

255-0650
255-5111
911
255-6757



IF IN DOUBT, ASK

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Laboratory Safety



Users are required to have passed Cornell's on-line laboratory safety training courses before they can use the Bruker X-ray detector on the Keck SEM.

Do not bring any chemicals or hazardous samples or materials into the lab.

Take specimens, stubs, and raw material with you when you leave. Do not store them in the bakeout chamber.

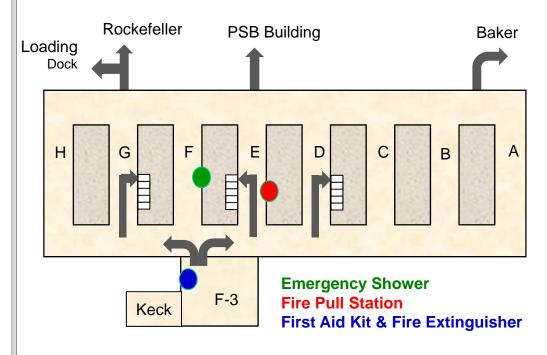
Isopropanol and Aero-Duster are supplied for your use. Wear safety glasses and read the appropriate MSDS before using them.

Nitrogen gas cylinders are changed by CCMR staff only.

Users failing to follow safety rules will be denied access to the lab.

Fire Safety

In case of fire leave immediately and close the door behind you. Do not use the elevators.





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1.05 Laboratory Policies

If ANYTHING does not even SEEM right, leave everything as it is and get or call Mick (255-0650)

DO NOT attempt to repair ANY part of the X-ray detector or remove panels, cables, or any part of the X-ray detector.

DO NOT add any software or hardware to the computer

Flash Drives (Memory Sticks) are forbidden Data MUST be removed by burning a CD or via Cornell DropBox

DO NOT change the room temperature or bring food or drink into F-3 or the SEM room

Wear gloves when venting the SEM, handling samples, or using the pumping station

If you sign up and then cannot use your time, cancel your time via Coral or contact Mick at least one weekday before your scheduled time.

Failure to show up without canceling, unless due to illness or other emergency, can result in use charges

If you are more than 30 minutes late then the microscope becomes available to other users.

If you feel sick, please reschedule your session. You will not be charged if you stay home due to illness.



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1.10 Data Storage

Data storage and safety is NOT guaranteed! - users are responsible for their data, and must copy their data in a timely fashion.

Data will only be kept on the hard drive for one year. Periodically the hard drive will be erased, leaving only one years' data there.

1.20 How to Acknowledge CCMR

When research is published using data from this microscope (including the X-ray detector), the facility and the grant number must be acknowledged:

This work made use of the electron microscopy facility of the Cornell Center for Materials Research (CCMR) with support from the National Science Foundation Materials Research Science and Engineering Centers (MRSEC) program (DMR 1120296).

2.00 Access

User Status	Signup	Room Entry	Access
Novice (< 2 sessions)	Email	Manager	M-F 8am – 5pm
Trained (>2 sessions)	Coral	Card Swipe	M-F 8am – 5pm
Experienced (>4 sessions)	Coral	Card Swipe	24/7

DO NOT use your card to let others into the SEM room



DO NOT lend your card to other users.

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3.00 About this manual

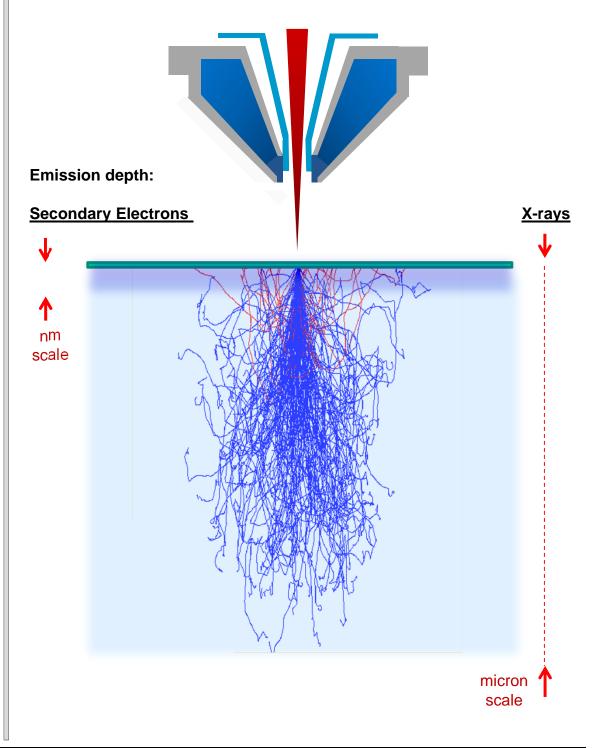
This is a **basic**, **limited use manual**, intended to help new users get started on the Bruker Quantax X-ray Detector. It provides information on basic data acquisition and X-ray mapping. It is not intended to be exhaustive, and there are many principles about X-ray analysis and features of this detector that are not covered by this manual.



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Understanding the excitation volume:

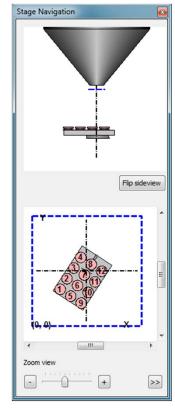
The volume that the x-rays emerge from is much larger than the spot size of the electron beam. This volume is a function of the accelerating voltage and the composition of the sample.





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How to position your sample for X-ray analysis:

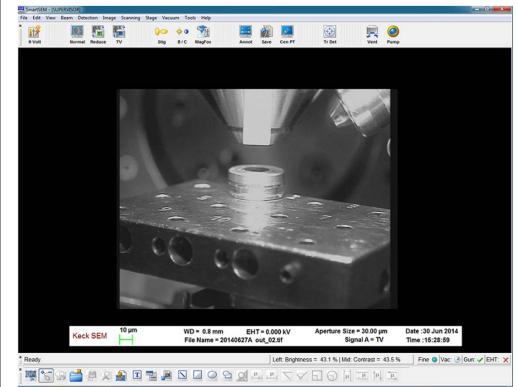


Move the sample in Z-direction with the joystick to about 7mm from the end of the pole piece as shown below.

Generally a distance between 6mm and 8mm will provide an adequate count rate, but beyond these values in either direction there will be a drop off of X-ray intensity.

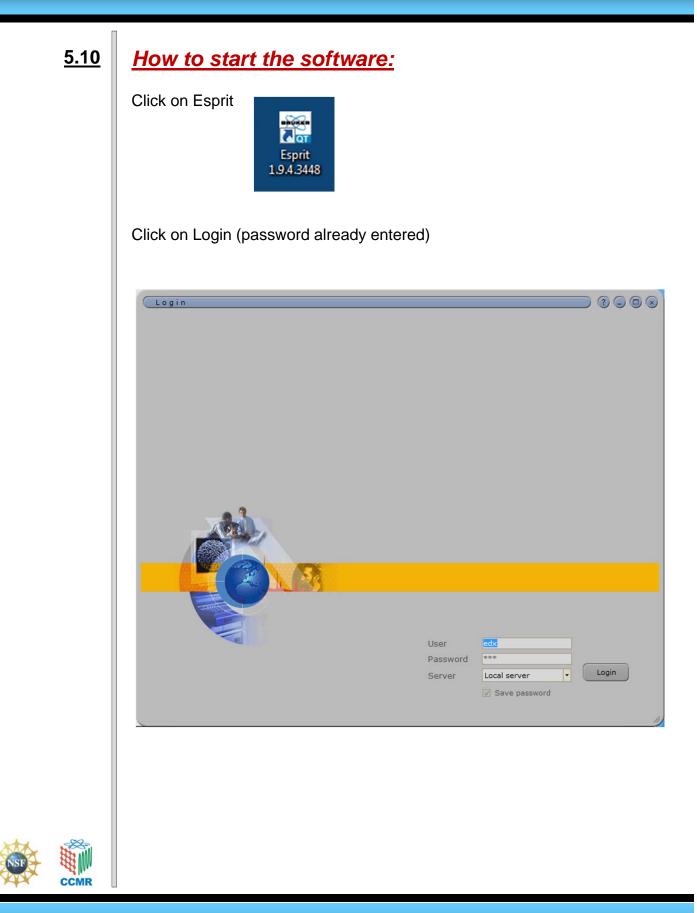
Do not trust the cartoon at left showing the gap between the sample and pole piece – your sample height will likely be different!

IF IN DOUBT, ASK





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6.00 Step 1 - setup:

Set the EHT to ~4X the highest energy edge you will be measuring.

Set the aperture to 60um to 230um aperture for high count rates.

Set the working distance to 7mm +/- 1mm.

Turn on beam in area of interest.

Click on **Spectra** Note the count rate (green circle).

Click on the triangle next to Spectrometer.

Databases	Spectra	Imaging	Feature	EBSD	Objects	Jobs	System	Report		
Spectr	a Mode)				_				
Devices										
O Spectron	(232 kcps	Rang	e 10 ke	V / 130 kcps	Dead time	40 %
	system v		1000 x 750				V 15.0		Magnification	

Choose the Max. pulse throughput to be just above the kcps value.

Choose the Maximum energy to include all the edges you are analyzing.

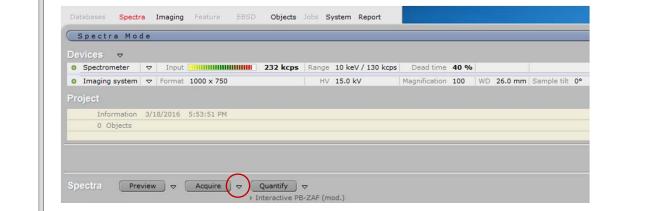
Max. pulse throughput 20 kcps 30 kcps	Maximum energy —	Mode Normal operat	ion	
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6.00 <u>Step 1 – setup (continued):</u>

Select the triangle next to Acquire



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Coating correction None Coating correction Carbon only Coating calibration Spectrum name Niobium Spectrum number: 1 Automatic numbering	Click on
ОК	

al time (or Live time) as desired. Time = Live Time + Dead Time

r a Spectrum name and check numbering then that name and al number will be assigned to ra.

OK



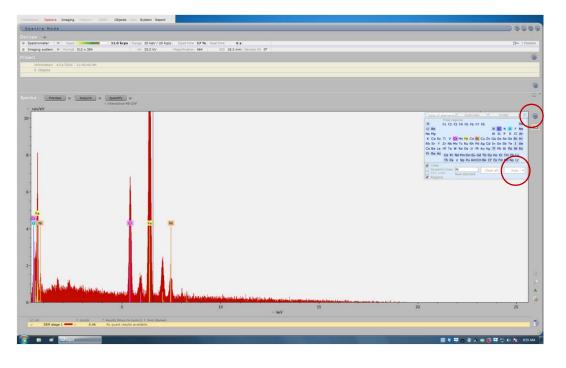
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6.00 Step 2 - acquisition:

Click on Acquire to collect spectra

The spectra will appear in real time until the acquisition has finished.

Clicking on the 🗀 icon on the right side of the screen will enable the periodic table to appear



Click on the Auto button v to automatically identify the peaks or click on the element in the Periodic Table to manually identify the peaks,



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6.00 Step 3 - quantify:

To quantify the elements present in a spectra click on Quantify

 Databases
 Spectra
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 EBSD
 Objects
 Jobs
 System Report

 Spectra
 Mode

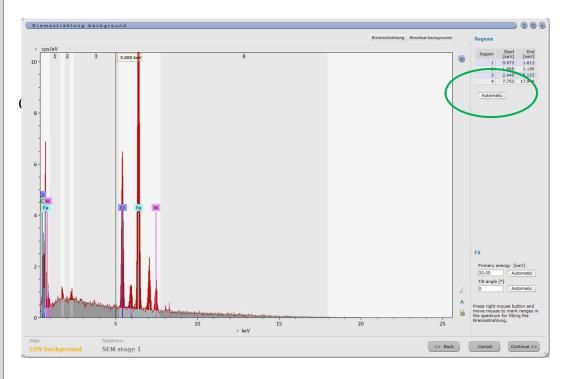
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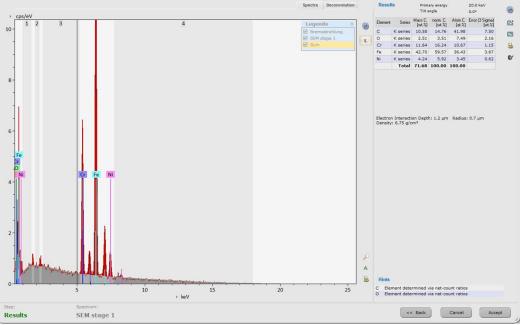


Area Mode

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6.00 Step 3 – quantify (continued):

Note a result table has showed up in the corner:



Click on Accept

Resul	ts	Prim Tilt a	ary energy ingle		20.0 ke∨ 0.0°
Element	Series	Mass C. [wt.%]	norm. C. [wt.%]	Atom C. [at.%]	Error (3 Sigma) [wt.%]
С	K series	10.58	14.76	41.98	7.50
0	K series	2.51	3.51	7.49	2.16
Cr	K series	11.64	16.24	10.67	1.15
Fe	K series	42.70	59.57	36.43	3.67
Ni	K series	4.24	5.92	3.45	0.62
	Total	71.68	100.00	100.00	

Electron Interaction Depth: 1.2 μm $\,$ Radius: 0.7 μm Density: 6.75 g/cm³ $\,$

The software displays the results in several ways.

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Note also that the electron interaction depth and radius are provided.





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<u>Step 4 – saving the data:</u>

Click on the \bigcirc icon as shown.

Area Mode

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There are four types of data that can be saved:	\langle	Element selection
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3) Result table		Сору
4) Graphic		Paste
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	\langle	Result table
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		Сору
		Save
		Add to report
	\langle	Graphic
		Сору
		Save
		Twain export



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<u>Step 4 – saving the data (continued):</u>

Sav	ving the spectra:	
		Spectrum
Und	der Spectrum click on Save…	Open
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		Save
	Save spectrum	Print
		Add to report
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		Open
	edx	Add to project
		Сору
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	Public	Save
	🍊 C:	Result table
	을 D: 금 Y:	Add to project
	⊊ Z: File name Information	Сору
•		Save
	File type Bruker Nano spectra (*.spx)	Add to report
	Bruker Nano spectra (*.spx) Text (*.txt)	Graphic
	Excel 97/2000/XP (*.xls) EMSA spectra (*.msa)	Сору
		Save
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3.		
4.	•	
	File name Information	
	File type	Save
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	Excel 97/2000/XP (*.xls) EMSA spectra (*.msa)	//
5	Under Spectrum click on Save again	
6.		
7.	The same file name should populate the File name	e box
CCMR 8.	Click on Save	

Area Mode

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<u>Step 4 – saving the data (continued):</u>

Saving the quantified results: Spectrum Under Result table click on Save... Open . . . Add to project Save results Save . . . edx\Data\ Volumes 📸 🖬 👻 🐰 🖿 💼 Preview Print . . . 눱 user Add to report £ Element selection edx Open . . . 1 Add to project Public Copy Paste 🚢 C: 🔐 D: Save . . . 👝 Y: Result table 🖵 Z: File name Information Add to project File type Copy Text table (*.txt) Save . . . Excel table (*.xls) Add to report Bitmap images (*.bmp) JPEG images (*.jpg) PNG images (*.png) TIF images (*.tif) Graphic Copy Save . . . Twain export 1. Make sure the directory is yours

- 2. Set to Text table (*.txt)
- 3. The same file name should populate the box with a .txt extension
- 4. Click on Save



Area Mode

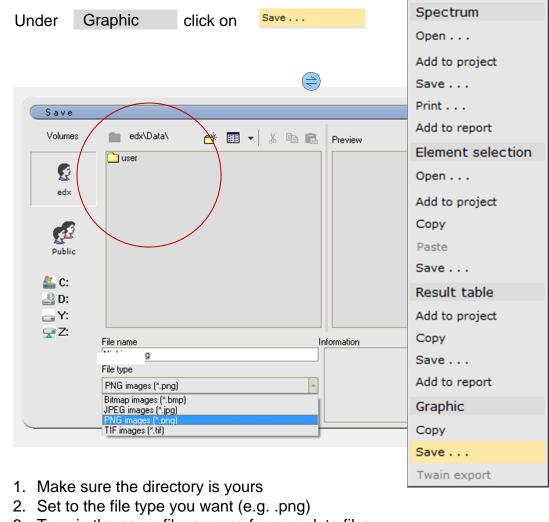
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6.00

<u>Step 4 – saving the data (continued):</u>

Saving an image of the spectra:

Important!! Saving an image file does NOT save the data!!! Image files CANNOT be quantitatively analyzed!!



- 3. Type in the same filename as for your data files
- 4. Click on Save



Area Mod

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<u>Step 1 - setup:</u>



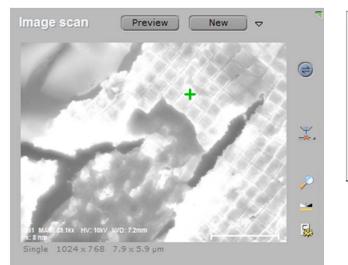
Select	Objects	and cli	ick on (Point					
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O Spectro	ometer 🛛 🗢	Input			29.4 kcps	Range	10 keV	/ 60 kcps	
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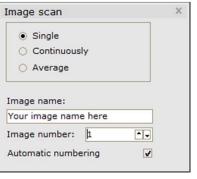
Set the EHT to ~4X the highest energy edge you will be measuring.

Set the aperture to 60um to 230um aperture for high count rates.

Set the working distance to 7mm +/- 1mm.

Click on the triangle next to New







Click on the image to set the measurement point (green crosshair).

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<u>7.00</u> <u>Step 1 – setup (continued):</u>

Note the count rate (green circle).

Click on the triangle next to Spectrometer.

Databases	Spectra	Imaging	Feature	EBSD	Objects	Jobs <mark>S</mark>	System	Report	
Point	Mode								
Devices	▽								
 Spectror 	meter (🗢	Input			29.4 kcps	Range	10 ke\	/ / 60 kcps	
Imaging	system 🗢	Format	1024 x 768			HV	10.0 k	V	N
Project									
Info	rmation 4/	1/2016 6	:35:12 PM						
0 0	bjects								
Point	MultiPoint	Line s	can Map	ping	HyperMap				

Choose the Max. pulse throughput to be just above the kcps value.

Choose the Maximum energy to include all the edges you are analyzing.

Max. pulse throughput –	Maximum energy —	Mode	
30 kcps	10 keV	 Normal opera 	ition
0 kcps	O 20 keV	Standby	Do not change
 130 kcps 275 kcps 	0 40 keV	Cooling	Mode or Cooling
0 400 kcps	0 80 keV	 Thermostat 	
 600 kcps Automatic 	O Automatic	O Maximum	
Quelles en here			
Cooling system:	On		
Detector temperature:	-30.1 °C		



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7.00 Step 1 – setup (continued):



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Measuring time	
 Automatic 	Precise
Manual	Treelse
Real time [s]	60
• Live time [s]	5
O Counts	2000
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Region end [k	eV] 20
Cyclic acquisition	n
Activate	
Count	100
Pause [s]	0
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	л
Coating correcti	on
None	
O Coating correct	tion Carbon only
O Coating calibra	ation
Spectrum name	
Niobium	
Spectrum number	: 1
Automatic number	
Automatic number	ing 😢

Set the Real time (or Live time) as desired. Note Real Time = Live Time + Dead Time

If you enter a Spectrum name and check Automatic numbering then that name and incremental number will be assigned to your spectra.

Click on OK



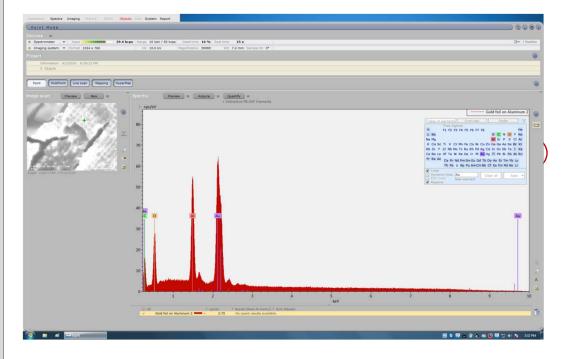
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7.00 Step 2 - acquisition:

Click on Acquire to collect spectra

The spectra will appear in real time until the acquisition has finished.

Clicking on the 🗀 icon on the right side of the screen will enable the periodic table to appear



Click on the Auto button v to automatically identify the peaks or click on the element in the Periodic Table to manually identify the peaks,



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<u>7.00</u> Step 3 - quantification:

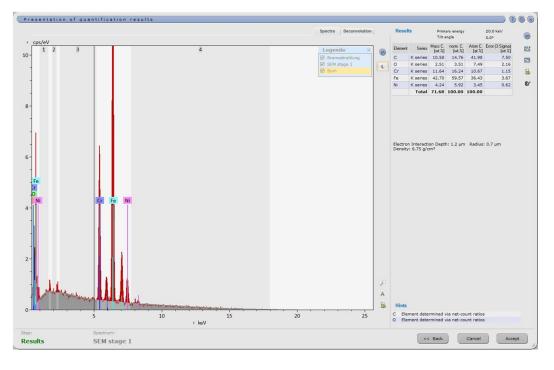
To quantify the elements present in a spectra click on Quantify

Databases Spectra Imaging Feature EBSD Objects Jobs System Report Point Mode Devices 🗢 29.4 kcps | Range 10 keV / 60 kcps | Dead time 16 % | Real time 15 s HV 10.0 kV Magnification 39088 WD 7.2 mm Sample tilt 0° Information 4/1/2016 6:35:12 PM 0 Objects MultiPoint Line scan Mapping HyperMap Point Spectra Preview Preview New 🗢 Interactive PB-ZAF linemarks Automatic Click on Bremsstrahlung background 000 Bremsstrahlung Residual background Regions Region Start [keV] End [keV] 1 0.973 1.613 2 1.888 2.188 3 2.995 5.153 10 3 4 Automatic (N Primary energy [keV] Tilt angle [°] 10 15 20 25 keV << Back Cancel Continue >>) EDS background SEM stage 1



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7.00 Step 3 – quantification (continued):



Note a result table has showed up in the corner:

Point Mode

Click on	Accept
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Resul	ts	Prim Tilt a	ary energy ingle		20.0 keV 0.0°
Element	Series	Mass C. [wt.%]	norm. C. [wt.%]	Atom C. [at.%]	Error (3 Sigma) [wt.%]
С	K series	10.58	14.76	41.98	7.50
0	K series	2.51	3.51	7.49	2.16
Cr	K series	11.64	16.24	10.67	1.15
Fe	K series	42.70	59.57	36.43	3.67
Ni	K series	4.24	5.92	3.45	0.62
	Total	71.68	100.00	100.00	

Electron Interaction Depth: 1.2 μm $\,$ Radius: 0.7 μm Density: 6.75 g/cm³ $\,$

The software displays the results in several ways.

Note also that the electron interaction depth and radius are provided.

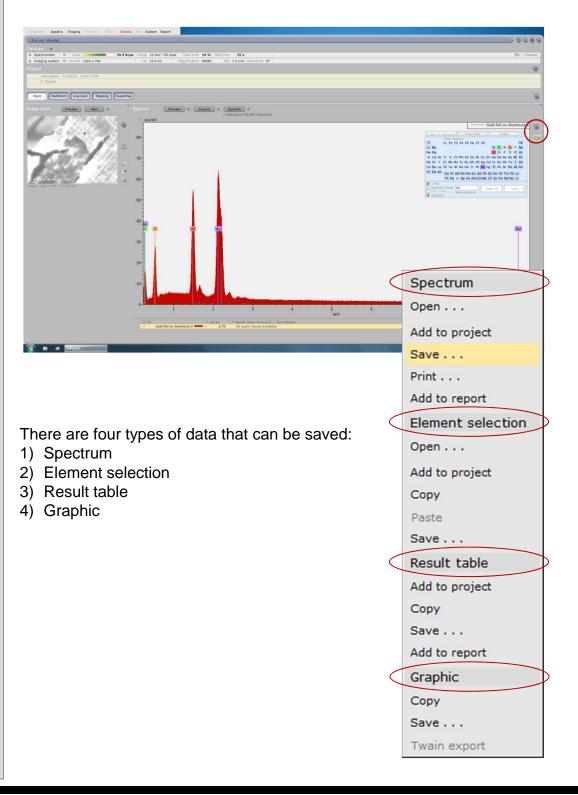


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<u> Step 4 – saving data:</u>

Click on the 😑 icon as shown.

Point Mode



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CCMI

<u>Step 4 – saving the data:</u>

Point Mode

Under Spectrum click on Save	Spectrum
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Text (*.txt) Excel 97/2000/XP (*.xls) EMSA spectra (*.msa)	Add to report
	Graphic
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 Make sure the directory is yours Set the file type to Bruker Nano spectra (*.spx) 	Save
3. Give the spectra a name	Twain export
•	
4. Click on Save	
4. CIICK ON Save	
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File name Information File type Excel 97/2000/XP (*.xls)	Save
File name Information File type Excel 97/2000/XP (*.xls) Text (*.txl) Excel 97/2000/XP (*.xls)	
File name Information File type Excel 97/2000/XP (*.xls) Bruker Nano spectra (*.spx) Text (*.txt)	
File name Information File type Excel 97/2000/XP (*.xls) Text (*.txl) Excel 97/2000/XP (*.xls)	
 File name Information File type Excel 97/2000/XP (*.xls) Bruker Nano spectra (*.spx) Text (*.txt) Excel 97/2000/XP (*.xls) EMSA spectra (*.msa) 5. Under Spectrum click on Save again	
 5. Under Spectrum click on Save again 6. Set the file type to Excel 97/2000/XP (*.xls) 	Cancel
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Mode

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<u>Step 4 – saving the data:</u>

Saving the quantified results: Spectrum Under Result table click on Save... Open . . . Add to project Save results Save . . . edx\Data\ Volumes ₩. 🎟 👻 🐰 🖿 💼 Preview Print . . . 🗋 user Add to report S Element selection edx Open . . . 22 Add to project Public Copy Paste 🚢 C: 🔐 D: Save . . . 👝 Y: Result table 🖵 Z: File name Information Add to project File type Copy Text table (*.txt) Save . . . Excel table (*.xls) Add to report Bitmap images (*.bmp) JPEG images (*.jpg) PNG images (*.png) TIF images (*.tif) Graphic Copy Save . . . Twain export 1. Make sure the directory is yours

- 2. Set to Text table (*.txt)
- 3. The same file name should populate the box with a .txt extension
- 4. Click on Save

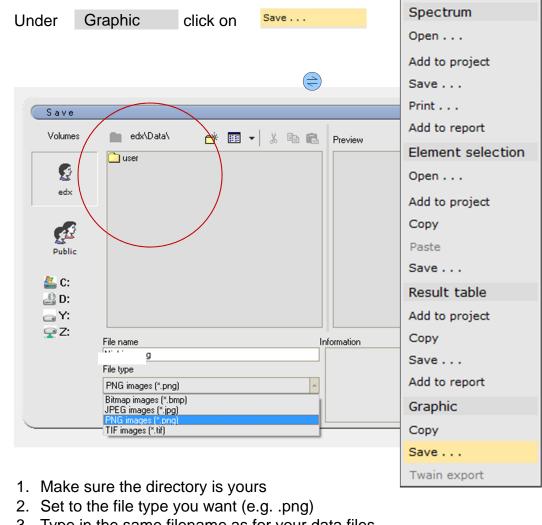


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7.00 Step 4 saving the data (continued):

Saving images of spectra:

Important!! Saving an image of the spectra does NOT save the raw data!!! Image files CANNOT be quantitatively analyzed!!



- 3. Type in the same filename as for your data files
- 4. Click on Save



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