

Elettra_PEPICO_VG — Data Analysis in Python

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Software Structure

Main program:

main.py runs the code with the parameters provided in the input section

Classes:

VG_tof.py reads in the raw files and creates lists with electron coordinates, ion

stop times, start times, and assigns corresponding events

VG_tof_only.py ion TOF spectra started by anode or random trigger

VG_figures.py plots histograms, images, spectra

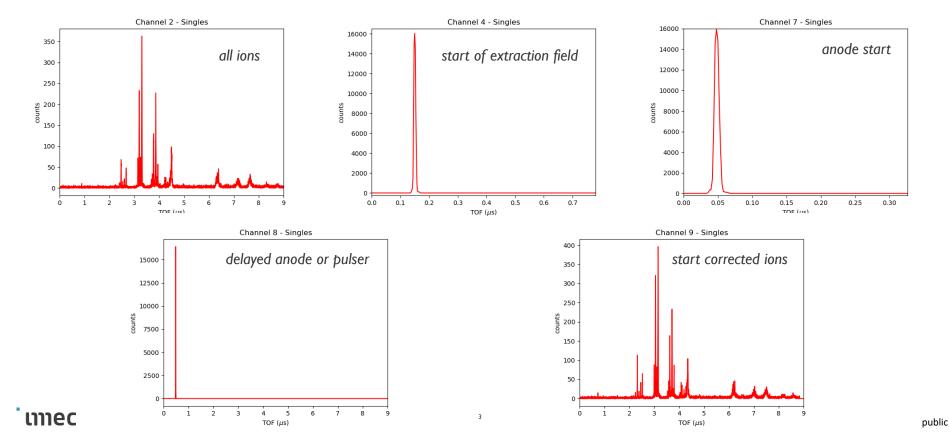
 VG_Movie_Tof.py produces electron energy selected TOF spectra and subtracts weighted TOF spectrum of randoms



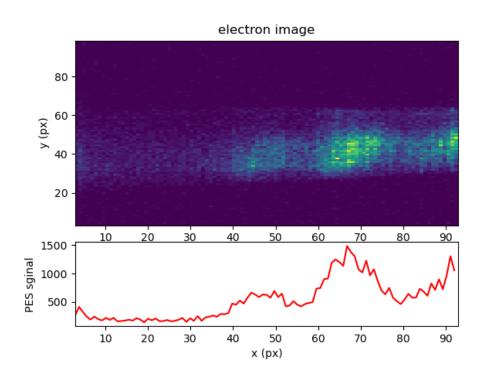
public

Output: channel histograms

Channel stops in coincidence with a complete electron event



Output: electron image



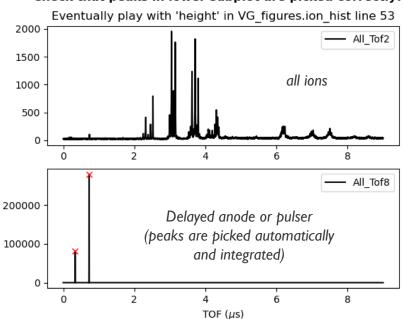
electron image in px

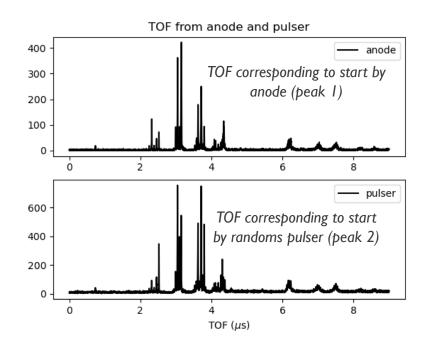
Projection on x-axis corresponding to electron energy



Output: TOF spectra (no coincidences)

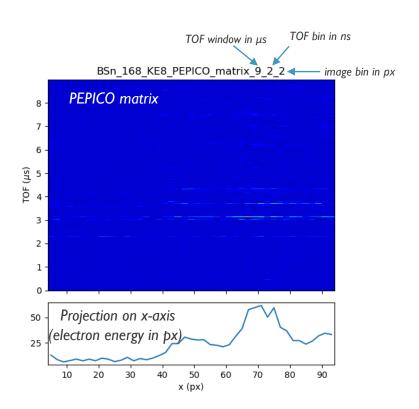
Check that peaks in lower subplot are picked correctly!

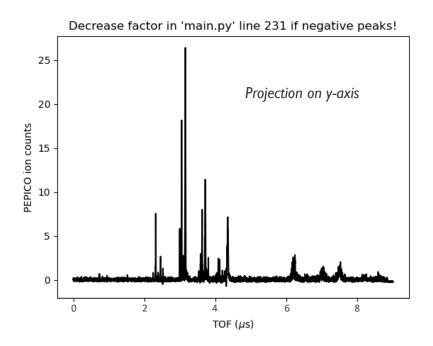






Output: PEPICO matrix (final result)







public

Output: Files

* PEPICO matrix*.dat

text file containing the final PEPICO matrix (columns = TOF, rows = electron energy)

_PEPICO_matrix.png

.png image of the PEPICO matrix figure

The 3 numbers at the end of the file name give the TOF upper limit in μs , the binning of the TOF spectrum in ns, and the binning of the electron image energy axis in px.

 $_{9}_{2}_{2}$.dat contains therefore a (45,4500) matrix, for which the energy axis runs from 0 to 90 px in steps of 2 px, and the TOF axis runs from 0 to 9 μ s in steps of 2 ns.

Putting together the kinetic energy groups

PEPICO_complete_sp.py

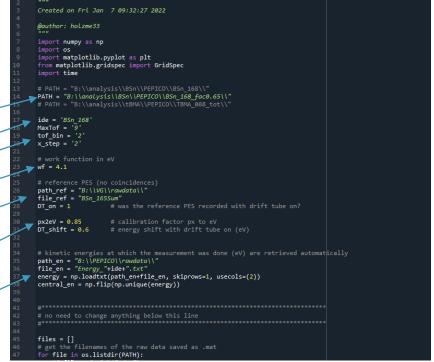
Puts together the PEPICO matrices for the different energy regions to obtain the complete matrix (overlapping regions are averaged)

Input needed in the first section

path to folder containing the input matrices
identifier of the mesurement
parameters used in the previous analysis
work function

path and file to a reference PES
energy calibration factor for PEPICO

central energies of the different regions are retrieved automatically from the Energy .txt file

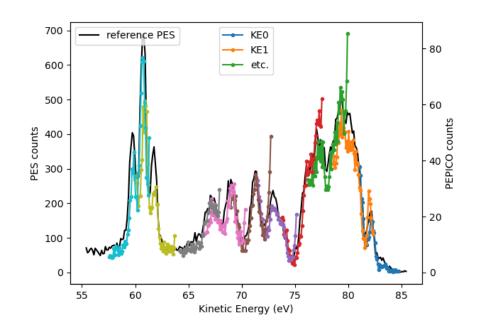


-*- coding: utf-8 -*

Check parameters, calibration, etc.

Compares the mass-integrated electron spectra for the individual energy regions to the pure photoelectron spectrum.

- Adjust drift tube shift if needed
- Adjust px2eV calibration parameter if needed

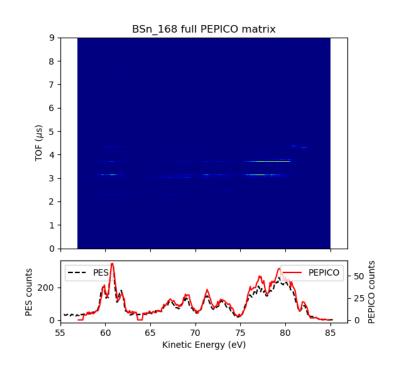




public

Combined PEPICO matrix

- Plots the combined PEPICO matrix and compares the mass-integrated electron spectrum to the reference PES
- Saves the figure as .png
- Saves the matrix and the kinetic energy axis, and the TOF axis in .txt files
- 'cut' (line 85) specifies how many points are dismissed for each energy region at the high energy side
- 'z_max' (line 88) defines the upper limit of the z axis (lower limit is by default 0)





Contact

Address questions, suggestions for extension, bug reports etc. to:

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