Quantum Metrology with Photoelectrons Vol. 3 *Analysis methodologies*

Paul Hockett

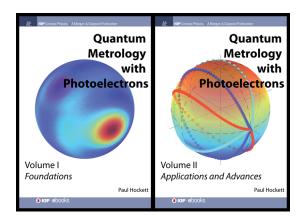
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Quantum Metrology with Photoelectrons Volume 3: *Analysis methodologies*, an open source executable book. This repository contains the source documents (mainly Jupyter Notebooks in Python) and notes for the book, as of Jan 2022 writing is in progress, and the current HTML build can be found online. The book is due to be finished in 2023, and will be published by IOP Press - see below for more details.

Series abstract

Photoionization is an interferometric process, in which multiple paths can contribute to the final continuum photoelectron wavefunction. At the simplest level, interferences between different final angular momentum states are manifest in the energy and angle resolved photoelectron spectra: metrology schemes making use of these interferograms are thus phase-sensitive, and provide a powerful route to detailed understanding of photoionization. In these cases, the continuum wavefunction (and underlying scattering dynamics) can be characterised. At a more complex level, such measurements can also provide a powerful probe for other processes of interest, leading to a more general class of quantum metrology built on phase-sensitive photoelectron imaging. Since the turn of the century, the increasing availability of photoelectron imaging experiments, along with the increasing sophistication of experimental techniques, and the availability of computational resources for analysis and numerics, has allowed for significant developments in such photoelectron metrology.

About the books



- Volume I covers the core physics of photoionization, including a range of computational examples. The material is presented as both reference and tutorial, and should appeal to readers of all levels. ISBN 978-1-6817-4684-5, http://iopscience.iop.org/book/978-1-6817-4684-5 (IOP Press, 2018)
- Volume II explores applications, and the development of quantum metrology schemes based on photoelectron measurements. The material is more technical, and will appeal more to the specialist reader. ISBN 978-1-6817-4688-3, http://iopscience.iop.org/book/978-1-6817-4688-3 (IOP Press, 2018)

Additional online resources for Vols. I & II can be found on OSF and Github.

Volume III in the series will continue this exploration, with a focus on numerical analysis techniques, forging a closer
link between experimental and theoretical results, and making the methodologies discussed directly accessible via
new software. The book is due for publication by IOP due in 2023; this volume is also open-source, with a live
HTML version at https://phockett.github.io/Quantum-Metrology-with-Photoelectrons-Vol3/ and source available
at https://github.com/phockett/Quantum-Metrology-with-Photoelectrons-Vol3.

For some additional details and motivations (including topical video), see the ePSdata project.

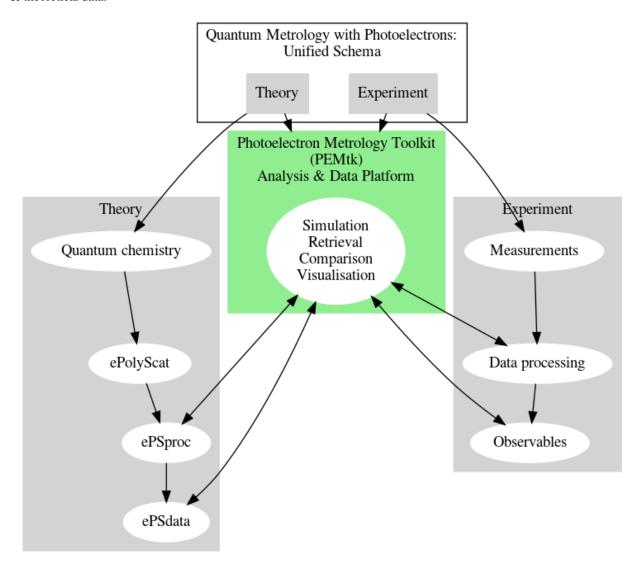
Technical details

This repository contains:

- doc-source: the source documents (mainly Jupyter Notebooks in Python)
- notes: additional notes for the book,
- the gh-pages branch contains the current HTML build, also available at https://phockett.github.io/ Quantum-Metrology-with-Photoelectrons-Vol3/

The project has been setup to use the Jupyter Book build-chain (which uses Sphinx on the back-end) to generate HTML and Latex outputs for publication from source Jupyter notebooks & markdown files.

The work *within* the book will make use of the Photoelectron Metrology Toolkit platform for working with experimental & theoretical data.



Running code examples

Each Jupyter notebook (*.ipynb) can be treated as a stand-alone computational document. These can be run/used/modified independently with an appropriately setup python environment (details to follow).

Building the book

The full book can also be built from source:

- 1. Clone this repository
- 2. Run pip install -r requirements.txt (it is recommended you do this within a virtual environment)
- 3. (Optional) Edit the books source files located in the doc-source/directory
- 4. Run jupyter-book clean doc-source/ to remove any existing builds
- 5. For an HTML build:
 - Run jupyter-book build doc-source/
 - A fully-rendered HTML version of the book will be built in doc-source/_build/html/.
- 6. For a LaTex & PDF build:
 - Run jupyter-book build doc-source/ --builder pdflatex
 - A fully-rendered HTML version of the book will be built in doc-source/_build/latex/.

See https://jupyterbook.org/basics/building/index.html for more information.

Credits

This project is created using the open source Jupyter Book project and the executablebooks/cookiecutter-jupyter-book template.

To add: build env & main software packages (see automation for this...)



Part I

Frontmatter

OVERVIEW

1.1 General overview

Vol. 3. will focus on analysis techniques for quantum metrology with photoelectrons, including:

- Interpreting experimental data.
- Extraction/reconstruction/determination of quantum mechanical properties (matrix elements, wavefunctions, density matrices) from experimental data.
- Comparison of experimental and theoretical data.
- New analysis methodologies & techniques.
- Introduction to newly-developed software platform (see below).

1.2 Provisional contents

1.2.1 Part 1: theory & software

General review & update of the topic, including recent theory developments.

- 1. Introduction
 - a. Topic overview.
 - b. Context of vol. 3 (following vols. 1 & 2).
 - c. Aims: Vol. 3 in the series will continue the exploration of quantum metrology with photoelectrons, with a focus on numerical analysis techniques, forging a closer link between experimental and theoretical results, and making the methodologies discussed directly accessible via a new software platform/ecosystem.
- 2. Quantum metrology software platform/ecosystem overview
 - a. Introduction to python packages for simulation, data analysis, and open-data.
 - b. Photoelectron metrology toolkit (PEMtk) package/platform for experimental data processing & analysis. (See *pemtk.readthedocs.io.*)
 - c. ePSproc package for theory & simulation. (See epsproc.readthedocs.io.)
 - d. ePSdata platform for data/results library (see ePSdata motivations).
- 3. General method development: geometric tensor treatment of photoionization, fitting & matrix-inversion techniques
 - a. Theory development overview tensor methods (e.g. ePSproc tensor methods)

- b. Direct molecular frame reconstruction via matrix-inversion methods (see Gregory, Margaret, Paul Hockett, Albert Stolow, and Varun Makhija. "Towards Molecular Frame Photoelectron Angular Distributions in Polyatomic Molecules from Lab Frame Coherent Rotational Wavepacket Evolution." *Journal of Physics B: Atomic, Molecular and Optical Physics* 54, no. 14 (July 2021): 145601.*DOI: 10.1088/1361-6455/ac135f.*)
- 4. Numerical implementation & analysis platform tools
 - a. Tensor methods implementation in ePSproc/PEMtk.
 - b. Information content analysis (inc. basis-set exploration, e.g. *PEMtk fitting demo*), see also vol. 2, sect. 12.1.
 - c. Density matrix analysis. (e.g. ePSproc density matrix method dev notes)
 - d. Generalised bootstrapping implementation in PEMtk (see vol. 2, sects. 11.3 & 12.3)

1.2.2 Part 2: numerical examples

Open-source worked examples using the new software platform.

- 1. Quantum metrology example: generalised bootstrapping for a homonuclear diatomic scattering system (N2)*
 - a. Experimental data overview & simulation.
 - b. Matrix element extraction (bootstrap protocol, see vol. 2, sects. 11.3 & 12.3) & statistical analysis.
 - c. Direct molecular frame reconstruction via matrix-inversion methods.
 - d. Comparison of methods.
 - e. Information content/quantum information analysis. (See vol. 2, sect. 12.1.)
- 2. Quantum metrology example: generalised bootstrapping for a heteronuclear scattering system (CO)*
 - a. Experimental data overview & simulation.
 - b. Matrix element extraction (bootstrap protocol, see vol. 2, sects. 11.3 & 12.3) & statistical analysis.
 - c. Direct molecular frame reconstruction via matrix-inversion methods.
 - d. Comparison of methods.
 - e. Information content/quantum information analysis. (See vol. 2, sect. 12.1.)
- 3. Quantum metrology example: generalised bootstrapping and matrix-inversion methods for a complex/general asymmetric top scattering system (C2H4 (ethylene))*
 - a. Experimental data overview & simulation.
 - b. Matrix element extraction (bootstrap protocol, see vol. 2, sects. 11.3 & 12.3) & statistical analysis.
 - c. Direct molecular frame reconstruction via matrix-inversion methods.
 - d. Comparison of methods.
 - e. Information content/quantum information analysis.
- 4. Future directions & outlook
- 5. Summary & conclusions
- * Exact choice of "simple" and "complex" systems may change, but should include a homonuclear diatomic and/or heteronuclear diatomic, and symmetric and asymmetric top polyatomic systems. May also include an atomic example.

Part II Theory & software

CHAPTER

TWO

INTRODUCTION

The overall aim of Vol. 3 is to expand, explore, and illustrate, quantum metrology with photoelectrons: specifically, the application of new python-based tools to tackle problems in matrix element retrieval. All the material is available as Jupyter Notebooks, which can be run locally to further explore the topic, or adapt the methodology to new problems

A brief introduction to the core physics is presented herein, although the unfamiliar reader is referred to Volume 1 of the series for a more detailed presentation, and gateway to the literature [Hoc18].

Quantum Metrology with Photoelectrons Vol. 3 *Analysis methodologies*						

Part III

Test pages

PLOTLY SURFACE RENDER TEST

Quick tests for plotly compatibility.

For JupyterLab, need additional extensions - see https://plotly.com/python/getting-started/#jupyterlab-support:

- conda install -c conda-forge -c plotly jupyter-dash
- jupyter labextension install jupyterlab-plotly

In some cases may get partially working installation with, e.g., blank surface plots, or plots via HV only. This usually means JupyterLab needs a restart (and maybe a rebuild). For more see https://plotly.com/python/troubleshooting/

Exports to HTML:

• Manual from JupyterLab OK.

3.1 Plotly example

From https://plotly.com/python/3d-surface-plots/#passing-x-and-y-data-to-3d-surface-plot

On blank plots or Javascript error "t is null". try a browser refresh and/or Jupyterlab restart. Seeing this frequently in testing March 2022, Firefox 98.0.2, JupyterLab 3.3.2, see end of page for other versions - running via Holoviews wrapper seems to be more robust here?

```
# From https://www.tutorialspoint.com/plotly/plotly_3d_scatter_and_surface_plot.htm
# Also empty
import numpy as np
```

(continues on next page)

(continued from previous page)

```
x = np.outer(np.linspace(-2, 2, 30), np.ones(30))
y = x.copy().T # transpose
z = np.cos(x ** 2 + y ** 2)
trace = go.Surface(x = x, y = y, z = z)
data = [trace]
layout = go.Layout(title = '3D Surface plot')
fig = go.Figure(data = data)
# iplot(fig)
fig.show()
```

3.2 Holoviews 3D surface test

From: https://holoviews.org/gallery/demos/plotly/surface_3d.html

See also

- http://holoviews.org/reference/elements/matplotlib/Surface.html
- http://holoviews.org/reference/elements/plotly/Surface.html

This wrapper seems robust, and works even when Plotly native rendering is giving issues?

```
import numpy as np
import holoviews as hv
hv.extension('plotly')
```

3.2.1 Define data

```
# Make data.
X = np.arange(-5, 5, 0.25)
Y = np.arange(-5, 5, 0.25)
X, Y = np.meshgrid(X, Y)
R = np.sqrt(X**2 + Y**2)
Z = np.sin(R)
surface = hv.Surface(Z, bounds=(-5, -5, 5, 5))
```

3.2.2 Plot

```
surface.opts(colorbar=True, width=500, height=500)
```

```
:Surface [x,y] (z)
```

3.2.3 Plotting with coords

BUT only supports regularly gridded coord systems (not x,y as grids). Trying to pass (X,Y,Z) in latter case gives ValueError: Surface coordinates must be 1D arrays, x and y dimension(s) were found to have multiple dimensions. Either supply 1D arrays or use the QuadMesh element for curvilinear coordinates..

```
xs = np.arange(-4, 4, 0.25)
ys = np.arange(-4, 4, 0.25)
X, Y = np.meshgrid(xs, ys)
R = np.sqrt(X**2 + Y**2)
Z = np.sin(R)
surface = hv.Surface((xs, ys, Z))
surface.opts(cmap='fire', height=500, width=500)
```

```
:Surface [x,y] (z)
```

3.3 Versions

```
import scooby
scooby.Report(additional=['xarray', 'jupyterlab','plotly','holoviews'])
```

```
Date: Mon Nov 07 17:27:07 2022 UTC
              OS : Linux
          CPU(s) : 64
        Machine : x86_64
    Architecture : 64bit
            RAM : 62.8 GiB
     Environment : Jupyter
     File system : btrfs
Python 3.9.7 | packaged by conda-forge | (default, Sep 29 2021, 19:20:46)
[GCC 9.4.0]
          xarray : 2022.3.0
      jupyterlab : 3.2.1
         plotly : 5.11.0
       holoviews: 1.15.2
          numpy: 1.20.3
           scipy : 1.7.1
        IPython : 7.28.0
      matplotlib : 3.4.3
         scooby : 0.7.0
```

March 2022 testing: Docker with jupyter/scipy-notebook base + Firefox 98.0.2.

3.3. Versions



EPSPROC BASE AND MULTIJOB CLASS INTRO

16/10/20

As of Oct. 2020, v1.3.0-dev, basic data classes are now implemented, and are now the easiest/preferred method for using ePSproc (as opposed to calling core functions directly, as illustrated in the functions guide).

A brief intro and guide to use is given here.

Aims:

- Provide unified data architecture for ePSproc, ePSdata and PEMtk.
- Wrap plotting and computational functions for ease of use.
- Handle multiple datasets inc. comparitive plots.

4.1 Setup

```
# For module testing, include path to module here, otherwise use global installation
local = True
if local:
    import sys
    if sys.platform == "win32":
        modPath = r'D:\code\github\ePSproc' # Win test machine
        winFlag = True
    else:
        modPath = r'/home/femtolab/github/ePSproc/' # Linux test machine
        winFlag = False
    sys.path.append(modPath)
# Base
import epsproc as ep
# Class dev code
from epsproc.classes.multiJob import ePSmultiJob
from epsproc.classes.base import ePSbase
```

```
* pyevtk not found, VTK export not available.
```

4.2 ePSbase class

The ePSbase class wraps most of the core functionality, and will handle all ePolyScat output files in a single data directory. In general, we'll assume:

- an ePS *job* constitutes a single ionization event/channel (ionizing orbital) for a given molecule, stored in one or more output files.
- the data dir contains one or more files, where each file will contain a set of symmetries and energies, with either
 - one file per ionizing event. In this case, each file will equate to one job, and one entry in the class datastructure.
 - a single ionizing event, where each file contains a different set of energies for the given event (*energy chunked* fileset). In this case, the files will be stacked, and the dir will equate to one job and one entry in the class datastructure.

The class datastructure is (currently) a set of dictionaries, with entries per job as above, and various data for each job. In general the data is stored in Xarrays.

The multiJob class extends the base class with reading from multple directories.

4.2.1 Load data

Firstly, set the data path, instantiate a class object and load the data.

```
# Instantiate class object.
# Minimally this needs just the dataPath, if verbose = 1 is set then some useful—
output will also be printed.
data = ePSbase(dataPath, verbose = 1)
```

```
# ScanFiles() - this will look for data files on the path provided, and read from-
them.
data.scanFiles()
```

```
*** Job orb6 details
Key: orb6
Dir D:\code\github\ePSproc\data\photoionization\n2_multiorb, 1 files.
{    'batch': 'ePS n2, batch n2_1pu_0.1-50.1eV, orbital A2',
    'event': ' N2 A-state (1piu-1)',
    'orbE': -17.096913836366,
    'orbLabel': '1piu-1'}

*** Job orb5 details
Key: orb5
Dir D:\code\github\ePSproc\data\photoionization\n2_multiorb, 1 files.
{    'batch': 'ePS n2, batch n2_3sg_0.1-50.1eV, orbital A2',
    'event': ' N2 X-state (3sg-1)',
    'orbE': -17.341816310545997,
    'orbLabel': '3sg-1'}
```

In this case, two files are read, and each file is a different ePS job - here the $3\sigma_g^{-1}$ and $1\pi_u^{-1}$ channels in N2. The keys for the job are also used as the job names.

4.2.2 Basic info & plots

A few basic methods to summarise the data...

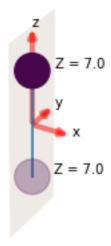
```
# Summarise jobs, this will also be output by scanFile() if verbose = 1 is set, asuillustrated above.
data.jobsSummary()
```

```
*** Job orb6 details
Key: orb6
Dir D:\code\github\ePSproc\data\photoionization\n2_multiorb, 1 files.
{    'batch': 'ePS n2, batch n2_1pu_0.1-50.1eV, orbital A2',
    'event': ' N2 A-state (1piu-1)',
    'orbE': -17.096913836366,
    'orbLabel': '1piu-1'}

*** Job orb5 details
Key: orb5
Dir D:\code\github\ePSproc\data\photoionization\n2_multiorb, 1 files.
{    'batch': 'ePS n2, batch n2_3sg_0.1-50.1eV, orbital A2',
    'event': ' N2 X-state (3sg-1)',
    'orbE': -17.341816310545997,
    'orbLabel': '3sg-1'}
```

```
# Molecular info
# Note that this is currently assumed to be the same for all jobs in the data dir.
data.molSummary()
```

```
*** Molecular structure
```



```
*** Molecular orbital list (from ePS output file)

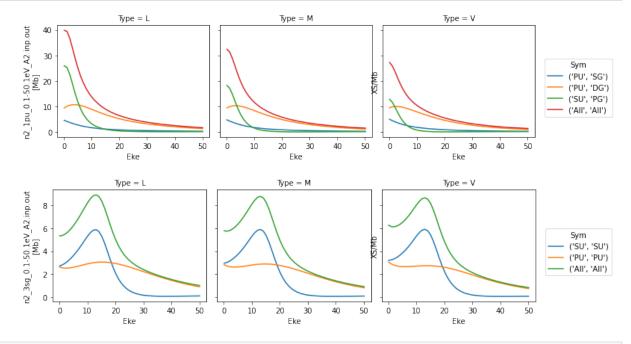
EH = Energy (Hartrees), E = Energy (eV), NOrbGrp, OrbGrp, GrpDegen = degeneracies_
and corresponding orbital numbering by group in ePS, NormInt = single centre_
expansion convergence (should be ~1.0).
```

props	s Sym	EH	0cc	E	NOrbGrp	OrbGrp	GrpDegen	NormInt
1	SG	-15.6719	2.0 -	-426.454121	1.0	1.0	1.0	0.999532
2	SU	-15.6676	2.0 -	-426.337112	1.0	2.0	1.0	0.999458
3	SG	-1.4948	2.0	-40.675580	1.0	3.0	1.0	0.999979
4	SU	-0.7687	2.0	-20.917392	1.0	4.0	1.0	0.999979
5	SG	-0.6373	2.0	-17.341816	1.0	5.0	1.0	1.000000
6	PU	-0.6283	2.0	-17.096914	1.0	6.0	2.0	1.000000
7	PU	-0.6283	2.0	-17.096914	2.0	6.0	2.0	1.000000

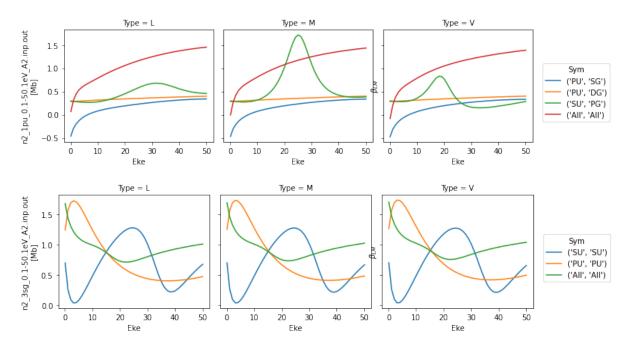
4.2.3 Plot cross-sections and betas

These are taken from the GetCro segments in the ePS output files, and correspond to results for an isotropic ensemble of molecules, i.e. observables in the lab frame (LF) for 1-photon ionization (see the ePS tutorial for more details).

```
# Minimal method call will plot cross-sections for all ePS jobs found in the data-
directory.
data.plotGetCro()
```



```
# Plot beta parameters with the 'BETA' flag
data.plotGetCro(pType = 'BETA')
```



TODO: fix labelling here.

4.2.4 Compute MFPADs

The class currently wraps just the basic numerical routine for MFPADs. This defaults to computing MFPADs for all energies and (z, x, y) polarization geometries (where the z-axis is the molecular symmetry axis, and corresponds to the molecular structure plot shown above).

```
# Compute MFPADs...
data.mfpadNumeric()
```

To plot it's advisable to set an enery slice, Erange = [start, stop, step], since MFPADs are currently shown as individual plots in the default case, and there may be a lot of them.

We'll also just set for a single key here, otherwise all jobs will be plotted.

```
data.padPlot(keys = 'orb5', Erange = [5, 10, 4])
#TODO: fix plot layout!
```

```
Found dims ('Labels', 'Phi', 'Theta', 'Eke', 'Sym'), summing to reduce for plot...

Pass selDims to avoid.

Sph plots: Pol geom: z

Plotting with mpl

Data dims: ('Phi', 'Theta', 'Eke'), subplots on Eke

Sph plots: Pol geom: x

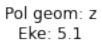
Plotting with mpl

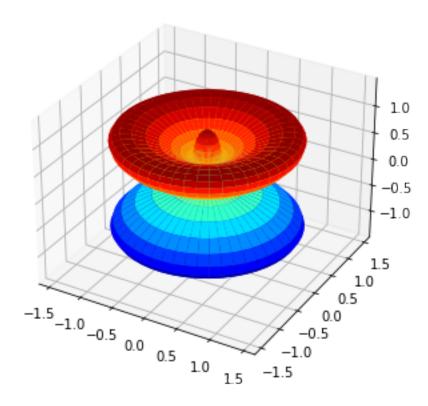
Data dims: ('Phi', 'Theta', 'Eke'), subplots on Eke

Sph plots: Pol geom: y

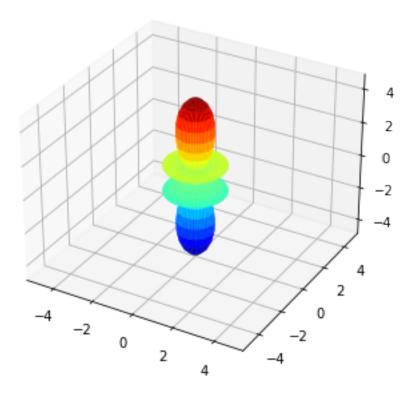
Plotting with mpl

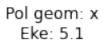
Data dims: ('Phi', 'Theta', 'Eke'), subplots on Eke
```

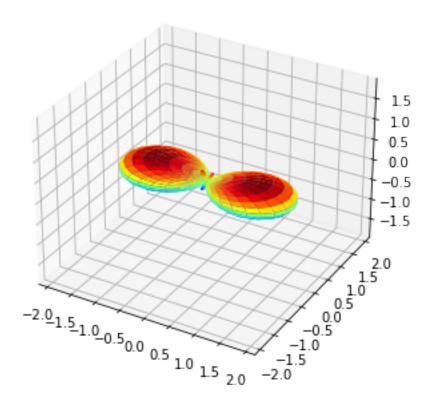


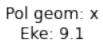


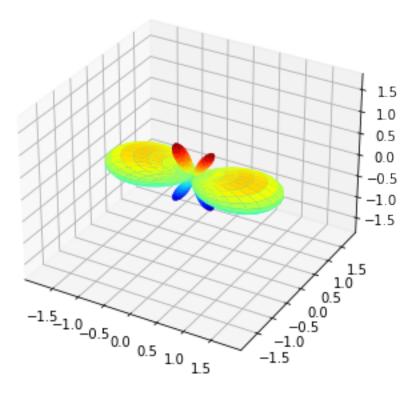
Pol geom: z Eke: 9.1

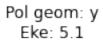


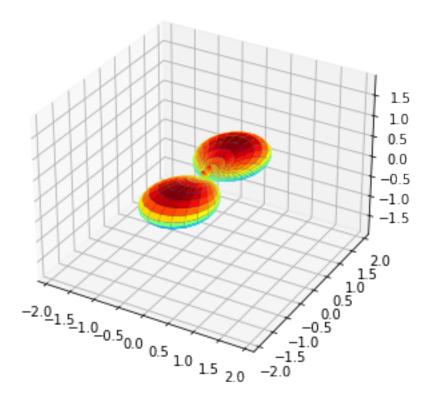


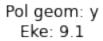


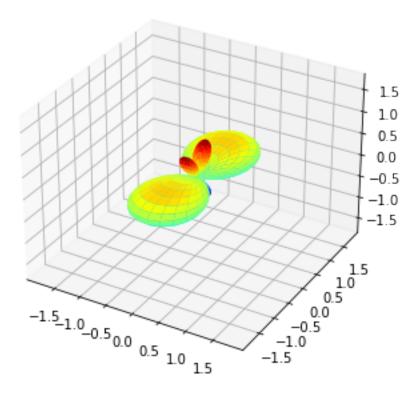










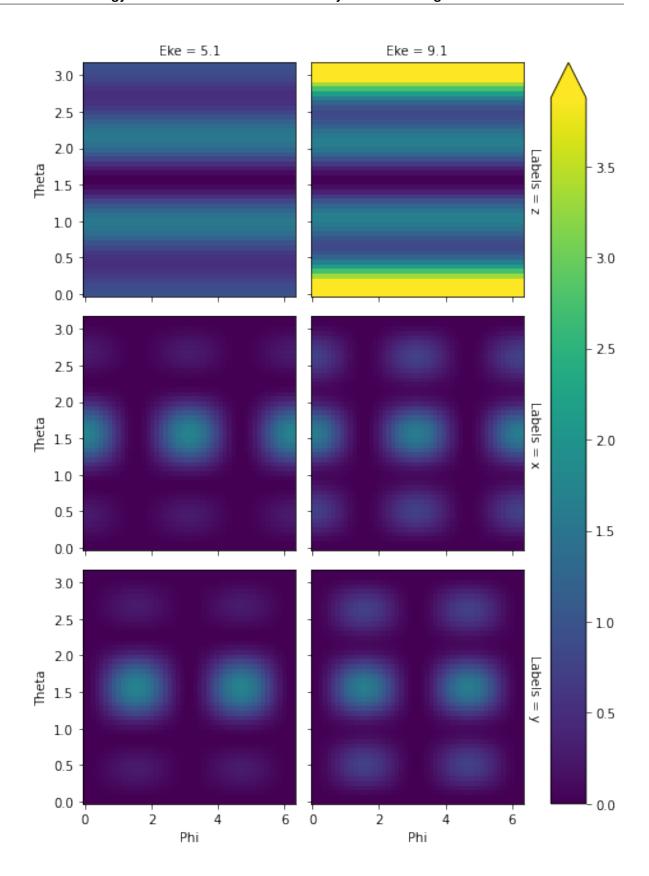


To view multiple results in a more concise fashion, a Cartesian gridded output is also available.

```
data.padPlot(keys = 'orb5', Erange = [5, 10, 4], pStyle='grid')
```

Found dims ('Labels', 'Phi', 'Theta', 'Eke', 'Sym'), summing to reduce for plot. Pass selDims to avoid.

Grid plot: 3sg-1



```
# Various other args can be passed...
# Set a plotting backend, currently 'mpl' (Matplotlib - default) or 'pl' (Plotly --
interactive, but may give issues in some environments)
backend = 'pl'
# Subselect on dimensions, this is set as a dictionary for Xarray selection (see-
http://xarray.pydata.org/en/stable/indexing.html#indexing-with-dimension-names)
selDims = {'Labels':'z'} # Plot z-pol case only.

data.padPlot(Erange = [5, 10, 4], selDims=selDims, backend = backend)
```

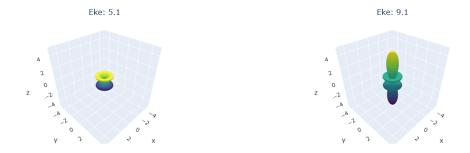
```
Found dims ('Phi', 'Theta', 'Eke', 'Sym'), summing to reduce for plot. PassuselDims to avoid.

Sph plots: 1piu-1
Plotting with pl
```



```
Found dims ('Phi', 'Theta', 'Eke', 'Sym'), summing to reduce for plot. PassuselDims to avoid.

Sph plots: 3sg-1
Plotting with pl
```



4.2.5 Compute β_{LM} parameters

For computation of β_{LM} parameters the class wraps functions from epsproc.geom, which implement a tensor method. This is quite fast, although memory heavy, so may not be suitable for very large problems. (See the method development pages for more info, more concise notes to follow).

- Functions are provided for MF and AF problems (which is the general case, and will equate to the LF case for an unaligned ensemble).
- For the MF the class wraps ep.geom.mfblmXprod, see the method development page for more info, more concise notes to follow.
- For the AF the class wraps ep.geom.afblmXprod, see the method development page for more info, more
 concise notes to follow.

Compute MF β_{LM} and PADs

Here's a quick demo for the default MF cases, which will give parameters corresponding to the (z, x, y) polarization geometries computed by the numerical routine above.

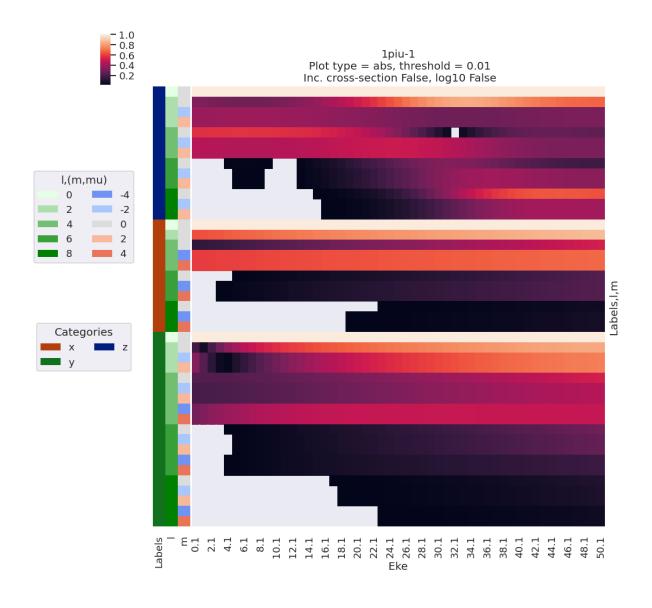
```
Calculating MF-BLMs for job key: orb6
Return type BLM.

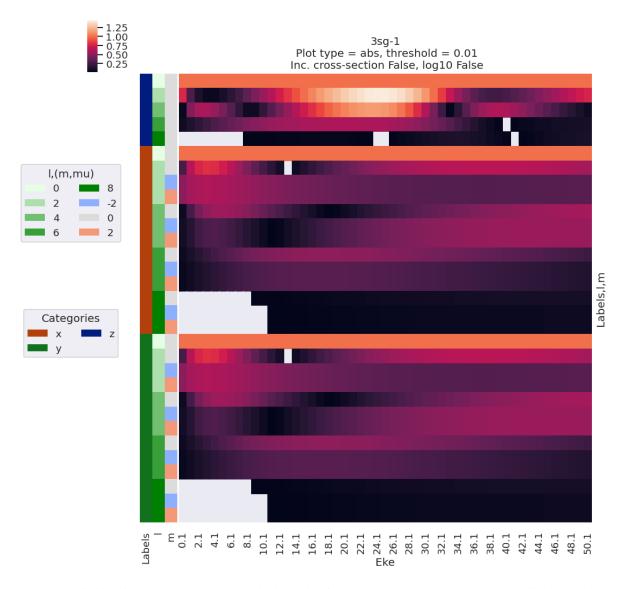
Calculating MF-BLMs for job key: orb5
Return type BLM.
```

Plotting still needs to improve... but ep.lmPlot() is a robust way to plot everything.

```
# data.BLMplot(dataType = 'MFBLM') # HORRIBLE OUTPUT at the moment!!!
data.lmPlot(dataType = 'MFBLM')
```

```
Plotting data n2_1pu_0.1-50.1eV_A2.inp.out, pType=a, thres=0.01, with Seaborn Plotting data n2_3sg_0.1-50.1eV_A2.inp.out, pType=a, thres=0.01, with Seaborn
```

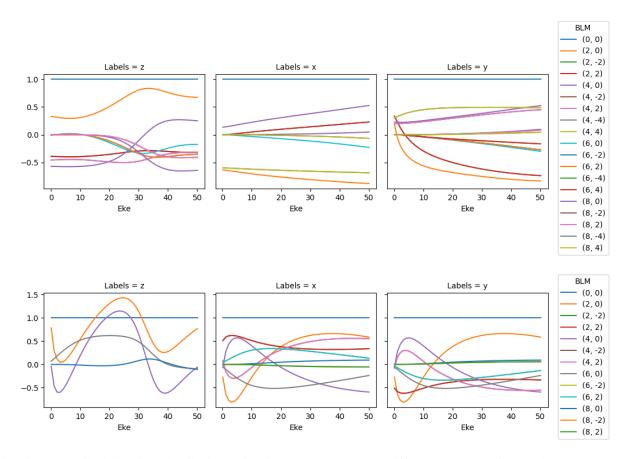




Line-plots are available with the BLMplot method, although this currently only supports the Matplotlib backend, and may have problems with dims in some cases (work in progress!).

```
data.BLMplot(dataType='MFBLM', thres = 1e-2) # Passing a threshold value here will → remove any spurious BLM parameters.
```

```
Dataset: orb6, 1piu-1, MFBLM Dataset: orb5, 3sg-1, MFBLM
```



Polar plots are available for these distributions using the padPlot () method if the dataType is passed.

```
data.padPlot(keys = 'orb5', Erange = [5, 10, 4], dataType='MFBLM')
```

```
Using default sph betas.

Sph plots: Pol geom: z

Plotting with mpl

Data dims: ('Eke', 'Phi', 'Theta'), subplots on Eke

Sph plots: Pol geom: x

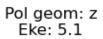
Plotting with mpl

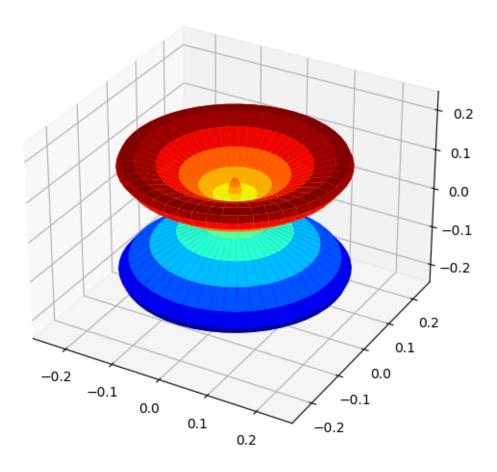
Data dims: ('Eke', 'Phi', 'Theta'), subplots on Eke

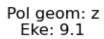
Sph plots: Pol geom: y

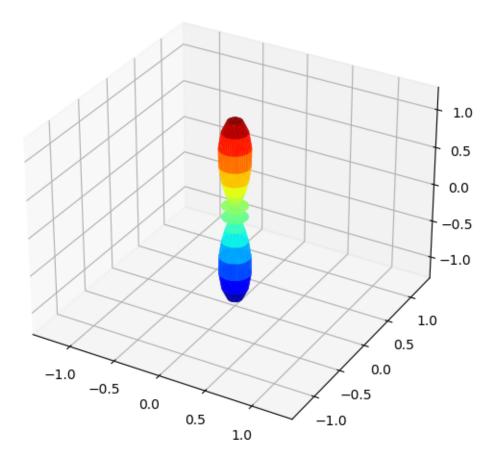
Plotting with mpl

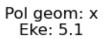
Data dims: ('Eke', 'Phi', 'Theta'), subplots on Eke
```

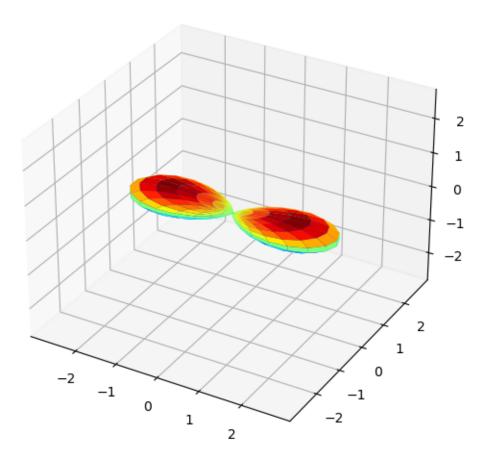


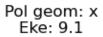


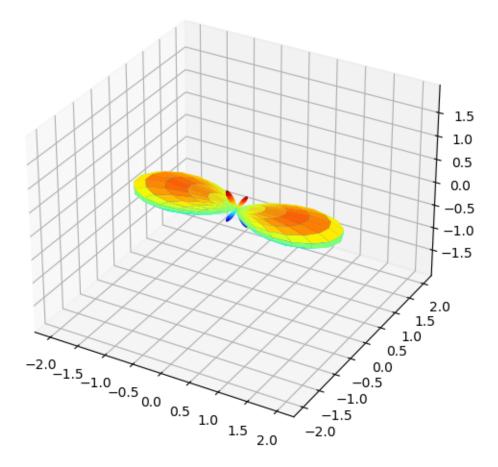


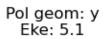


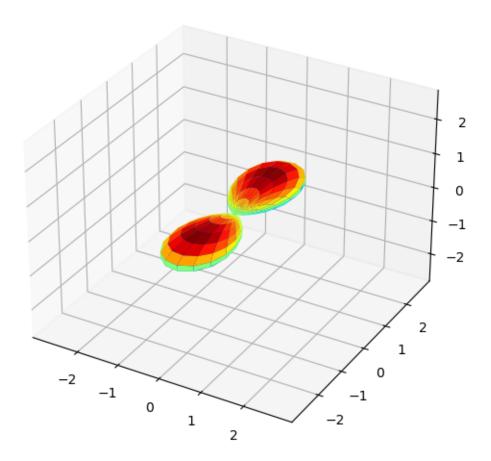


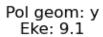


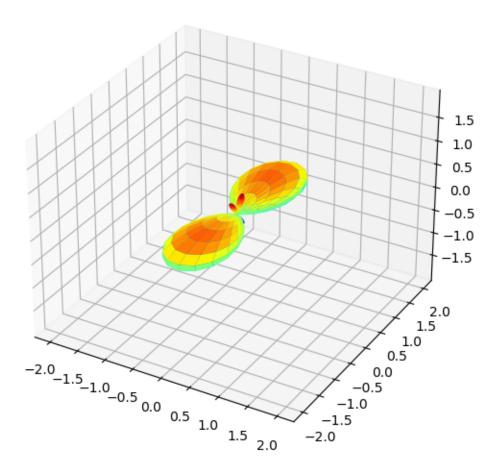












Compute LF/AF β_{LM} and PADs

Here's a quick demo for the default AF case (isotropic distribution, hence == LF case).

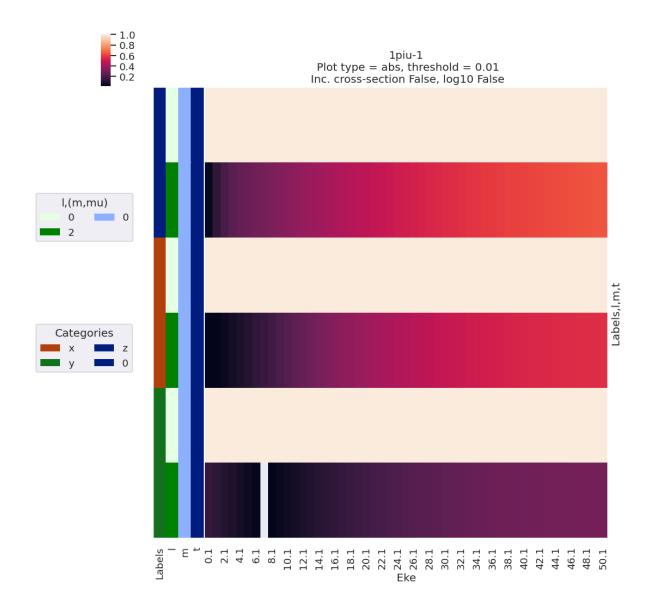
```
data.AFBLM()
```

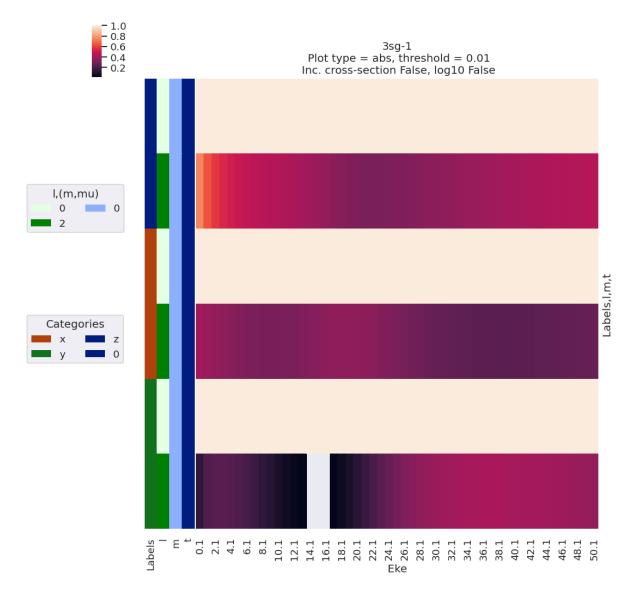
```
Calculating AF-BLMs for job key: orb5
```

Plotting still needs to improve... but ep.lmPlot() is a robust way to plot everything.

```
# data.BLMplot(dataType = 'MFBLM') # HORRIBLE OUTPUT at the moment!!!
data.lmPlot(dataType = 'AFBLM')
```

```
Plotting data n2_1pu_0.1-50.1eV_A2.inp.out, pType=a, thres=0.01, with Seaborn Plotting data n2_3sg_0.1-50.1eV_A2.inp.out, pType=a, thres=0.01, with Seaborn
```

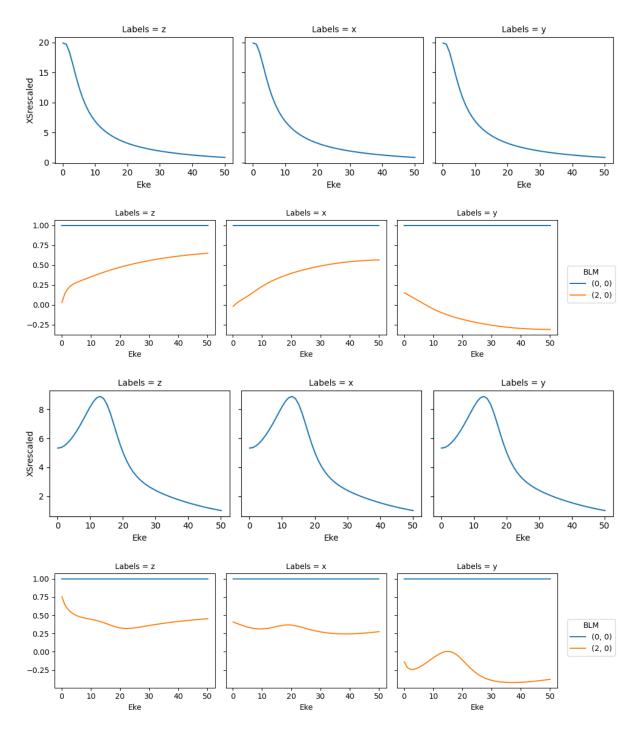




Line-plots are available with the BLMplot method, although this currently only supports the Matplotlib backend, and may have problems with dims in some cases (work in progress!).

```
data.BLMplot(dataType='AFBLM', thres = 1e-2) # Passing a threshold value here will-
-remove any spurious BLM parameters.
```

```
Dataset: orb6, 1piu-1, XS
Dataset: orb6, 1piu-1, AFBLM
Dataset: orb5, 3sg-1, XS
Dataset: orb5, 3sg-1, AFBLM
```



Polar plots are available for these distributions using the padPlot () method.

```
data.padPlot(keys = 'orb5', Erange = [5, 10, 4], dataType='AFBLM')

# NOTE - seem to have an inconsistency with (x,y) pol geometries here - should check-
source code & fix. Likely due to mix-up in frame defns., i.e. probably mixing LF-
and MF pol geom defn. - TBC.
```

```
Found dims ('Labels', 't', 'Eke', 'BLM'), summing to reduce for plot. Pass selDims—
to avoid.

Using default sph betas.

Sph plots: Pol geom: z

Plotting with mpl

Data dims: ('t', 'Eke', 'Phi', 'Theta'), subplots on Eke

Sph plots: Pol geom: x

Plotting with mpl

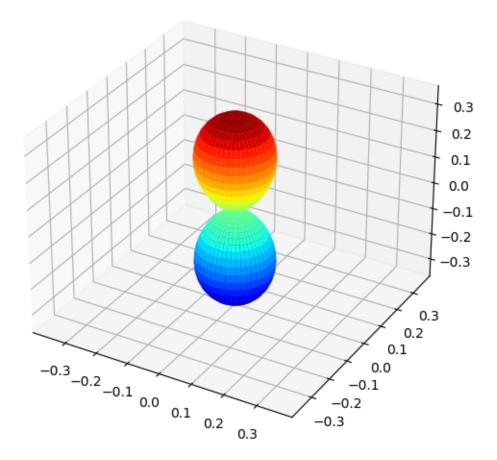
Data dims: ('t', 'Eke', 'Phi', 'Theta'), subplots on Eke

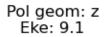
Sph plots: Pol geom: y

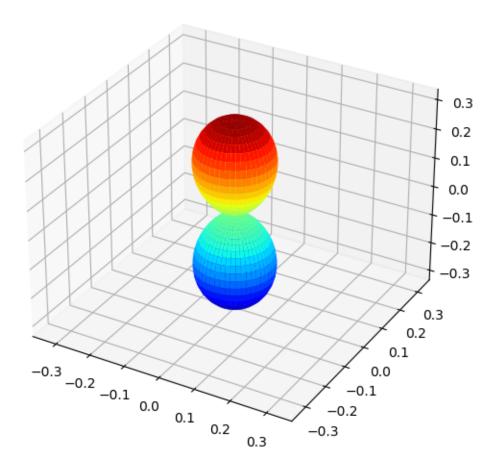
Plotting with mpl

Data dims: ('t', 'Eke', 'Phi', 'Theta'), subplots on Eke
```

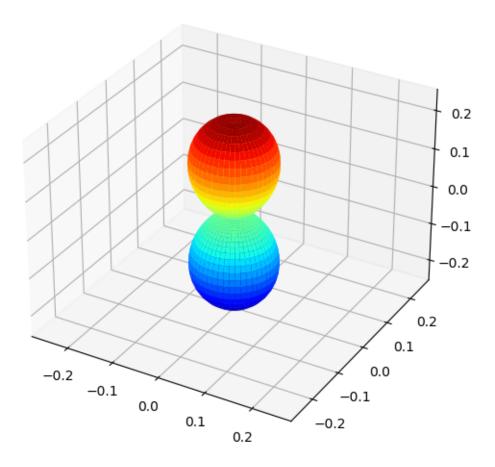
Pol geom: z Eke: 5.1

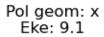


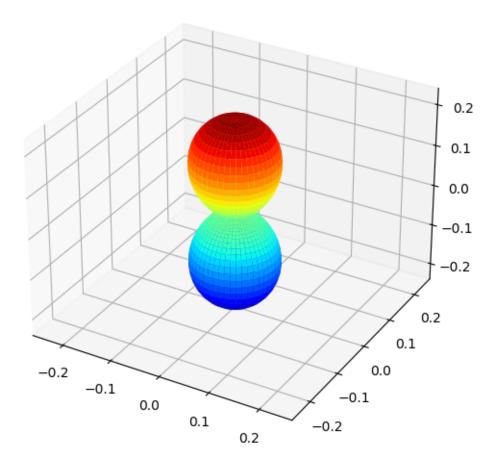


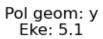


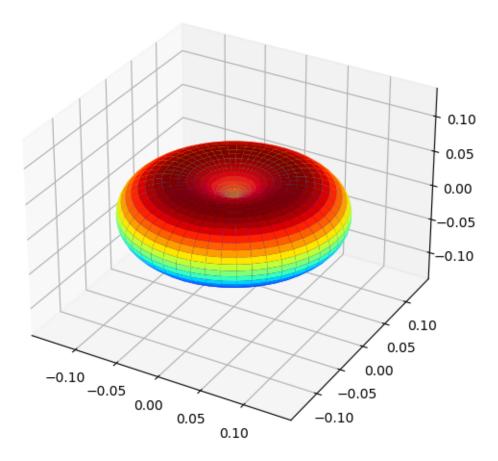
Pol geom: x Eke: 5.1



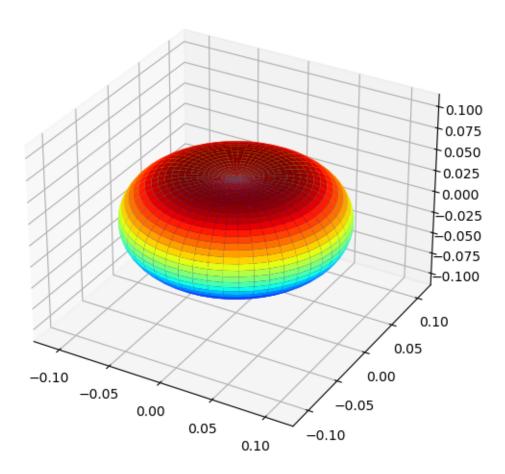












4.3 Additions

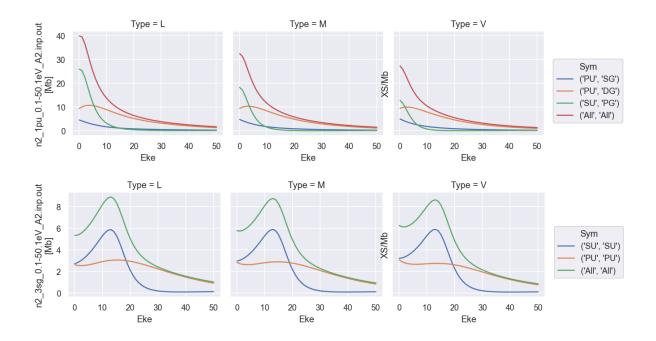
4.3.1 Plot styles for line-plots

To set to Seaborn plotting style, use ep.hvPlotters.setPlotters() (note this will be set for all plots after loading, unless overriden). Seaborn must be installed for this to function.

For more on Seaborn styles, see the Seaborn docs.

```
from epsproc.plot import hvPlotters
hvPlotters.setPlotters()

data.plotGetCro()
```



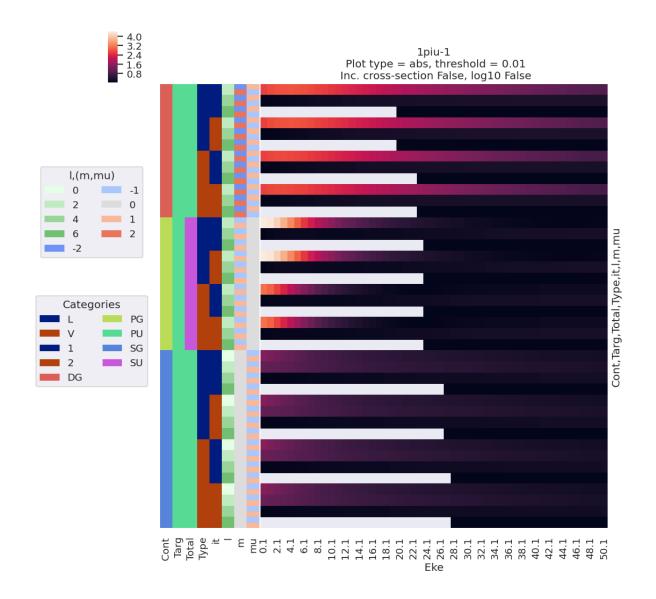
4.3.2 Matrix element plotting

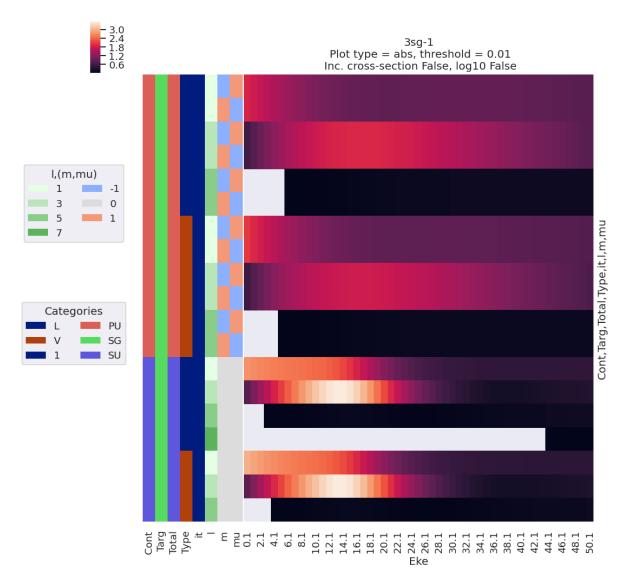
For a full view of the computational results, use the lmPlot() method with the default, which correspond to dataType=matE.

```
data.lmPlot()
```

Plotting data n2_1pu_0.1-50.1eV_A2.inp.out, pType=a, thres=0.01, with Seaborn Plotting data n2_3sg_0.1-50.1eV_A2.inp.out, pType=a, thres=0.01, with Seaborn

4.3. Additions 51



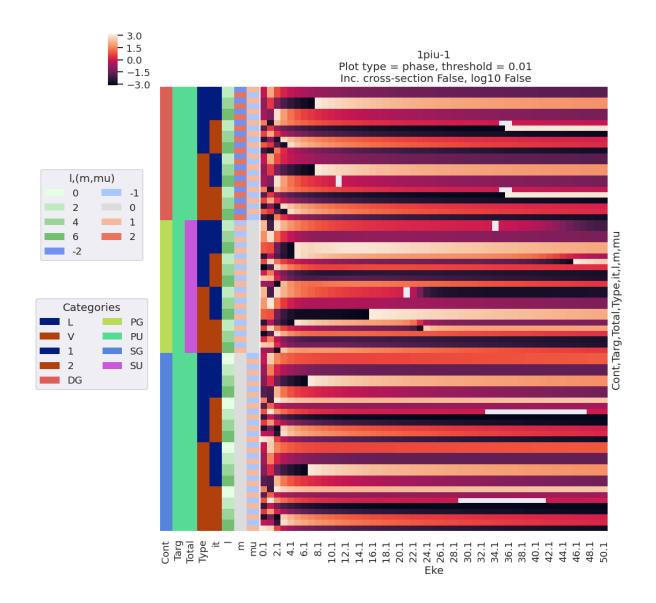


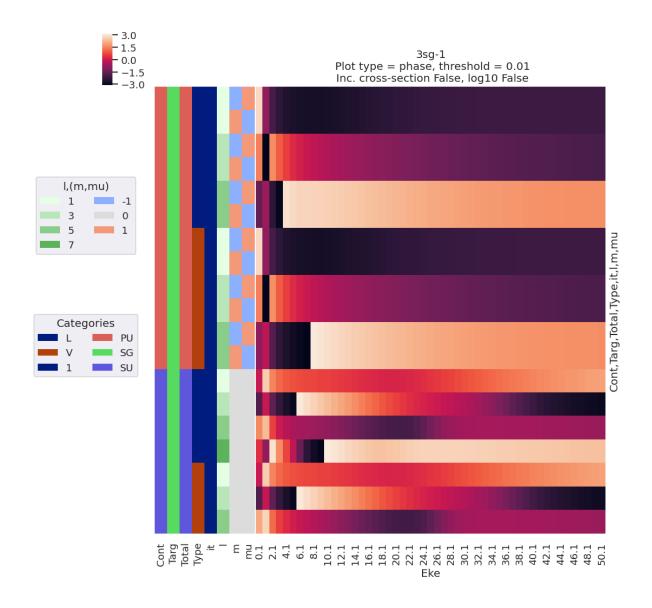
The default here plots abs values, but the same routine can be set for phase plotting.

```
data.lmPlot(pType='phase')
```

Plotting data n2_1pu_0.1-50.1eV_A2.inp.out, pType=phase, thres=0.01, with Seaborn Plotting data n2_3sg_0.1-50.1eV_A2.inp.out, pType=phase, thres=0.01, with Seaborn

4.3. Additions 53





4.4 Versions

```
import scooby
scooby.Report(additional=['epsproc', 'xarray', 'jupyter'])

Date: Tue Oct 20 15:57:33 2020 Eastern Daylight Time

OS: Windows
CPU(s): 32
Machine: AMD64
Architecture: 64bit
RAM: 63.9 GB
Environment: Jupyter

(continues on next page)
```

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CHAPTER

FIVE

EPSPROC MATLAB DEMO

As noted in the main readme/intro doc, ePSproc originally started as a set of Matlab routines, before migration to Python in 2019. Although no longer maintained/updated since ~2018, the Matlab routines are still available as part of the ePSproc distribution. This notebook demos the routines, following the demo script ePSproc_NO2_MFPADs_demo.m (to run this code in a Jupyter environment, a Matlab kernel is required, e.g. Calysto's Matlab kernel).

Functionality:

- Read raw photoionization matrix elements from ePS output files with "dumpIdy" segments
- Calculate MF-PADs from the matrix elements (ePSproc_MFPAD.m, see also ePSproc_NO2_MFPADs_demo.m)
- Plot MF-PADs
- · Plot X-sects
- (Beta testing): Calculate MF-BLMs from matrix elements, see ePSproc_MFBLM.m
- (Under development): Calculate AF-BLMs from matrix elements.

Source:

- /matlab: stable matlab code (as per release v1.0.1 https://github.com/phockett/ePSproc/releases).
 - a set of functions for processing (ePSproc*.m files)
 - a script showing demo calculations, ePSproc_NO2_MFPADs_demo.m
- /docs/additional contains:
 - the benchmark results from these calculations, ePSproc NO2 testing summary 250915.pdf
 - additional notes on ePS photoionization matrix elements, ePSproc_scattering_theory_ePS_notes_011015. pdf.

See ePSproc: Post-processing suite for ePolyScat electron-molecule scattering calculations _formore details">https://www.authorea.com/users/71114/articles/122402/_show_article>_formore details.

5.1 Setup

```
cd('~/github/ePSproc') % Change to ePSproc root dir.
  ans =
       '9.4.0.813654 (R2018a)'
%% *** SETTINGS
% Set up basic environment
% Name & path to ePS output file. In this version set full file name here, and
⇔working directory below.
% fileName='no2_demo_ePS.out' % OK for MFPAD testing, but only single E point
fileName='n2_3sg_0.1-50.1eV_A2.inp.out'
% Set paths for Linux or Win boxes (optional!)
if isunix
   dirSlash='/';
else
   dirSlash='\';
end
filePath=[pwd dirSlash 'data' dirSlash 'photoionization'];
→Root to working directory, here set as current dir/data/photoionization
fileBase=[filePath dirSlash fileName]; % Full path to ePS results file, here set as_
⇔current working direcory
scriptPath=[pwd dirSlash 'matlab' dirSlash]; % Add path to ePSproc scrips to Matlab_
⇒path list, here set as current dir/matlab
path (path, [scriptPath]);
  fileName =
       'n2_3sg_0.1-50.1eV_A2.inp.out'
%% *** Read data
% Variables:
      rlAll contains matrix elements (from DumpIdy segments)
      params contains various calculation parameters
       getCro contains cross-section (from GetCro segments), if present
[rlAll, params, getCro] = ePSproc_read(fileBase);
params.fileBase=fileBase;
params.fileName=fileName;
  *** Reading ePS output file
  /home/paul/github/ePSproc/data/photoionization/n2_3sg_0.1-50.1eV_A2.inp.out
  Found 102 sets of matrix elements
  Read 102 sets of matrix elements (0 blank records)
  Found 2 symmetries
      'SU' 'PU'
                                                                         (continues on next page)
```

(continued from previous page)

```
Found 51 energies
Found 2 atoms
Found 18 data records
Found 3 sets of cross sections
```

```
% Matrix elements are stored in a structure rlAll
```

```
rlAll =

2x51 struct array with fields:

    eKE
    PE
    symm
    symmSet
    MbNorm
    rawIdyHead
    rawIdy1
    rlnlHead
    rlnl1
    rawIdy2
    rlnl2
    pWaveAll
```

```
% GetCro outputs (total cross-secions, LF betas)
getCro
```

```
getCro =
  1x3 struct array with fields:
    GetCro
```

```
% General calculation params
params
```

```
params =

struct with fields:

symmList: {'SU' 'PU' 'All'}

    eKE: [1x51 double]

symmAll: {1x102 cell}

    eAll: [1x102 double]

nRecords: 102

nEnergies: 51

nSymms: 2

gLmax: 11

blankRec: 0

missingE: [1x0 double]
```

(continues on next page)

5.1. Setup 59

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```
pWaveAll: [144x51x3 double]
pWaveAllMb: [144x51x3 double]
LMallInd: [144x2 double]
    coords: {1x5 cell}
dataRecords: {18x2 cell}
    IP: 15.5800
GetCroHeader: {{1x1 cell}}
    fileBase: '/home/paul/github/ePSproc/data/photoionization/n2_3sg_0.1-50.
41eV_A2.inp.out'
    fileName: 'n2_3sg_0.1-50.1eV_A2.inp.out'
```

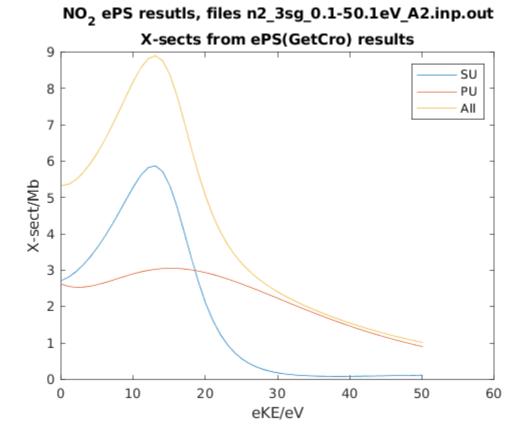
5.2 Plot cross-sections and betas

These are taken from the GetCro segments in the ePS output files, and correspond to results for an isotropic ensemble of molecules, i.e. observables in the lab frame (LF) for 1-photon ionization (see the ePS tutorial for more details).

```
Warning: MATLAB has disabled some advanced graphics rendering features by switching to software OpenGL. For more information, click <a href="matlab:opengl(solution-problems")">here</a>.

Warning: Ignoring extra legend entries.

In legend>set_children_and_strings (line 646)
In legend>make_legend (line 316)
In legend (line 259)
```



5.3 MFPADs

These are calculated numerically from the matrix elements, for a given polarization geometry and symmetry (the method is the same as the python version of the routine).

```
%% *** Calculate MFPADs - single polarization geometry, all energies and symmetries
% Calculate for specified Euler angles (polarization geometry) & energies
% Set resolution for calculated I (theta, phi) surfaces
res=100;
% ip components to use from ePS output (1=length gauge, 2=velocity gauge)
ipComponents=1;
% it components to use from ePS output (for degenerate cases), set an array here for
⇔as many components as required, e.g. it=1, it=[1 2] etc.
it=1;
% Set light polarization and axis rotations LF -> MF
                    % p=0 for linearly pol. light, +/-1 for L/R circ. pol.
                    % Eugler angles for rotation of LF->MF, set as [0 0 0] for z-pol,
eAngs=[0 0 0];
→[0 pi/2 0] for x-pol, [pi/2 pi/2 0] for y-pol
polLabel='z';
% Run calculation - outputs are D, full set of MFPADs (summed over symmetries); Xsect,
→ calculated X-sects; calcsAll, structure with results for all symmetries.
                                                                         (continues on next page)
```

5.3. MFPADs 61

(continued from previous page)

```
% Results are output as a structure, dims (symmetries, energies). calcsAll
```

```
calcsAll =

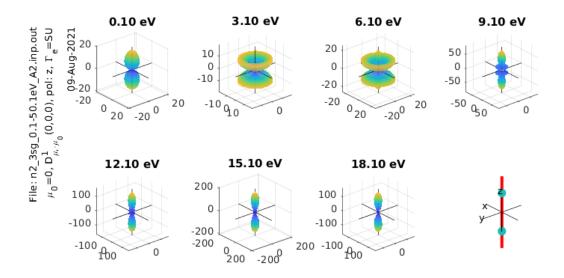
3x51 struct array with fields:

D
C
Cthres
eKE
symm
euler
Xsect
XsectD
Rlf
P
Cind
```

```
%plot -s 800,400
```

(Above is a line-magic for setting displayed plot size with the Calysto kernel - see the demo notebook for more.)

```
%% Plotting - MFPAD panel plots
% Set plot ranges
symmInd=1; % Select symmetry (by index into calcsAll rows). Final symmetry state_
→is set as sum over all symmetries
% eRange=1;
            % Select energies (by index into calcsAll cols)
eRange=1:3:20;
% Additional options (optional)
sPlotSet=[2 4];
                 % Set [rows cols] for subplot panels. The final panel.
will be replaced with a diagram of the geometry
% titlePrefix='NO2 testing'; % Set a title prefix for the figure
titlePrefix='';
ePSproc_MFPAD_plot(calcsAll, eRange, symmInd, params, sPlotSet, titlePrefix);
% ePSproc_MFPAD_plot(calcsAll, eRange, symmInd, params, sPlotSet, '', 'n', 'off');
% ePSproc_MFPAD_plot(calcsAll, eRange, symmInd, params, [2 4], '', 'n', 'off');
```



```
% Calculate & plot for a different polarization state
eAngs = [0 pi/2 0];  % x-pol case
polLabel = 'x';

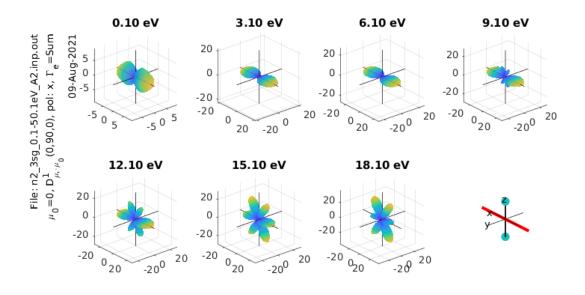
[Xsect, calcsAll, pWaves]=ePSproc_MFPAD(rlAll,p,eAngs,it,ipComponents,res);

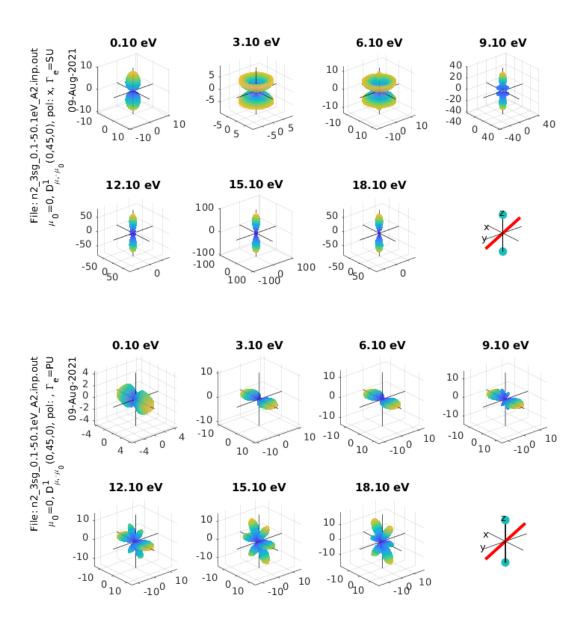
% Add pol labels - currently expected in plotting routine, but not set in MFPAD_
eroutine
for n=1:size(calcsAll,2)
    for symmIn=1:size(calcsAll,1)
        calcsAll(symmInd,n).polLabel=polLabel;
    end
end

symmInd=3;
ePSproc_MFPAD_plot(calcsAll,eRange,symmInd,params,sPlotSet,titlePrefix);
```

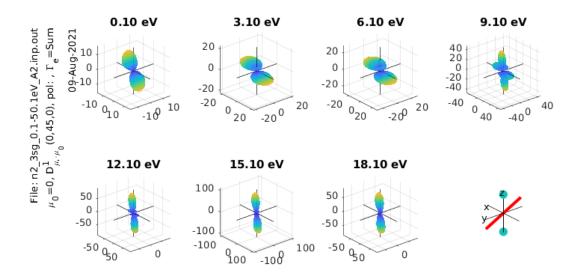
```
symmInd =
3
```

5.3. MFPADs 63





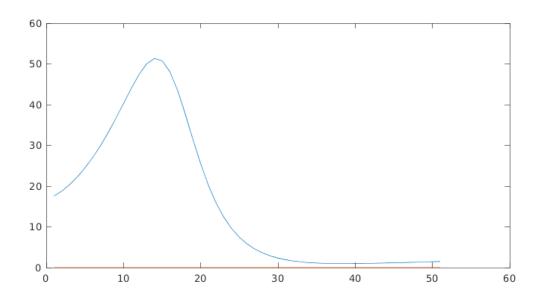
5.3. MFPADs 65



5.4 MF β_{LM}

```
%% *** Calculate MFPADs - single polarization geometry, all energies and symmetries
% Calculate for specified Euler angles (polarization geometry) & energies
% Set resolution for calculated I(theta, phi) surfaces
res=100;
% ip components to use from ePS output (1=length gauge, 2=velocity gauge)
ipComponents=1;
% it components to use from ePS output (for degenerate cases), set an array here for-
→as many components as required, e.g. it=1, it=[1 2] etc.
it=1;
% Set light polarization and axis rotations LF -> MF
                % p=0 for linearly pol. light, +/-1 for L/R circ. pol.
                  % Eugler angles for rotation of LF->MF, set as [0 0 0] for z-pol,_
eAngs=[0 0 0];
\rightarrow [0 pi/2 0] for x-pol, [pi/2 pi/2 0] for y-pol
polLabel='z';
% Run calculation - outputs are D, full set of MFPADs (summed over symmetries); Xsect,
- calculated X-sects; calcsAll, structure with results for all symmetries.
calcsAll=ePSproc_MFPAD(rlAll,p,eAngs,it,ipComponents,res);
% Add pol labels - currently expected in plotting routine, but not set in MFPAD.
⇔routine
% for n=1:size(calcsAll,2)
      for symmIn=1:size(calcsAll,1)
          calcsAll(symmInd,n).polLabel=polLabel;
양
응
      end
% end
```

plot(calcsAll.')



 $\overline{$ 5.4. MF β_{LM}

Quantum Metrology with Photoelectrons Vol. 3 *Analysis methodologies*	

BIBLIOGRAPHY

[Hoc18] Paul Hockett. *Quantum Metrology with Photoelectrons, Volume 1: Foundations.* IOP Publishing, 2018. ISBN 978-1-68174-684-5. URL: http://iopscience.iop.org/book/978-1-6817-4684-5, doi:10.1088/978-1-6817-4684-5.