrum
- Complete a reference XES scan before and after the experiment per edge.
- If running multi-day experiment, complete at least once per day.

8 Collecting XEOL Spectra

- Collecting XEOL presents a challenge in that the optical output is not correlated to x-ray absorption.
- Special care needs to be taken to ensure you take quality XEOL spectra.
 - 1. Set the desired energy using *moveE*.
 - Typically XEOL is strongest at lower energy and before the XAS edge.
 - 2. Determine the correct frame rate. Set the frame rate using *setXEOL rate*, where rate is in seconds.
 - Set the rate such that the detector is not clipping at 65,000 counts, but sufficient that the spectra are good quality, > 30,000 counts.
 - For example: *setXEOL rate 10* sets the frame rate to 10s.
 - 3. Collect XEOL spectra using Tscan
 - Ensure the exposure time matches the frame rate. For example *Tscan 300 10*.
 * You can see the exposure time before the *Tscan* using *setXEOL rate*.
 - The exposure time can be a multiple of the frame rate also.
 - 4. Take a back ground spectrum with the same time rate, typically just complete the same scan with the beam off.
 - setFLUX off
 - Tscan 300 10
 - setFLUX on
 - * The background spectra need to be in the same file as the data you want to use it for, then specify background scan in notebook plotting.