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# **pypec Documentation**

***Release 0.1.1***

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**Nov 08, 2023**



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## GETTING STARTED

### 1.1 Description

**pypec** aims at fitting PEC data with an graphical interface.

In terminal enter the following command:

```
python -m pypec
```

A pdf version of the documentation can be found here [pypec](#). The source code can be viewed on [GitHub](#).

### 1.2 How to install

```
$ python setup.py install
```

or

```
$ pip install pypec
```

### 1.3 Dependencies

```
numpy>=1.17  
scipy>=1.5  
matplotlib>=3.0.0
```

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## 2.1 PhotoElectrochemistry

### 2.1.1 Basics

Photo-electrochemistry characterizations are used to study films at macroscopic, mesoscopic, and microscopic scales. The latter advances were used to support (photo-)electrochemical studies of the electronic and optical properties of passive films and oxidized metals, and of their interfaces with electrolytes, providing informations on the nature and structure of these materials and to use properties such as the oxidation behaviour of a metallic substrate.

Basically, two kinds of curves are recorded in the course of photoelectrochemical characterization experiments, photocurrent voltammograms and photocurrent energy spectra. In photocurrent voltammograms, photocurrents are measured as a function of the potential,  $V$ , applied to the semiconducting electrode, at a given photon energy,  $E = h\nu$ . In photocurrent energy spectra, photocurrents are recorded, at a given applied potential,  $V$ , as a function of the photon energy,  $E$ . The analysis of the shapes of photocurrent voltammograms may allow to obtain informations such as the semiconducting type of the material, the energy of the surface band levels, the presence of macroscopic defects inducing photogenerated electron-hole pairs recombinations.

However, despite attempts to refine the Gartner-Butler model by taking into account surface or volume recombination, a complete description of the photocurrent voltammograms remains difficult, for the latter developments make use of a high number of adjustable parameters, most of them being very difficult to assess. The analysis of the photocurrent energy spectra is intended to identify the chemical nature of the material constituting the semiconducting electrode, through the value of their bandgap energies,  $E_g$  as, on the one hand, bandgap energy values have been reported in the literature for numerous compounds, and as, on the other hand, bandgap values may be estimated from thermodynamic extensive atomic data. Practically, photocurrent energy spectra are usually analyzed by means of linear transforms to take benefit of the fact that, using the simplified form of the Gartner-Butler model, the quantum yield,  $\eta$ , of the photocurrent is proportional to the light absorption coefficient.

In such conditions,  $\eta$ , obeys to the following relationship:

$$(\eta * E)^{1/n} = K(E - E_g)$$

where  $C$  is a constant (things other than  $E$  being equal),  $E_g$  is the bandgap energy of the semiconductor, and  $n$  depends on the band to band transition type,  $n = 1/2$  for an allowed direct transition, and  $n = 2$  for an allowed indirect transition. Direct transitions are rarely observed in more or less disordered thin oxide films.

## 2.1.2 Fitting

Linear transformations were successfully performed for oxides made of one or two constituents. However, for complex oxide scales formed of several p-type and n-type phases, the complete description of the photocurrent energy spectra could not be achieved, and only semi-quantitative and/or partial informations could be obtained on the oxides present in the scales.

As  $I_{PH}^*$  is measured under modulated light conditions and thus actually is a complex number, the real and the imaginary parts of the photocurrent should be considered simultaneously when analyzing and fitting the photocurrent energy spectra, rather than their modulus [1].

$$\begin{aligned} I_{PH}^* &= |I_{PH}^*| \cos \theta + j |I_{PH}^*| \sin \theta \\ I_{PH}^* &= \sum_i^{i=N} J_{PH,i} \cos \theta_i + j \sum_i^{i=N} J_{PH,i} \sin \theta_i \end{aligned} \quad (2.1)$$

where  $J_{PH,i}$  and  $\theta_i$  represent the modulus and phase shift, respectively, of the photocurrent issued from the  $i$ th semiconducting constituent of the oxide layer. For thin semiconducting films, the space charge regions are low compared to penetration depth of the light.  $J_{PH,i}$  may thus be expected, at a given applied potential, to follow the simplified form of the Gartner–Butler model.

$$(J_{PH,i} * E)^{1/n} = K_i (E - E_{g,i})$$

where  $E_{g,i}$  and  $K_i$  represent the energy gap and a value proportional to  $C$  ( $I_{PH}^*$  is proportional to but not equal to  $\eta$ ) for the  $i$ th semiconducting constituent.

For a given vector of  $m$  ( $K_i, \theta_i, E_{g,i}$ ) triplets,  $m$  representing the supposed number of semiconducting phases contributing to the photocurrent, the scalar function to be minimized by the Nelder-Mead function was defined as the product of the square roots of two quantities:

$$\begin{aligned} D_{Re} &= \sqrt{\sum_E (Re I_{PH,exp}^* - Re I_{PH,calc}^*)^2} \\ D_{Im} &= \sqrt{\sum_E (Im I_{PH,exp}^* - Im I_{PH,calc}^*)^2} \\ D &= D_{Re} \cdot D_{Im} \end{aligned}$$

The 3  $m$  variables can be locked or not by the user. Initial estimates can be provided by the user or can be randomly generated. Several successive calls of the Nelder-Mead procedure are necessary to reach the minimum of the scalar function and a stable set of the output parameters. The user is free to set the number of successive calls of the Nelder-Mead procedure. Constraints on the 3  $m$  variables can be set by the user.

## 2.2 GUI

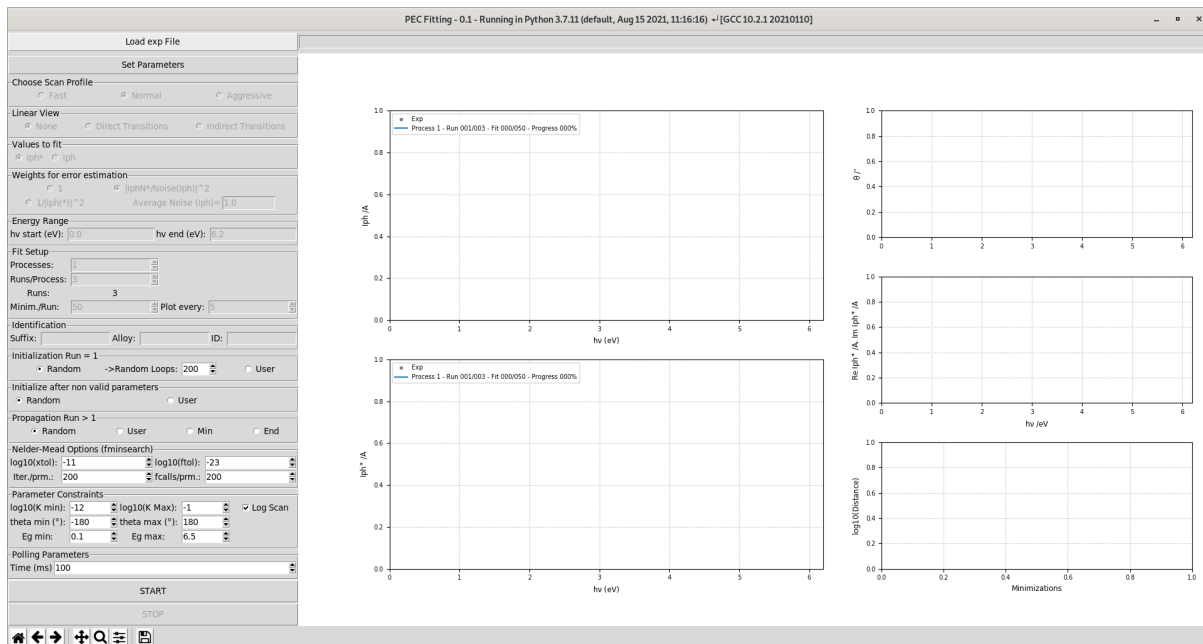
The main window contains all the elements necessary to run the fit. The design is cluttered but it allows exposing directly all the fitting settings to the user without deep menus.

The different steps, presented in the left pane, for performing the fit are:

- **load data: the accepted formats are:**
  - \*.dot files which are ascii files developed in the SiMaP Lab
  - \*.data files which are generic ascii files
- set the number of semiconductive contributions (*Parameter Table*)

At this point the fit can be ran.

If needed select custom choice for all the fitting parameter in the left pane.



## 2.2.1 \*.dot files

They have a specific formatting and they are provided by the PEC setup in the SiMaP Lab.

## 2.2.2 \*.data files

The \*.data files are generic ascii files where:

- the first column is the energy of the incident light in eV.
- the second column is the modulus of the photocurrent in A.
- the third column is the phase shift of the photocurrent in degrees.

## 2.2.3 Parameter Table

The parameter table allows for fitting the  $3m$  variables. The table is structured as shown below:

Ki	Fit Kgi	theta i	Fit Phase i	Egi	Fit Egi
K 1	0 or 1	Phase 1	0 or 1	Eg 1	0 or 1
K 2	0 or 1	Phase 2	0 or 1	Eg 2	0 or 1
...	...	...	...	...	...
K n	0 or 1	Phase n	0 or 1	Eg n	0 or 1

Each parameter  $K_i$ ,  $\theta_i$  and  $Eg_i$  can be locked by setting the **Fit X** column to 0.



## RELEASE NOTES

### 3.1 pypec 0.1 Release Notes

#### 3.1.1 Highlights

pypec aims at fitting PEC data with an graphical interface. The program was initially developped in Python 2.7 during my PhD from 2012 to 2015.

In 2016, the code was rewritten in Python 3 at the end of my PhD with slight modifications of the original code. The code remained untouched for almost 3 years and is now again under development in this repository.

This is the first release after migrating to github which will serve as a backup for the version developed during my PhD with completed documentation. No modification to the original code.

#### 3.1.2 New Features

- Load data: \*.dot files (data format developped in Grenoble Lab SiMaP)
- multi-process fitting: N processes x M fits x L loops
- automatic creation of result folder based on sample name and fitting parameters
- results saved in a folder

#### 3.1.3 Contributors

Milan Skocic

#### 3.1.4 Commits

N/A





## AUTOGENERATED DOCUMENTATION

### 4.1 Graphical FrontEnd

Graphical frontend for fitting photo-current spectra.

**class** `pypec.Analyse_PEC.Analyse_PEC`(*master=None*)

Construct a frame widget with the parent MASTER.

Valid resource names: background, bd, bg, borderwidth, class, colormap, container, cursor, height, highlightbackground, highlightcolor, highlightthickness, relief, takefocus, visual, width.

#### Methods

`update_nb_runs()`

Update the number of cpu and run per process on the graphical interface.

AddFiles_cb
Fit_cb
ask_quit
autoscale
create_fit_lines
get_progress
on_Run_Fit
on_hv_limits
on_start_workers
on_stop_button
plot_Graph
plot_Re_Im
plot_fit_lines
plot_ligne_V
prm_binary
process_queue
remove_fit_folder
remove_fit_lines
start
update_figure
update_legend
update_nb_fit_in_run

AddFiles\_cb()

Fit\_cb()

`ask_quit()`  
`autoscale(chart)`  
`create_fit_lines()`  
`get_progress(run=1, fit=0)`  
`on_Run_Fit()`  
`on_hv_limits(*args)`  
`on_start_workers()`  
`on_stop_button()`  
`plot_Graph()`  
`plot_Re_Im()`  
`plot_fit_lines(*args)`  
`plot_ligne_V()`  
`prm_binary()`  
`process_queue()`  
`remove_fit_folder()`  
`remove_fit_lines()`  
`start()`  
`update_figure()`  
`update_legend(axes, run=1, fit=0, location='upper left')`  
`update_nb_fit_in_run()`  
`update_nb_runs()`

Update the number of cpu and run per process on the graphical interface.

**class** `pypec.Analyse_PEC.ParameterTable(master, prm, last_prm_folder, **kwargs)`

Construct a frame widget with the parent MASTER.

Valid resource names: background, bd, bg, borderwidth, class, colormap, container, cursor, height, highlightbackground, highlightcolor, highlightthickness, relief, takefocus, visual, width.

**class** `pypec.Analyse_PEC.ParameterWindow(master, prm_init, last_prm_folder)`

Construct a toplevel widget with the parent MASTER.

Valid resource names: background, bd, bg, borderwidth, class, colormap, container, cursor, height, highlightbackground, highlightcolor, highlightthickness, menu, relief, screen, takefocus, use, visual, width.

## Methods

<b>get_paths</b>
<b>get_prm</b>

**get\_paths()**

**get\_prm()**

**class** pypec.Analyse\_PEC.ScrolledFrame(*master*, **\*\*kwargs**)

Construct a frame widget with the parent MASTER.

Valid resource names: background, bd, bg, borderwidth, class, colormap, container, cursor, height, highlightbackground, highlightcolor, highlightthickness, relief, takefocus, visual, width.

## 4.2 Iph Functions

Modules - Iph Functions

This module contains functions required for computing and optimizing the photo-current values from input parameters i.e. triplets  $(K_i, \theta_i, Eg_i)$  and experimental data for each semi-conductive phase [1].

**pypec.iph\_functions.get\_Iph\_calc**(*hv*, *prm\_array*, *phi\_N*)

Compute the complex values of *Iph* based on the values and states of the triplets  $(K_i, \theta_i, Eg_i)$  [1].

$$Iph^* = \frac{Iph}{\Phi_N}$$

### Parameters

**hv: 1d array**

Vector of energies for which the complex *Iph* has to be computed.

**prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

**phi\_N: 1d array**

Represents the values of the normalized photon flux to the maximum value. If *nphf* is a unity vector, the true photo-current is returned otherwise the as-measured photo-current is returned.

### Returns

**iph\_calc\_complex: 1d array**

Vector of the computed complex values of *Iph*.

**pypec.iph\_functions.get\_LCC**(*iph\_exp\_complex*, *iph\_calc\_complex*)

**pypec.iph\_functions.get\_distance**(*iph\_exp\_complex*, *iph\_calc\_complex*)

Compute the distance *D* between  $Iph_{exp}$  and  $Iph_{calc}$ . The distance is computed by multiplying the distances on real and imaginary parts of *Iph*:

$$\Delta Re = Re\ Iph_{exp} - Re\ Iph_{calc}$$

$$\Delta Im = Im\ Iph_{exp} - Im\ Iph_{calc}$$

$$D_{Re} = \sqrt{\sum \Delta Re^2}$$

$$D_{Im} = \sqrt{\sum \Delta Im^2}$$

$$D = D_{Re} \cdot D_{Im}$$

**Parameters**

**iph\_exp\_complex: 1d numpy array**  
Contains the complex values of the  $Iph_{exp}$ .

**iph\_calc\_complex: 1d numpy array**  
Contains the complex values of the  $Iph_{calc}$ .

**Returns**

**D: float**  
The computed distance on real and imaginary parts of  $Iph$ .

`pypec.iph_functions.get_exp_data(filepath)`

Get the data array of data files according to their extension.

Supported files are .dot files recorded by PECLab software and .data files were the first three columns represent  $h\nu$ ,  $|Iph^*|$ ,  $\theta$ .

**Parameters**

**filepath: string**  
Path to the data file.

**Returns**

**data\_array: 2d array**  
Experimental data.

`pypec.iph_functions.get_header_footer_dot_file(filepath)`

Find the number of lines in header and footer in .dot files.

**Parameters**

**filepath: path to the dot file**

**Returns**

**skip\_header: int**  
number of lines in header

**skip\_footer: int**  
number of lines in footer

**nbpoints: int**  
number of data lines

`pypec.iph_functions.get_random_prm_values(prm_array, K_bound=(1e-12, 0.1), theta_bound=(-180.0, 180.0), Eg_bound=(0.1, 6.2), phase_flag=True)`

Generates random values for the triplets  $(K_i, \theta_i, Eg_i)$  to be fitted based on the states given by the *prm\_array*.

By default, the limits are:

- $K_i$ :  $[10^{-12}, 10^{-1}]$
- $\theta_i$ :  $[-\pi, +\pi]$
- $Eg_i$ :  $[0.1, 6.0]$

**Parameters**

**prm\_array: 2d array**  
Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

**K\_bound: tuple**  
Contains the lower and upper limits for the  $K_i$  values.

**theta\_bound: tuple**  
Contains the lower and upper limits for the  $\theta_i$  values.

**Eg\_bound:tuple**

Contains the lower and upper limits for the  $Eg_i$  values.

**phase\_flag: bool**

Indicates if the values of  $\theta_i$  have to be randomized.

**Returns****random\_prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

`pypec.iph_functions.get_results_array(hv, iph_exp_complex, iph_calc_complex)`

Build the data array of the experimental and calculated data:  $h\nu$ ,  $|Iph_{exp}|$ ,  $\theta_{exp}$ ,  $|Iph_{calc}|$  and  $\theta_{calc}$

**Parameters****hv: 1d numpy array**

Contains the energy vector.

**iph\_exp\_complex: 1d array**

Contains the complex values of  $Iph_{exp}$ .

**iph\_calc\_complex: 1d array**

Contains the complex values of  $Iph_{calc}$ .

**Returns****data\_array: 2d array**

Array containing the .

`pypec.iph_functions.get_summary(fit_folder)`

List result files for the triplets  $(K_i, \theta_i, Eg_i)$  at the end and the minimum of each run.

Compute the distance, the LCCs for the energy interval that was used for minimizing the the triplets  $(K_i, \theta_i, Eg_i)$ .

The results are saved in 4 files: .SumEnd, .SumEndEg, \*.SumMin, \*.SumMinEg.

**Parameters****fit\_folder: string**

Path of the fit folder.

`pypec.iph_functions.import_prm_file(filepath)`

Import the triplets  $(K_i, \theta_i, Eg_i)$  from text file where each line represents a contributing semi-conductive phase.

**Parameters****filepath: string**

Absolute or relative file path to the text file.

**Returns****prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

`pypec.iph_functions.minimize(hv, iph_exp_complex, phi_N, weights, prm_array, Ki_log_flag=True, maxiter=None, maxfun=None, xtol=1e-11, ftol=1e-23, full_output=True, retall=False, disp=False, callback=None)`

Execute the Nelder-Mead algorithm based on parameter values given by `prm_array` and energy vector  $h\nu$ .

First, the `prm_array` is flattened and the parameters  $(K_i, \theta_i, Eg_i)$  to be fitted are extracted and sent to the `target_function()` through the Nelder-Mead algorithm.

Once the parameters were computed by the Nelder-Mead algorithm, the `prm_array` is updated with the new values.

**Parameters**

**hv: 1d numpy array**

Contains the energy vector.

**iph\_exp\_complex: 1d numpy array**

Contains the complex values of the experimental photo-current.

**phi\_N: 1d array**

Contains the normalized photon spectrum.

**weights: 1d array**

Contains the weights of the data.

**prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

**Ki\_log\_flag: bool**

Indicates if the  $K_i$  values are in logarithmic space.

**maxiter**

[int, optional] Maximum number of iterations to perform.

**maxfun**

[number, optional] Maximum number of function evaluations to make.

**xtol**

[float, optional] Relative error in xopt acceptable for convergence.

**ftol**

[number, optional] Relative error in func(xopt) acceptable for convergence.

**full\_output**

[bool, optional] Set to True if fopt and warnflag outputs are desired.

**retall**

[bool, optional] Set to True to return list of solutions at each iteration.

**disp**

[bool, optional] Set to True to print convergence messages.

**callback**

[callable, optional] Called after each iteration, as callback(xk), where xk is the current parameter vector.

## Returns

**prm\_array: 2d array**

Represents the updated values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

**fopt**

[float] Value of function at minimum:  $fopt = func(xopt)$ .

`pypec.iph_functions.plot_summary(fit_folder)`

Plot the result files that were created by the `get_summary()` for the triplets  $(K_i, \theta_i, Eg_i)$  at the end and the minimum of each run.

The results are saved in 2 files: -0-End.pdf, -0-Min.pdf.

## Parameters

**fit\_folder: string**

Path of the fit folder.

`pypec.iph_functions.save_pdf(filepath, hv, iph_exp_complex, iph_calc_complex, mask, all_results)`

`pypec.iph_functions.save_results(run, process_id, fit_folder, datafilepath, suffix, hv, mask, iph_exp_complex, phi_N, prm_min_run, prm_end_run, distance_min_run, distance_end_run, minimization_results, header_minimization_results)`

`pypec.iph_functions.scatter_logpolar(ax, theta, r_, ticks=5, bullseye=0.0, **kwargs)`

`pypec.iph_functions.shift_phase(prm_array, theta_bound=(-180.0, 180.0))`

Compute the modulo of  $\theta_i$  values with  $2\pi$  and then shift the values of  $\theta_i$  by the amplitude of the boundaries in order to be in between the boundaries.

By default, the boundaries for  $\theta_i$  are set to  $[-\pi, +\pi]$ .

#### Parameters

**prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

**theta\_bound: tuple**

Contains the lower and upper limits for the  $\theta_i$  values.

#### Returns

**prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$  where the  $\theta_i$  values were shifted.

`pypec.iph_functions.sort_prm_Eg(prm_array)`

Sort the `prm_array` based on values of  $Eg_i$ .

#### Parameters

**prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

#### Returns

**prm\_array: 2d array**

Represents the sorted values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

`pypec.iph_functions.target_func(p, hv, prm_array, iph_exp_complex, phi_N, weights, Ki_log_flag=True)`

Update the triplets  $(K_i, \theta_i, Eg_i)$  from the flattened parameter vector `p` sent by the optimization algorithm. The `prm_array` will be flattened and the indexes of the parameters to be fitted will be updated with `p` vector.

The calculated complex values of  $Iph$  will be sent along the experimental values to the `get_distance()` function. The value of the distance between the experimental and calculated data will sent back to the optimization algorithm.

#### Parameters

**p: 1d array**

Parameter vector sent by the optimization algorithm which is always flattened.

**hv: 1d array**

Vector of energies for which the complex values of  $Iph$  have to be calculated.

**prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

**iph\_exp\_complex: 1d numpy array**

Contains the complex values of the experimental  $Iph$ .

**phi\_N: 1d array**

Represents the values of the normalized photon flux to the maximum value.

**weights: 1d array**

Contains the values of the data weights.

**Ki\_log\_flag: bool**

Indicates if the  $K_i$  values are in logarithmic space.

#### Returns

**distance: float**

Calculated distance between experimental and calculated data values. See the [`get\_distance\(\)`](#) function.

`pypec.iph_functions.validate_prm(prm_array, K_bound=(1e-12, 0.1), Eg_bound=(0.1, 6.2))`

Check if the values of  $K_i$  and  $Eg_i$  are within the boundaries.

#### Parameters

**prm\_array: 2d array**

Represents the values and states of the triplets  $(K_i, \theta_i, Eg_i)$ .

**K\_bound: tuple**

Contains the lower and upper limits for the  $K_i$  values.

**Eg\_bound: tuple**

Contains the lower and upper limits for the  $Eg_i$  values.

#### Returns

**valid: bool**

Set to True if value of  $K_i$  or  $Eg_i$  is out of the boundaries.

## 4.3 Parallel Processes

Module for controlling the parallel processes running during the fitting procedure

```
class pypec.Parallel_Process.MinimizationProcess(output_queue, name, prm_init, nb_run, nb_SC,
                                                init_type, random_loops, hv, iph_exp_complex,
                                                iph_exp_complex_CI, phi_N, phi_N_CI,
                                                weights, hv_limits, nb_fit_in_run, fit_folder,
                                                filepath, suffix, NelderMead_options=(1e-11,
                                                1e-23, 200, 200),
                                                ParameterConstraints=((1e-12, 0.1), (-180,
                                                180), (0.1, 6.2), True), update_every=5)
```

#### Methods

`run()`

Method to be run in sub-process; can be overridden in sub-class

**shutdown**

`run()`

Method to be run in sub-process; can be overridden in sub-class

`shutdown()`

```
class pypec.Parallel_Process.PlotSummaryProcess(fit_folder)
```

#### Parameters

**fit\_folder: str**

Path to the folder where the summary files will be saved.

#### Attributes

**fit\_folder: str**

Path to the folder where the summary files will be saved.



## Methods

<code>run()</code>	Method to be run in sub-process; can be overridden in sub-class
--------------------	---

### `run()`

Method to be run in sub-process; can be overridden in sub-class

`pypec.Parallel_Process.get_cpu_number()`

DocString

`pypec.Parallel_Process.get_queue()`

DocString

`pypec.Parallel_Process.initialize_processes(n, daemon=True, **kwargs)`

Create N processes as daemons and return them in a list.

#### Parameters

**n: int**

Number of processes to be created.

**daemon: bool**

Flag for creating daemon process.

**kwargs: dict**

See [MinimizationProcess](#)

`pypec.Parallel_Process.start_processes(workers)`

#### Parameters

**workers: subprocess workers**

#### Returns

**0: return integer**



## BIBLIOGRAPHY

### 5.1 Bibliography



## INDICES AND TABLES

- genindex
- modindex
- search



## BIBLIOGRAPHY

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