**FLAP User’s Guide**

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# Introduction

The Fusion Library of Analysis Programs (FLAP) is a Python framework to work with large multi-dimensional data sets especially for turbulence data evaluation in fusion experiments. Data are stored in data objects together with coordinates, data names and coordinates, thus the built-in plotting functions create figures with correct axes. The data set can be sliced to reduce dimensions and thus enable visualization of more than 2D data sets. FLAP is a modular package: data are read through modules which register themselves to the FLAP framework. This way data are read through a uniform interface by defining data source, experiment ID and data name. Also coordinate conversion is done by functions in the data source modules.

# Configuration

FLAP can be adapted to the local environment using a configuration file. When flap is imported the default configuration file “flap\_defaults.cfg” is loaded from the working directory. If the file is not found a warning is printed. A configuration file can also be read explicitly using the flap.config.read() function.

The configuration file is a windows-style configuration file consisting of sections and elements. Sections start with their name in []. The elements follow on individual lines. The name of the element and the value is separated by =. Space can be used in both the section names and element names. Lowercase and uppercase characters are different. Usually names start with upper case but this is not a requirement. An example:

**[PS]**

**Resolution = 1e3**

**Range = [1e3, 1e6]**

**[Module TESTDATA]**

**PS/Resolution = 100**

**Name = ‘This is a string’**

In the above example section “PS” contains two elements. Section “Module TESTDATA” refers to the TESTDATA data source module. The element PS/Resolution refers to the resolution element in the PS section and enables overriding section settings with module specific values.

All elements in a section can be read with the flap.config.get\_all\_section() function. This returns a dictionary with keys referring to element names. The values are converted using the following rules:

* True and Yes is converted to boolean True.
* False and No is converted to boolean False.
* An element enclosed in single or double quotes is handled as string (without the quotes).
* Elements which can be converted to int, float or complex are interpreted accordingly.
* An element enclosed in square brackets ([]) is interpreted as a list. List elements should be separated by commas. Each list element is interpreted using the same rules as one element of the section.
* If all the above interpretation attempt fails the element is handled as string.

# Options and defaults

Standard flap functions (like flap.get\_data(), flap.apsd, ...) take an “options” keyword argument. A default options list is defined inside the routine, ehich contains all possible option keys understood by the function and default values for them. If no options are passed to the function these will take effect. The function might also be linked to a section in the configuration file. Options read from this section override the default options. The data object processed by the function might contain a data\_source variable. If that is present and not None configuration file elements in the section of that module are read. (That is in the above example for TESTDATA data\_source Module TESTDATA section is read.) elements starting with the section name of the function followed by / override settings in the section. Finally values in the function input options dictionary override all the above. As all possible options of the function are known from the default options dictionary it is allowed to abbreviate the option names in the function input options list up to the point where it matches only one key. (In the configuration file full option keys should be used.) This also means that any key in the default key list cannot be an abbreviation of another one. (E.g. ‘A’ and ‘A1’ are not allowed.)

This procedure is handled by the flap.config.merge\_options() function.

# Coordinates in FLAP

In the FLAP program package coordinates are stored with the data. This document describes the implementation of this feature.

## Data storage and coordinates

Data are stored in an n-dimensional numpy array in the FLAP.DataObject class variable. This n-dimensional space we call *data sample space*. Different dimensions of the array are associated with primary coordinates, like sample number, channel number, or e.g. for simulated data x, y. However, these primary coordinates are often not useful and we need to make plots along physical coordinates. This can be handled by adding other coordinates to the DataObject. Also during processing some coordinates might be turned to others. An example is calculating power spectra. From a 2D measurement data with channel, time coordinates spectrum calculation creates another 2D array with channel, frequency as coordinates.

In order to be more general by coordinate we will consider all information related to the data, like measurement times, spatial locations, frequency, etc, especially what is variable for the data array elements. However, this is not necessarily the case, a single scalar coordinate value can be assigned to all elements as well. Coordinate information is not necessarily of numeric type, e.g. channel name can also be considered as coordinate information. On the other hand, other information (e.g. date of the measurement, measurement device configuration information) are not considered as coordinate but stored in the info dictionary of DataObject.

Multiple coordinate information may be present in the DataObject but all of them assign a value to all the array data elements. Storage of the coordinate information is designed by considering that a coordinate mostly changes along one or a few dimensions of the data, in a lot of cases coordinate values are equidistant but in special cases a coordinate value might change along all dimensions of the data array. This way the simple cases are described with minimal amount of data, while enabling even the most complicated case when data is practicably doubled by adding a randomly varying coordinate. Data processing, plotting is optimal if a coordinate changes along one dimension only.

## Representation of coordinates

Coordinates have a name and unit, both described by a string. Standard names are Channel name, Channel number, Signal name, Time, Sample, Device x, Device y, Device z, Device R, Device Z, Device phi, Flux r, Flux Theta, Flux phi, Image x, Image y, Frequency, Time lag. Any other names and units can be used, but it is preferred to use the above where possible. The names are case sensitive as usual in Python. The type of the coordinate values is dependent on the coordinate type. E.g. Sample, is integer, Time is either float or Decimal, Signal name is string.

The following variables are defined in the Coordinate class, but not all of them are used in all definitions: name, unit, mode, shape, step, start, values, value\_index, value\_ranges, dimension\_list. Unused variables are set to None.

Coordinates are not stored in the data matrix but each coordinate description is contained in a FLAP.Coordinate class object. Such an object describes the coordinate values in a d-dimensional rectangular *coordinate sample space* described by the shape variable what is a tuple of sample numbers (s1,s2,...sd) in each dimension, similarly to shape in numpy arrays. If shape has 0 elements it means that the coordinate value is constant and described by the ‘values’ and ‘value\_ranges’ variables. The coordinate sample space is a subarray of the data sample space. As an example consider measurements on a 2D spatial mesh. At each measurement point a time signal is collected, thus the data sample space is 3D. If the 2D mesh is rotated relative to physicsal x,y coordinates then these physical coordinates will change on the 2D mesh. This way the coordinate sample space of x and y will be 2D, while the coordinate samples space for the time coordinate will be 1D. The link between the coordinate sample space and the data sample space is established by the dimension\_list element of FLAP.Coordinate. This has number of elements equal to the dimension of the coordinate sample space and each element contains the index of the related data sample space.

The coordinate values are described in the coordinate sample space [0...s1-1, 0...s2-1, ..., 0...sd-1] in one of two ways.

* If FLAP.Coordinate.mode.equidistant is False samples of the coordinate value are given on a regular or irregular grid in the coordinate sample space. The following cases are considered:
  + If value\_index is None and the shape of the coordinate sample space is identical to the corresponding subspace of the data sample space, then there is a one-to-one correspondence between data samples and coordinate samples. The coordinate values do not change along dimensions which are not in FLAP.Coordinate.dimension\_list.
  + If the two above shapes are different but value\_index is None interpolation is done in the directions with different number of elements assuming that first and last samples match.
  + If value\_index is not None than coordinate samples are on an irregular grid. The coordinate sample locations are given in the ‘value\_index’ (d by Nsamp) array where Nsamp is he number of coordinate samples. The coordinates in the sample space are between 0 and si in the i-th dimension. The coordinate values are given in the 1D ‘values’ array which has Nsamp elements. To calculate the coordinate value for the data array points a (multi-dimensional) interpolation is done between the sample coordinate system and the data sample coordinates.
* If FLAP.Coordinate.mode.equidistant is True then the coordinate sample space is assumed to be identical to the subspace of the data sample space selected by ‘dimension\_list’. (FLAP.Coordinate.shape is not used.) The coordinates change linearly in each dimension: c=b+s1x1+...sdxd , where b is the ‘start’ element of FLAP.Coordinate and si is the step size in dimension i of the data sample space. The si values are stored in the ‘step’ element which is a d long 1D array.

The coordinate values may have a range which is either symmetric or asymmetric around the values. This can be considered either as an error of the coordinate or measurement range, and it is described by a value\_ranges variable. If FLAP.Coordinate.mode.range\_symmetric is True the range is symmetric around the coordinate values, otherwise there is a low and high range. For the equidistant coordinate description ‘value\_ranges’ is either a scalar or 2-element array depending whether the range is symmetric or asymmetric. For the non-equidistant coordinate description in the symmetric case ‘value\_ranges’ has the same shape as ‘values’, for the asymmteric case it is a dictionary with ‘low’ and ‘high’ keys. Each dictionary element has the same shape as values.

The ‘coordinates’ variable of the FLAP.DataObject is a list of FLAP.Coordinate class objects.

## Converting coordinates

Each data source may name a function in the registration process in the add\_coord\_func keyword variable. The add\_coordinate() method of FLAP.DataObject gets coordinate name(s) (string, or string list) and options dictionary. It calls the function registered for the given data source with the data object, the new coordinate name(s) and options arguments. The function should add the named coordinate(s) to the data object or raise a ValueError. The function knows the experiment ID and other information about the data, therefore it should be possible to calculate the new coordinate.

## Explanations and examples

The above definition is complex but it has a reason. It contains all possibilities from the most simple to the most complex. The coordinate descriptions are usually prepared in the data read module and the coordinate values accessed by the data() method of the Coordinate class, therefore the user should not take care of details of the coordinate description. Additionally, the most often encountered cases are very simple, difficulty arises only e.g. when random points are measured in time dependent flux coordinates at random time samples.

In the examples below we do not indicate the coordinate ranges, it can be simply added as described above.

Some typical situations:

* **Constant coordinate.** This is useful where e.g. a measurement is done with all measuring points in the Device z=const. coordinate. This constant can be entered in the DataObject description to be used later when e.g. mapping is done from device to flux coordinates.

shape = []

values = <*z>*

* **Independent equally spaced coordinates along each dimension of the data array.** In this case a coordinate is defined for each dimension of the data array. The definition of each coordinate contains a scalar start and a step value. The shape variable is one number, only the number of elements of shape is used showing that the coordinate description is 1D.

shape = 1

mode.equidistant = True

start = <start>

step = <step>

dimension\_list = [0]

In the above example the coordinate changes along the first dimension of the data array.

* **Array of N temporal signals measured at N different points in the device coordinate space.** The data is stored in a 2D array, one dimension (0) is time, the other is channel. In this case a ‘Time’ coordinate is described with equidistant spacing as shown in the previous example. To describe the measurement spatial coordinates additionally to ‘Time’ 3 coordinates are entered in the coordinates list of the DataObject: ‘Device x’, ‘Device y’ and ‘Device z’. The description for the x coordinate is:

shape = N

mode.equidistant = False

values = *<array with N elements of coordinate values>*

The other two coordinates are entered similarly. The time vector and x,y,z coordinates of measuring channel i can be obtained from the d DataObject as:

time = d.coordinate(‘Time’,(...,i))

x = d.coordinate(‘Device x’,(i,0))

y = d.coordinate(‘Device y’,(i,0))

z = d.coordinate(‘Device z’,(i,0))

In this example it is also useful to additionally define a ‘Signal name’ and maybe a ‘Channel’ coordinate. Signal has normally string values (that is non-equidistant array, values is a list of strings).

* **Fast measurement signals at an array of spatial points mapped to a temporally slowly variable flux coordinate system.**

The data are stored in a 2D array, 1-st dimension is channel, second is time. The data read routine enters the device coordinates into the DataObject. From this the flux coordinate calculation method generates the flux coordinates of the measurement points at a few time points (Nt) during the measurement time. 3 coordinates are added to DataObject, the three flux coordinates. For each coordinate the calculated values are put into a 1D array. The value\_index will be a 2xNt array, at each time point the channel number and the flux coordinate calculation time will be entered. The time is normalized to (t-tstart)/(tend-tstart)\*(Nt-1). The shape variable is (Nch, Nt), where Nch is the number of channels, mode is set to 0 and dimension\_list to [0,1]. As in the channel direction the mapping is 1:1 from the coordinate sample coordinate and the data matrix coordinate no interpolation will occur. In the time direction interpolation will be done and the flux coordinates of each measurement channel will be interpolated values between the sparsely known flux coordinates. The Time coordinate is entered as an equidistant coordinate description.

# Slicing

Slicing means selecting certain elements in the data matrix and optionally taking their sum, minimum, maximum, or doing some other operation on them. Description of the silicing operation is based on coordinates. (Although originally it was foreseen to do slicing along data dimensions, this is not considered useful now.)

Slicing is performed with the slice\_data method. In the slicing argument it takes a dictionary with keys referring to coordinates. The values describe how slicing is done. If the slicing dictionary has multiple keys the slicing operations are done sequentially, except a special case, see below. Summing is done after slicing. (If the slicing argument is omitted only summing is done.) Summing again defined by a dictionary where keys refer to coordinate names.

If the slicing coordinate changes only along one dimension of the data array slicing is done on the data along the associated dimension (see dimension\_list). Other coordinates changing along this dimension are adjusted. It has to be noted that coordinate changes might result in changing from equidistant to non-equidistant type, which can cause more data in the data object. If only one data remains in the sliced dimension that dimension is dropped from the data and also coordinate dimension lists are adjusted correspondingly.

If the slicing coordinate changes along multiple dimensions the situation is more complex as shown in the 2D example below. Here x,y are the original coordinates in a 2D array and R is some coordinate derived from them. The points are arranged in an x-y coordinate system. The orange lines indicate constant R contours. The two solid lines indicate slicing in the R coordinate, the red filled dots are the selected points. Selecting elements in the data in the range of a coordinate which changes in multiple dimensions means that the selected sub-array becomes non-rectangular. In this case the data along these dimensions will be flattened to 1D before slicing and the slicing operation will be done on the flattened dimensions.

x

y

To illustrate this further let us consider a 2D image. A polar coordinate system with origin in the center of the image is introduced and slicing is done in the radial coordinate. Selecting one radial area results in a 1D array. The angle coordinate will change non-monotonically on this. However, coordinates are corrected accordingly and it is still possible to plot as a function e.g. of polar angle.

Two basic slice types are distinguished:

* Simple slice is an operation when a single interval or individual elements are selected along a coordinate. This is described above.
* Multi-slice is an operation when multiple intervals are selected from the data. In this case the dimensions along which the slicing coordinate changes are flattened as described at simple slice and the intervals are selected. Two new dimensions are added to the data matrix. Along one the interval number, along the other the data index inside the intervals change. The interval data are distributed into these new dimensions and the original flattened dimension is removed. With this procedure it becomes possible to plot/sum data in individual intervals or across intervals.

Multi-slice is a complicated procedure and the above described scheme breaks down when multi-slice is intended on two coordinates which change on (partly) common dimensions. E.g. in the above described case of multi-slicing the data on an x-y grid to a r-phi grid poses problems. After multi-slicing with r one gets the two dimensions along and accross the intervals. However, the multi-slicing along phi would flatten these into one dimension and create new intervals in phi. To avoid this multi-slicing along multiple coordinates with common dimensions is done in one step and data are distributed into boxes arranged along each dimension. In case of n such slicing operations n+1 dimensions are added with one dimension where the interval number changes along each coordinate and a single dimension where the sample index in one interval box changes. This case is not implemented yet.

After a multi-slice operation coordinates changing along the flattened dimensions are split into two coordinates: “Rel. <coord> in int(<sl\_coord>” and “Start <coord> in int(<sl\_coord>”. (Except for string coordinates where this is not possible and the original coordinate shape will be changed.) Here <coord> is the name of the coordinate and <sl\_coord> is the name off the slicing coordinate. Also coordinates with names “Interval(<sl\_coord>)” and “Interval(<sl\_coord> index)” are added storing the interval number (along one coordinate) and the sample index in one interval.

If multi-slice operation results in different interval length, the dimension along the samples in the intervals will be set to the longest. Where data is shorter in one interval np.NaN values will be filled in case of float data and 0 for int. (There is no integer Not-a-number value in Python.) In the coordinate matrix missing elements will be filled similarly to data.)

Slicing can be described with the following objects:

1. For simple slice:
   1. A Python slice or range object, to select a sequence of regularly spaced elements.
   2. A scalar value or a list of scalars to select random elements.
   3. A numpy array to select random elements.
   4. flap.DataObject without error and with data unit.name equal to the slicing coordinate name.
   5. flap.DataObject with the data unit.name not equal to the slicing coordinate, but one of the coordinate names equal to the slicing coordinate and the coordinate has no value\_ranges.
   6. flap.Intervals object with one interval.

In cases 1a-e the data elements with closest coordinates will be selected, while in case 1f all elements in the interval will be selected. In case a string type slicing coordinate matching between the slicing and the coordinate value is required instead of close match. (There is no sense in close match for strings.) However, extended wildcards can also be used, e.g. slicing=’{Signal name’:’TEST-\*-3’} is a valid slicing expression.

1. For multi-slice:
   * + - 1. flap.Intervals object with more than one interval.
         2. flap.DataObject with data unit.name equal to the slicing coordinate. The error values give the intervals.
         3. flap.DataObject with the data unit.name not equal to the slicing coordinate name but name of one coordinates equal to the slicing coordinate. The value\_ranges select the intervals.

The summing input argument to the slice\_data method can be used for processing the sliced data. This is also a dictionary with coordinate names as keys. Before processing the dimensions where the summing coordinate changes will be flattened. The values of the dictionary can be the following:

* ‘Sum’: Add all elements.
* ‘Mean’: Take the mean of all elements
* ‘Min’: Take the minimum of all elements
* ‘Max’: Take the maximum of all elements

As the result of the processing is a single value along the summing coordinate, this dimension will be removed from the data. After processing the data the coordinate changes will be done. In the case of ‘Sum’ and ‘Mean’ the mean of the coordinates of the summed data will be taken, while in the case of ‘Min’ and ‘Max’ the coordinate of the minimum or maximum value will be selected.

The slice\_data method of flap.DataObject is complemented by the flap.slice\_data function which operates on data objects in flap storage.

## Examples

As an example we read all signals from the TESTDATA module for a 1 ms piece and store under name TESTDATA in flap storage:

**d=flap.get\_data('TESTDATA',name='\*', options={'Scaling':'Volt'},**

**object\_name='TESTDATA', coordinates={'Time':[0,0.001]})**

This results in a 3D data object where signals are arranged in row and column and the third dimension is time. ‘Time’, ‘Sample’, ‘Row’, ‘Column’ and ‘Signal name’ coordinates are supplied by the data read routine. Then we add spatial coordinates:

**flap.add\_coordinate('TESTDATA',**

**coordinates=['Device x','Device z','Device y'])**

We can list the content of the data object using the flap.list\_data\_objects() call:

**TESTDATA(exp\_id:None) "Test data" shape:[15,10,1001]**

**Coords:**

**'Sample'[n.a.](Dims:2) [<Equ.><R. symm.>] Start: 0.000E+00, Steps: 1.000E+00**

**'Time'[Second](Dims:2) [<Equ.><R. symm.>] Start: 0.000E+00, Steps: 1.000E-06**

**'Signal name'[n.a.](Dims:0,1, Shape:15,10) [<R. symm.>] Val:TEST-1-1, TEST-1-2, TEST-1-3, TEST-1-4, TEST-1-5, TEST-1-6, TEST-1-7, TEST-1-8, TEST-1-9, TEST-1-10, ...**

**'Column'[n.a.](Dims:0, Shape:15) [<R. symm.>] Val. range: 1.000E+00 - 1.500E+01**

**'Row'[n.a.](Dims:1, Shape:10) [<R. symm.>] Val:1, 2, 3, 4, 5, 6, 7, 8, 9, 10**

**'Device x'[cm](Dims:0,1, Shape:15,10) [<R. symm.>] Val. range: -1.112E+00 - 6.657E+00**

**'Device z'[cm](Dims:0,1, Shape:15,10) [<R. symm.>] Val. range: 0.000E+00 - 5.587E+00**

**'Device y'[cm](Dims:, Shape:) [<R. symm.>] Val: 0.000E+00**

The above list shows that Sample and Time changes along dimension 2 (3-rd dimension), Signal names change on the first tow dimensions, Column and Row changes on dimension 0 and 1, respectively. Device y does not change at all (emply dimension list) as the measurement channels are in the y=0 plane. Device x and y both change on dimensions 0,1 as the measurement matrix is inclined in the x, y plane.

A simple slice to select one signal looks like:

**flap.slice\_data('TESTDATA', slicing={'Signal name': 'TEST-1-3'},**

**output\_name='TESTDATA\_slice')**

The result is a 1D array:

**TESTDATA\_slice(exp\_id:None) "Test data" shape:[1001]**

**Coords:**

**'Sample'[n.a.](Dims:0) [<Equ.><R. symm.>] Start: 0.000E+00, Steps: 1.000E+00**

**'Time'[Second](Dims:0) [<Equ.><R. symm.>] Start: 0.000E+00, Steps: 1.000E-06**

**'Signal name'[n.a.](Dims:, Shape:1) [<R. symm.>] Val:TEST-1-3**

**'Column'[n.a.](Dims:, Shape:1) [<R. symm.>] Val:1**

**'Row'[n.a.](Dims:, Shape:1) [<R. symm.>] Val:3**

**'Device x'[cm](Dims:, Shape:1) [<R. symm.>] Val:-2.472E-01**

**'Device z'[cm](Dims:, Shape:1) [<R. symm.>] Val: 7.608E-01**

**'Device y'[cm](Dims:, Shape:) [<R. symm.>] Val: 0.000E+00**

Extended regular expressions can also be used. In the following expression 3 signals are selected, resulting in a 3x1001 data array:

**TESTDATA\_slice(exp\_id:None) "Test data" shape:[3,1001]**

**Coords:**

**'Sample'[n.a.](Dims:1) [<Equ.><R. symm.>] Start: 0.000E+00, Steps: 1.000E+00**

**'Time'[Second](Dims:1) [<Equ.><R. symm.>] Start: 0.000E+00, Steps: 1.000E-06**

**'Signal name'[n.a.](Dims:0, Shape:3) [<R. symm.>] Val:TEST-1-8, TEST-1-9, TEST-1-10**

**'Column'[n.a.](Dims:0, Shape:3) [<R. symm.>] Val:1, 1, 1**

**'Row'[n.a.](Dims:0, Shape:3) [<R. symm.>] Val:8, 9, 10**

**'Device x'[cm](Dims:0, Shape:3) [<R. symm.>] Val:-8.652E-01, -9.889E-01, -1.112E+00**

**'Device z'[cm](Dims:0, Shape:3) [<R. symm.>] Val: 2.663E+00, 3.043E+00, 3.424E+00**

**'Device y'[cm](Dims:, Shape:) [<R. symm.>] Val: 0.000E+00**