
PIGOR Documentation

Nico Einsidler

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Pigor is a lightweight analysis tool for the the polarimeter instrument NEPTUN beam port of the 250kW research reactor hosted at [Atominstitut](#) of [TU Wien](#), Austria. For more information visit [our homepage](#).

HOW TO INSTALL PIGOR

1.1 Installing PIGOR

Assuming you have a running installation of **Python 3.7** or higher, to install PIGOR please follow these steps:

1. **Download the source from Bitbucket with `git clone https://github.com/nicoeinsidler/pigor.git`. This**

- `measurement.py`
- `pigor.py`
- `README.md`
- `requirements.py`
- `requirements-dev.py`
- `requirements-measurement.py`

2. Install the requirements with:

```
pip install -r requirements.txt
```

Note: If you are using two separate python 2.x and 3.x installations, you might need to use `pip3` instead of `pip`.

You can now run PIGOR with the following command in the folder where the `pigor.py` and `measurement.py` files are located:

```
python pigor.py
```

Note: If you only want to use the `measurement.Measurement` class, you can only install it's dependencies with `pip install -r requirements-measurement.txt`. But since most dependencies are shared, it won't make much of a difference.

1.2 Installing only the Measurement Class

Warning: This is not maintained yet. Please use this feature with caution and build it first.

In some cases the user may only want to install the measurement module to use the Measurement class as one of their python installations modules. A zip or tarball file should be provided for this use case. See deployment-measurement-module.

To install (and copy into the user's installed modules) simply unpack the zip or tarball. Head over to its directory and type from there:

```
python setup.py install
```

1.3 Installation for Developers

A known working set of python modules used to create PIGOR can be found in `requirements-dev.txt`. To install all these requirements, simply type:

```
pip install -r requirements-dev.txt
```

It is recommended to use a virtual environment.

MEASUREMENT NAMING CONVENTION

In order for the Measurement class to work properly, the user should follow this naming convention:

YYYY_MM_DD_HHmm_IDENTIFIER_TYPE.dat (not case sensitive)

where...

- **IDENTIFIER** denotes the type of measurement or the reason for the measurement. The following values are common and
 - dcX#: DC coil (for the magnetic field in x-direction) scan for the DC coil with the number #. A sine fit is automatically applied, when TYPE is not set.
 - dcZ#: DC coil scan in z-direction (compensation field) for DC coil number #. Fitted with a polynomial of order 5 by default.
 - pos#: position scan with stage number #. Fitted with a Gaussian by default.
- **TYPE** specifies and overrides the type of fitting/analysis that should be applied by PIGOR. Possible values:
 - **fitting types**
 - * poly5: polynom of order 5
 - * sine_lin: sine plus additional added linear function
 - * sine: sine
 - * gauss: Gauß fit
 - **special analysis types**
 - * pol: specifies that this measurement is a degree of polarisation measurement. This will automatically calculate the degree of polarisation for any given number of points (number of data points must be multiple of four)

Types can be combined within a types group (gauss and pol for example, like “gauss_pol” or “pol_gauss”).

QUICKSTART GUIDE

Note: Please refer to the [Installing PIGOR](#) page for more information on the installation.

Suppose you have some measurement data files with in a directory. The hierachy looks like this:

- **measurements**
 - **2019-01**
 - * 2019-01-01-dc1x.dat
 - * 2019-01-04-dc1z.dat
 - * 2019-01-05-dc1z.dat
 - **2019-02**
 - * 2019-02-10-dc2x.dat
 - * 2019-02-11-dc2x.dat
 - * 2019-02-14-dc2z.dat
 - * 2019-01-19-dc2z.dat
 - * 2019-01-20-dc2z.dat
 - * 2019-01-28-dc1x.dat
 - **2019-03**
 - * 2019-02-02-dc3x.dat
 - * 2019-02-03-dc3z.dat

Note: Here we will assume that `python` will start Python 3. On some installations it must be explicitly started with `python3`.

In order to analyse these files with PIGOR, head to the directory where PIGOR is located. Start it by executing the command `python pigor.py`. If it is the first startup it will go through some questions regarding your analysis setup. The most important question is the first one: Where should PIGOR look for files to analyse? This is the folder where all your measurements are located in and will be referred as PIGOR's root folder. In our example type in the path to `measurements/`. You can either use absolute paths or relative ones. PIGOR will automatically check if the path actually exists, if not you will be prompted to enter a new one again. It also does not matter if you are using a POSIX type (IEEE Std 1003.1-1988) or as a DOS type path. (POSIX uses `/` and DOS uses `\` as separators)

If PIGOR does not ask these question on startup, they have already been configured. Either delete the `pigor.config` file in the directory where `pigor.py` is located, or just use the command `[i]` within PIGOR to reconfigure it.

Now you should see something like that, depending on PIGOR's version:

```
$ python pigor.py

=====
Welcome to PIGOR v1.1.
=====

i ... init
h ... print_help
a ... analyse_files
j ... create_index
r ... remove_generated_files
x ... print_root
q ... quit PIGOR v1.1

PIGOR v1.1 will look for measurement files in D:\measurements.

If you need more information about a command, just type h + [command] + <ENTER>
to get more help. For example: h + a + <ENTER>.

Please type a command you want to perform and press <ENTER>.
```

On startup PIGOR is printing the root, which in this case is `D:\measurements`. It will also include a list of available commands. Each command can be triggered by pressing the corresponding letter followed by an <ENTER>. Some commands allow more options, each separated by a space.

We will describe these commands by `[cmd]` where `cmd` stands for the specific command.

To get more help on a specific command, just type `[h] + [cmd]`. For example typing `h a` will give us the help on `analyse_files()`. Let's try it:

```
Please type a command you want to perform and press <ENTER>.
h a
a:

Analyses all given files in list. This function can be used by the command [a].

    :param filepaths:    list of files to analyse with
                        their relative dir path added

    .. todo:: Change to no override mode. measurement.Measurement.plot(override=False)
    .. todo:: a + today => only analyse files for today
    .. todo:: a + override => override=True
```

This is the same help text as found in this documentation.

We can now analyse our files with `[a]`. You can see on which file PIGOR is currently working on. If an error occurs, you will see it as well and PIGOR will skip the file.

After PIGOR is done analysing files, you may want to access these files. Use the `[j]` command to create an index to quickly go through all the files that have been analysed. Now you should see a new file has been created:

- **measurements**
 - `index_pigor.html`

- 2019-01
- 2019-02
- 2019-03

Open `index_pigor.html` to see the list. From there you can review the original data and the files that PIGOR created from this.

4.1 What PIGOR does

PIGOR ('Python IGOR' = PIGOR) aims to help physicists on the NEPTUN beamline to quickly extract the needed information when measuring and configuring or preparing an experiment. It will go through all files in its root folder and will continue to **look for files in all subdirectories recursively** as well. It will then **auto detect¹ the type of measurement** and guess what the user wants to know. After analysis of all files, **additional files will show up alongside the original measurement files**:

- .png file: plot of the data
- .md file: usefull information gathered about the measurement in plain text as markdown
- .html file: same content as the markdown file, but nicely viewable in a modern browser

4.2 PIGORs inner workings

`pigor.analyse_files (filepaths='all')`

Analyses all given files in list. This function can be used by the command [a].

Parameters `filepaths` – list of files to analyse with their relative dir path added

Todo: Change to no override mode. `measurement.Measurement.plot(override=False)`

Todo: a + today => only analyse files for today

Todo: a + override => `override=True`

`pigor.bool2yn (b)`

Converts a boolean to yes or no with the mapping: y = True, n = False.

`pigor.create_index ()`

Creates an index.html listing all directories and subdirectories and their HTML and Markdown files. If a default browser is found, it will automatically open index.html. This function can be used by the command [j].

¹ This will only work if the correct naming convention is used.

`pigor.find_all_files()`

Finds all dat files recursively in all subdirectories ignoring hidden directories and Python specific ones.

Returns a list of filepaths.

`pigor.init(create_new_config_file=True)`

This function will read the config file and initialize some variables accordingly. This function can be used by the command [i].

Available options:

- root directory where PIGOR will start to look for measurement files
- Should PIGOR look for files to analyse recursively?
- Which file extension do the measurement files possess?
- What plot output format should PIGOR use?
- Should PIGOR automatically create an html file?
- Should PIGOR automatically create a md file?
- Should PIGOR create a txt file containing all used fit functions for the use in Mathematica?

Note: If no config file can be found, it will create one.

`pigor.is_valid_theme(theme)`

Checks if this theme exists.

`pigor.list_themes()`

Returns a list of all themes available.

`pigor.main()`

Main Loop

`pigor.print_header(text)`

This function prints a beautiful header followed by one empty line.

Parameters `text` – text to be displayed as header

`pigor.print_help(display='all')`

Prints a help menu on the screen for the user. This function can be used by the command [h].

Parameters `display` – specify the length of the help menu, options are 'all' or 'quick' (Default value = "all")

`pigor.print_root()`

Prints the root for PIGOR, e.g. where it will look for files to analyse. This function can be used by the command [x].

`pigor.remove_generated_files(files='all')`

Removes the generated png, html and md files. This function can be used by the command [r].

Parameters `files` – list of Path objects to files that should be removed; if set to 'all' it will delete all generated files (Default value = 'all')

Todo: Cover the case when files are not a list of path, e.g. wrong input given.

`pigor.show_user(func)`

Register a function to be displayed to the user as an option

`pigor.yn2bool(s)`

Converts yes and no to True and False.

USING PIGOR IN JUPYTER

Note: Please make sure that you have a running Jupyter or Jupyter Lab installation on your system. For more information, please visit <https://jupyter.org/install>.

Note: Make sure to install all the requirements for the measurement class in *requirements-measurement.txt*. See *Installing PIGOR* for more details.

First, open a new Jupyter notebook. Make sure that these following files are within the directory where your newly created Jupyter notebook is located:

- measurement.py
- fit_functions.py

If not, you can just copy those two files.

Now we have to import the measurement module in the Jupyter notebook by using the *import* command:

```
import measurement
```

This should already be enough to work with the measurement class within a Jupyter notebook. However there are a few extra steps to take, if one wishes to use inline graphics. To enable this functionality one line has to be added as the first cell to the notebook:

```
%matplotlib inline
```

Now, whenever the plot method is called, an additional argument has to be passed on.

```
m.plot(enable_jupyter=True)
```

Note: A test notebook is provided in the root of the PIGOR project at <https://github.com/nicoeinsidler/pigor/blob/master/PIGOR-Test-Notebook.ipynb>.

THE MEASUREMENT CLASS

6.1 Flow at Startup

1. read the data
2. detect measurement type
3. read position file if a position file exists
4. clean up and description gathering
5. select columns for plotting

6.2 Flow when plotting

6.3 Class Usecases

There are many ways to interact with or use the Measurement class. Here are the three main ways:

6.4 Methods

Todo: Method attributes are shown, but value is always None.

class `measurement.Measurement` (*path*, *type_of_measurement*='default', *type_of_fit*='gauss')

This class provides an easy way to read, analyse and plot data from text files.

There are two different file formats, which are used on the interferometry as well as on the polarimeter station at Atominstitut of TU Wien. For more information on the conventions please head to the docs or take a look at the example files provided.

FIT_RESOLUTION = None

number of points to calculate the fit for

N_HEADER = None

number of lines of header of measurement file

__init__ (*path*, *type_of_measurement*='default', *type_of_fit*='gauss')

The Measurement class provides an easy and quick way to read, analyse and plot data from text files. When creating a new instance, the following parameters have to be provided:

param self the object itself

param path pathlib.Path object

param type_of_measurement used to hard set the type of measurement on instance creation (default value = 'default')

param type_of_fit sets an initial fit type, which may be overridden by `detect_measurement_type()` later (default value = 'gauss') TODO: change to be permanent?

The startup sequence is as follows:

1. try to read the data
2. measurement type (either set it, when given as input argument or try to detect it)
3. if measurement is POL, try to find a position file and read it
4. clean up the given data
5. select columns => write into `self.x` and `self.y`
6. if measurement is POL, calculate degree of polarisation

Todo: Is `type_of_fit` really needed?

Returns nothing.

clean_data()

Splits the `raw` data into `head` and `data` vars.

contrast (*source='fit'*)

Calculates the contrast of source as:

$$\text{contrast} = (\text{max} - \text{min}) / (\text{max} + \text{min})$$

where `min` and `max` are the minima and maxima of the given data.

Parameters **source** – defines the source of the data to calculate the contrast from, can be either set to 'fit' or 'data' (Default value = 'fit')

Returns a list of contrasts.

Todo: When calculation of contrast fails, what should this function return? Now it returns `[0]`.

degree_of_polarisation()

Calculates the degree of polarisation for each position in `pos_data`.

detect_measurement_type()

This function auto detects the type of measurement based on the file name. This works best with a meaningful file name convention. For more information please refer to the docs.

Several `type_of_measurement` can be detected:

- DC#X: x-field of DC coil number # scan -> sets `type_of_fit` = 'sine_lin'
- DC#Z: z-field of DC coil number # scan -> sets `type_of_fit` = 'poly5'
- POS: scan of different linear stage positions -> sets `type_of_fit` = 'gauss'

`type_of_fit` can be overridden by explicitly mentioning a fit function to use in the name of the file. See docs for more information.

In addition to the type of fit and measurement type, some additional information about the measurement is gathered in the `settings` dict.

export_meta (*make_md=True, make_html=False, theme='github'*)

Exports all available information about the measurement into a markdown file.

Parameters

- **html** – if set to True, an HTML file will be additionally created (Default value = False)
- **theme** – set the default theme for html export, all available themes can be found in the `markdown_themes` directory (Default value = 'github')

find_bounds (*fit_function=None*)

Automatically finds usefull fit bounds and updates them in the `fit_function_list` dict.

Parameters **fit_function** – defines for which fit functions the bounds should be updated (Default value = None), if set to None, `type_of_fit` will be used

fit (*fit_function=None, fit_function_export=False*)

Fits the data in `x` and `y` using the default fit function of each `type_of_fit` if not specified further by passing a certain fit function as an argument.

Parameters

- **fit_function** – fit function to use to fit the data with (Default value = None)
- **fit_function_export** – exports the fit function as a txt file in a specified format (Mathematica is default and only implementation yet.).

Stores the optimal values and the covariances in `popt` and `pcov` for later use.

fit_function_list = None

list of fit functions that can be used; imported from `fit_functions.py`

measurement_type (*type_of_measurement='default'*)

Sets the type of the measurement if parameter `type_of_measurement` is set.

Parameters

- **self** – object itself
- **type_of_measurement** – default”: new type of measurement (default value = 'default')

Returns the current type of measurement.

Todo: Evaluate if this method (`measurement_type()`) is needed at all.

Todo: Set better default value for measurement type.

path = None

path (`pathlib.Path` object) to the measurement file

plot (*column1=(0, 1), column2=(1, 1), fit=True, type_of_plot="", override=True, file_extention='.png', enable_jupyter=False*)

Creates a plot for the data. If `fit` is set to False the data fit won't be plotted, even if there exists one. Following parameters are possible:

Parameters

- **self** – the object itself
- **column1** – (column, nth element) to choose the data from for x-axis (Default value = (0))
- **column2** – (column, nth element) to choose the data from for y-axis (Default value = (1))
- **fit** – if set to False plotting of the fit will be suppressed (Default value = True)
- **type_of_plot** – string to specify a certain plot type, which will be used in the file name as well as in the plot title (Default value = '')
- **override** – determines if plot image should be recreated if it already exists (Default value = True)
- **enable_jupyter** – if set to True, triggers `plt.show()` so the Measurement class can be used in a Jupyter Notebook

Todo: Make x and y labels more general, especially for interferometer files, where more than one y value list is needed.

pos_file_path = None

path (`pathlib.Path` object) to the corresponding position file

read_data (*path*)

Reads data from file and stores it in `raw`.

Parameters

- **self** – the object itself
- **path** – a `pathlib.Path` object pointing to a measurement file

read_pos_file ()

Looks for a position file and reads it into `pos_data`.

Todo: When searching for a position file, the length of the file should match. So it should be 1/4 of the size of the original measurement file.

reset_bounds (*fit_function=None*)

Resets the bounds of the measurement type's default fitting function if not specified otherwise.

Reset values are `(-np.inf, np.inf)`.

Parameters **fit_function** – specifies the fit function for which the bounds should be reset (Default value = None)

select_columns (*m=None*)

Selects columns of the `data` as specified in `m` (map) and saves in `x` and `y[]`.

..note:: `y_error[]` is calculated as `sqrt(y)`

Parameters **m** – map, e.g. list of tuples or None values; if `m=None` `select_columns` will be skipped (Default value = None)

The map `m` defines which columns of the original measurement data will be used later. Only one x-axis can be defined, but multiple y-axes may be used. The length of the map must not exceed the number of the columns in `data`, but can be less or equal.

Each map is a list of items, which can either be tuples or None values, if a column should be skipped. In the case of a tuple, the first value must be a string, either 'x' or 'y', which determines if the column should be interpreted as an x- or y-axis. Its second value describes what nth element of the columns should be selected.

A few examples:

```
m = [ ('x', 1), ('y', 1) ]
```

This will select the first column as x-axis and take every (1st) element of it, and the second column as y-axis, also using every element of that column.

```
m = [ ('y', 2), None, ('x', 2), ('y', 2) ]
```

Here we will take every second element of column 1, 3 and 4, but skip column 2.

Note: If the length of the map is less than the number of columns in `data`, every column that has no corresponding map element will be skipped.

settings = None

dict containing useful information read from files header in `clean_data()`

type_of_fit = None

type of fit to be applied to the data

MEASUREMENT CLASS 2.0

Due to some design flaws in the existing `measurement.Measurement` class, it will be rewritten from ground up using already existing libraries. This undertaking will lead to a cleaner version and will make it possible to adapt `measurement.Measurement` more easily for the interferometer experiments.

7.1 Structure

The new improved structure will make use of:

- LMFIT package
- pandas dataframe

So that the fitting will be done with LMFIT models, whereas the data is handled in pandas dataframes. This creates huge advantages for the developer as well as for the user.

7.2 Previous Development

There have been efforts previously to build a unique own modular `measurement.Measurement` class by Nico. These efforts can be examined in the branch `improvements/core`.

These following elements were attempted to build:

- data column: has data and a head, knows its name etc.; functions can easily be applied to it
- fit object: used for fitting and finding bounds; each instance can have its own bounds
- **data set: these objects can be plotted by Measurement, so Measurement will try to create one of those objects; they consist of**
 - data columns objects
 - fit objects

Column Class

Methods:

- `reverse()`: reverse order of data
- `__init__(self, desc, data)`
- `__repr__()`: plots '<column object 'desc' of length len(data)>' or something like that

Variables:

- `columns.data`: holds the data as numpy array in float64

- `columns.desc`: holds the name of the columns heading as string

Fit Function Class

Method:

- `fit_function()`
- `find_bounds()`: tries to find the bounds
- `bounds`: holds the bounds to be used when fitting as array of tuples

Variables:

- `parameters`: dictionary holding the names of the parameters and the parameters themselves

Fit Class

Variables:

- `type` with which function the fit should be carried out, string
- `popt`
- `pcov`

Methods:

- `fit()`

Data Set Class

This object stores data points (lists or Column objects) to form a data set. It must contain at least two data point lists. These lists must have the same number of elements, if not the lists that don't have enough elements will get padded with 0.

Variables:

- `desc`: a description of what the data set describes (optional)
- `data`: data is stored in list; `len(list) > 1`;

It is obvious to every reader that indeed almost all of this functionality is already included in the python package `pandas` and `lmfit`. This led to the conclusion that improvements/core won't be maintained any longer.

7.3 The brand new Measurement Class

The brand new measurement class will include only the following features:

- ready-to-use `lmfit` fit models
- `fit()` method to actually produce a fit using `lmfits minimize()`
- `plot()` method for implementing plotting functionality
- `contrast()` to calculate the contrast
- `degree_of_polarisation()` to calculate the degree of polarisation
- `read()` to read data in a very generic way
- `export_meta()` to export meta data

Its child classes will then implement the experiment specific data handling details like cleaning the data and extending the meta data that will be exported.

FIT FUNCTIONS

The following fit functions are implemented and can be used within PIGOR or the Measurement class. They can be found in `fit_functions.py`.

`fit_functions.gauss(x, a, x0, sigma, export=False)`
Gaussian function, used for fitting data.

Parameters

- **x** – parameter
- **a** – amplitude
- **x0** – maximum
- **sigma** – width
- **export** – enable text output of function

`fit_functions.poly(x, *args, export=False)`
Polynom nth degree for fitting.

Parameters

- **x** (*int, float*) – parameter
- ***args** – list of coefficients [*a_N, a_{N-1}, ..., a₁, a₀*]
- **export** (*bool or string, optional*) – enable text output of function, defaults to False

Returns returns the polynomial

Return type str, int, float

```
>>> poly(3.4543, 5, 4, 3, 2, 1, export='Mathematica')
'5*3.4543^5 + 4*3.4543^4 + 3*3.4543^3 + 2*3.4543^2 + 1*3.4543^1'
```

```
>>> poly(3.4543, 5, 4, 3, 2, 1)
920.4602110784704
```

`fit_functions.poly5(x, a5, a4, a3, a2, a1, a0, export=False)`
Polynom 5th degree for fitting.

Parameters

- **x** – parameter
- **a5** – coeff
- **a4** – coeff

- **a3** – coeff
- **a2** – coeff
- **a1** – coeff
- **a0** – coeff
- **export** – enable text output of function

Returns function – polynomial 5th degree

`fit_functions.register_fit_function(func, bounds=(-inf, inf))`

This decorator registers a new fit function and writes an entry to `fit_function_list`.

`fit_functions.sine(x, a, omega, phase, c, export=False)`

Sine function for fitting data.

Parameters

- **x** – parameter
- **a** – amplitude
- **omega** – frequency
- **phase** – phase
- **c** – offset
- **export** – enable text output of function

`fit_functions.sine_lin(x, a, omega, phase, c, b, export=False)`

Sine function with linear term added for fitting data.

Parameters

- **x** – parameter
- **a** – amplitude
- **omega** – frequency
- **phase** – phase
- **c** – offset
- **b** – slope
- **export** – enable text output of function

SPRINT PLANNING

This site gives a quick overview what will come next. Each sprint should take about 1 week to finish.

9.1 PIGOR

1. Sprint

- ✓ feature: remove all generated files (html, md, png)
- ✓ feature: introducing a config file (PIGOR start directory, ...)
- ✓ improvement: auto create config file if not present
- ✓ improvement: auto register all functions for help menu (decorators)

2. Sprint

- feature: remove last generated files (html, md, png)
- feature: remove all html/md or png files
- ✓ improvement: use JSON for config file

3. Sprint:

- feature: auto run command in specified intervals; syntax maybe: time + [cmd] + <ENTER>

9.2 Measurement Class

1. Sprint

- ✓ improvement: switching from `self.y` → `self.y[]` and `self.y_error` → `self.y_error[]`
- ✓ [not tested yet] improvement: plot multiple `self.y`'s
- feature: auto detect interferometer measurements

2. Sprint

- feature: remove all associated files from file system, except the measurement file itself
- ✓ improvement: auto register all available fit functions via decorators
- improvement: adding `__repr__`

3. Sprint: finish branch **feature/interferometer**

- fixing / understanding inheritance of instance variables (see python test file in branch)

- **creating subclasses from Measurement:**

- Interferometer: adding custom maps to COLUMN_MAPS and overriding `clean_data()` and `detect_measurement()`
- Polarimeter: adding custom maps to COLUMN_MAPS and overriding `clean_data()` and `detect_measurement()`

4. Sprint: not planned yet

9.3 Ideas

Building Measurement from ground up with custom objects like:

- data column: has data and a head, knows its name etc.; functions can easily be applied to it
- fit object: used for fitting and finding bounds; each instance can have its own bounds
- **data set: these objects can be plotted by Measurement, so Measurement will try to create one of those objects; they consist**
 - data columns objects
 - fit objects

9.4 Column Class

Methods:

- `reverse()`: reverse order of data
- `__init__(self, desc, data)`
- `__repr__()`: plots '<column object 'desc' of lenght len(data)>' or something like that

Variables:

- `columns.data`: holds the data as numpy array in float64
- `columns.desc`: holds the name of the columns heading as string

9.5 Fit Function Class

Method:

- `fit_function()`
- `find_bounds()`: tries to find the bounds
- `bounds`: holds the bounds to be used when fitting as array of tuples

Variables:

- `parameters`: dictionary holding the names of the parameters and the parameters themselves

9.6 Fit Class

Variables:

- `type` with which function the fit should be carried out, string
- `popt`
- `pcov`

Methods:

- `fit()`

9.7 Data Set Class

This object stores data points (lists or Column objects) to form a data set. It must contain at least two data point lists. These lists must have the same number of elements, if not the lists that don't have enough elements will get padded with 0.

Variables:

- `desc`: a description of what the data set describes (optional)
- `data`: data is stored in list; `len(list) > 1`;

TODO LIST

Todo: check if dependencies are correct with dependencies.txt

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/doc/index.rst, line 56.)

Todo: Method attributes are shown, but value is always None.

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/doc/measurement.rst, line 62.)

Todo: Is type_of_fit really needed?

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/measurement.py:docstring of measurement.Measurement.__init__, line 23.)

Todo: When calculation of contrast fails, what should this function return? Now it returns [0].

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/measurement.py:docstring of measurement.Measurement.contrast, line 12.)

Todo: Evaluate if this method (measurement_type()) is needed at all.

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/measurement.py:docstring of measurement.Measurement.measurement_type, line 9.)

Todo: Set better default value for measurement type.

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/measurement.py:docstring of measurement.Measurement.measurement_type, line 10.)

Todo: Make x and y labels more general, especially for interferometer files, where more than one y value list is needed.

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/measurement.py:docstring of measurement.Measurement.plot, line 13.)

Todo: When searching for a position file, the length of the file should match. So it should be 1/4 of the size of the original measurement file.

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/measurement.py:docstring of measurement.Measurement.read_pos_file, line 3.)

Todo: Change to no override mode. `measurement.Measurement.plot(override=False)`

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/pigor.py:docstring of pigor.analyse_files, line 6.)

Todo: `a + today => only analyse files for today`

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/pigor.py:docstring of pigor.analyse_files, line 7.)

Todo: `a + override => override=True`

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/pigor.py:docstring of pigor.analyse_files, line 8.)

Todo: Cover the case when files are not a list of path, e.g. wrong input given.

(The [original entry](#) is located in /Users/nicoeinsidler/Code/pigor/pigor.py:docstring of pigor.remove_generated_files, line 5.)

OLD TODOS

Note: This list of Todos should be integrated into the docstrings or into the sprint planning where those Todos belong.

Here are all Todos listed. Feel free to contribute and check this project out on Bitbucket.

- self.x_error: not yet implemented
- a lot of commenting
- error of fit
- verbose mode on/off
- getting PIGOR ready for shipment by creating a setup.py
- better display of self.pcov
- separation between pure functions and functions with context (methods)
- auto comment decorator for functions
- use decorators to auto register fit functions with their input argument list
- setuptools in setup.py
- <https://click.palletsprojects.com/en/7.x/quickstart/>

PROJECT DEPENDENCIES

Todo: check if dependencies are correct with dependencies.txt

- numpy
- re
- matplotlib
- os.path
- glob
- difflib
- datetime
- pathlib
- scipy
- markdown

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