genopt Documentation

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Tong Zhang

CONTENTS

1	Introduction	3		
2	Demonstrations2.1 Getting started2.2 Setup BPMs, correctors and reference orbit2.3 Setup variables2.4 Setup optimization engine2.5 Run optimization2.6 After optimization	6 7 8 9		
3	API 3.1 genopt package	15 15		
4	1 Indices and tables			
Рy	thon Module Index	39		
In	dex	41		

genopt Python package

genopt: general multi-dimensional optimization

PDF documentation: Download

Github repo: https://github.com/archman/genopt

Author Tong Zhang

E-mail zhangt@frib.msu.edu

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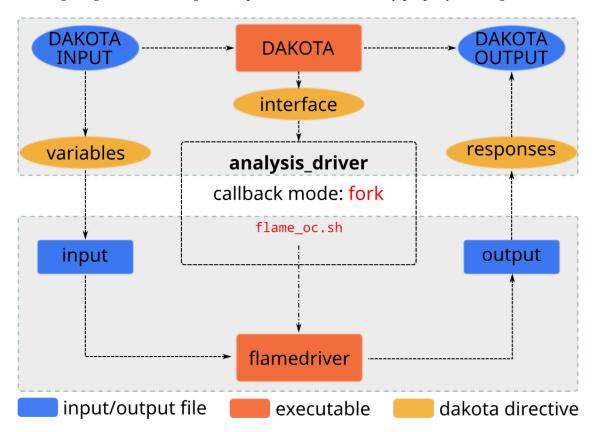
CONTENTS 1

2 CONTENTS

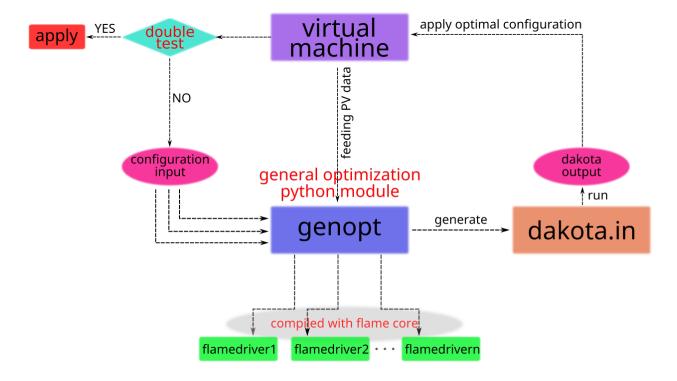
INTRODUCTION

genopt is a python package, trying to serve as a solution of general multi-dimensional optimization. The core optimization algorithms employed inside are mainly provided by DAKOTA, which is the brief for *Design Analysis Kit for Optimization and Terascale Applications*, another tool written in C++.

The following image illustrates the general optimization framework by properly utilizing DAKOTA.



To apply this optimization framework, specific analysis drivers should be created first, e.g. flamedriver1, flamedriver2... indicate the dedicated executable drivers built from C++, for the application in accelerator commissioning, e.g. FRIB.



Note: flame is an particle envolope tracking code developed by C++, with the capbility of multi-charge particle states mroementum space tracking, it is developed by FRIB; flamedriver(s) are user-customized executables by linking the flame core library to accomplish various requirements.

The intention of genopt is to provide a uniform interface to do the multi-dimensional optimization tasks. It provides interfaces to let the users to customize the optimization drivers, optimization methods, variables, etc. The optimized results are returned by clean interface. Dedicated analysis drivers should be created and tell the package to use. DakotaOC is a dedicated class designed for orbit correction for accelerator, which uses flame as the modeling tool.

DEMONSTRATIONS

Here goes some examples to use genopt package to do orbit correction, it should be noted that the more complicated the script is, the more options could be adjusted to fulfill specific goals.

Getting started

This approach requires fewest input of code to complete the orbit correction optimization task, which also means you only has very few options to adjust to the optimization model. Hopefully, this approach could be used as an ordinary template to fulfill most of the orbit correction tasks. Below is the demo code:

```
import genopt
latfile = 'test_392.lat'
oc_ins = genopt.DakotaOC(lat_file=latfile)
oc_ins.simple_run(method='cg', mpi=True, np=4, iternum=20)
# get output
oc_ins.get_orbit(outfile='orbit.dat')
# plot
oc_ins.plot()
```

The lattice file used here could be found from here, or from https://github.com/archman/genopt/blob/master/lattice/test_392.lat.

For this approach, the following default configuration is applied:

- 1. Selected all BPMs and correctors (both horizontal and vertical types);
- 2. Set the reference orbit with all BPMs' readings of x=0 and y=0;
- 3. Set the objective function with the sum of all the square of orbit deviations w.r.t. reference orbit.

By default, conmin_frcg optimization method is used, possible options for simple_run() could be:

- common options:
 - 1. mpi: if True, run in parallel mode; if False, run in serial mode;
 - 2. np: number of cores to use if mpi is True;
- gradient descent, i.e. method=cg:
 - 1. iternum: max iteration number, 20 by default;
 - 2. step: forward gradient step size, 1e-6 by default;
- pattern search, i.e. method=ps:
 - 1. iternum: max iteration number, 20 by default;

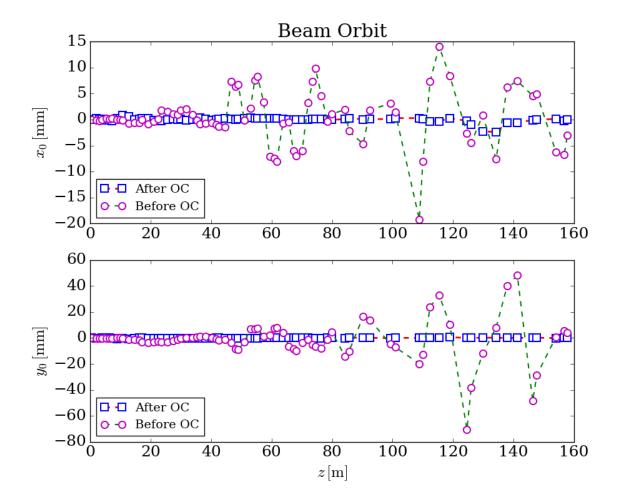
2. evalnum: max function evaulation number, 1000 by default;

There are two options for DakotaOC maybe useful sometimes:

- 1. workdir: root directory for dakota input and output files
- 2. keep: if keep working files, True or False

After run this script, beam orbit data could be saved into file, e.g. orbit.dat:

which could be used to generate figures, the following figure is a typical one could be generated from the optimized results:



Setup BPMs, correctors and reference orbit

For more general cases, genopt provides interfaces to setup BPMs, correctors, reference orbit and objective function type, etc., leaving more controls to the user side, to fulfill specific task.

Here is an exmaple to show how to use these capabilities.

```
import genopt
# lattice file
latfile = 'test_392.lat'
oc_ins = genopt.DakotaOC(lat_file=latfile)
```

```
# select BPMs
bpms = oc_ins.get_elem_by_type('bpm')
oc_ins.set_bpms(bpm=bpms)
# select correctors
hcors = oc_ins.get_all_cors(type='h')[0:40]
vcors = oc_ins.get_all_cors(type='v')[0:40]
oc_ins.set_cors(hcor=hcors, vcor=vcors)
# setup objective function type
oc_ins.ref_flag = "xy"
\# setup reference orbit in x and y
bpms_size = len(oc_ins.bpms)
oc_ins.set_ref_x0(np.ones(bpms_size)*0.0)
oc_ins.set_ref_y0(np.ones(bpms_size)*0.0)
# run optimizaiton
oc_ins.simple_run(method='cg', mpi=True, np=4, iternum=30)
oc_ins.get_orbit(outfile='orbit.dat')
oc_ins.plot()
```

The highlighted code block is added for controlling all these abovementioned properties.

Warning:

- 1. BPMs and correctos are distinguished by the element index, which could be get by proper method, e.g. get_all_cors();
- 2. The array size of selected BPMs and reference orbit should be same;
- 3. bpms, hcors, vcors are properties of DakotaOC instance.

Note: Objective functions could be chosen from three types according to the value of ref_flag:

```
1. ref_flag="xy": \sum \Delta x^2 + \sum \Delta y^2
```

2. ref_flag="x":
$$\sum \Delta x^2$$

3. ref_flag="y":
$$\sum \Delta y^2$$

where $\Delta x = x - x_0$, $\Delta y = y - y_0$.

Setup variables

By default the variables to be optimized is setup with the following parameters:

initial value	lower bound	upper bound
1e-4	-0.01	0.01

However, subtle configuration could be achieved by using set_variables() method of DakotaOc class, here is how to do it:

Parameter could be created by using DakotaParam class, here is the code:

plist_y could be created in the same way, then issue set_variables() with set_variables(plist=plist_x+plist_y).

Note: The emphasized line is to setup the variable labels, it is recommended that all parameters' label with the format like x001, x002, etc.

Setup optimization engine

The simplest approach, (see *Getting started*), just covers detail of the more specific configurations, especially for the optimization engine itself, however genopt provides different interfaces to make customized adjustment.

Method

DakotaMethod is designed to handle method block, which is essential to define the optimization method, e.g.

Interface

DakotaInterface is designed to handle interface block, for the general optimization regime, fork mode is the common case, only if the analysis driver is compile into dakota, direct could be used.

Here is an example of user-defined interface:

Note: Extra parameters could be added by this way: oc_inter.set_extra(deactivate="active_set_vector")

Responses

Objective function(s) and gradients/hessians could be set in responses block, which is handled by DakotaResponses class.

Typical example:

```
oc_responses = DakotaResponses(gradient='numerical', step=2.0e-7)
oc_ins.set_responses(oc_responses)
```

Environment

Dakota environment block could be adjusted by instantiating class DakotaEnviron, e.g.

```
datfile = 'dakotal.dat'
e = genopt.DakotaEnviron(tabfile=datfile)
oc_ins.set_environ(e)
```

tabfile option could be used to define where the dakota tabular data should go, will not generate tabular file if not set.

Model

DakotaModel is designed to handle model block, recently, just use the default configuration, i.e:

```
oc_ins.set_model()
# or:
m = genopt.DakotaModel()
oc_ins.set_model(m)
```

Run optimization

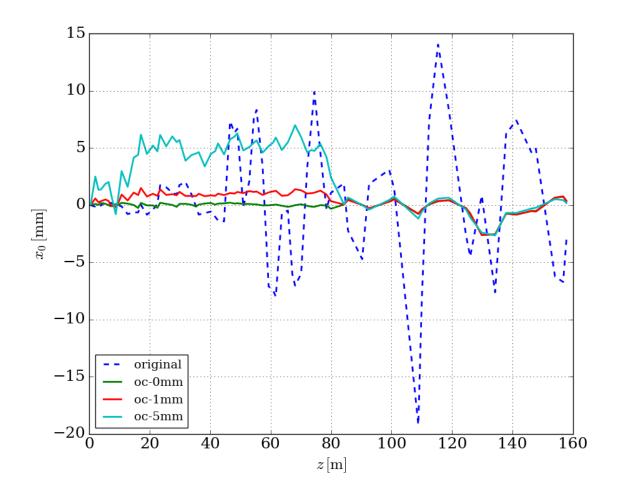
If running optimization not by simple_run() method, another approach should be utilized.

```
# generate input file for optimization
oc_ins.gen_dakota_input()
# run optimization
oc_ins.run(mpi=True, np=4)
```

Below is a typical user customized script to find the optimized correctors configurations.

```
# set interface
oc_ins.set_interface()
# set responses
r = genopt.DakotaResponses(gradient='numerical', step=2.0e-5)
oc_ins.set_responses(r)
# set model
m = genopt.DakotaModel()
oc_ins.set_model(m)
# set method
md = genoptDakotaMethod(method='ps',
       max_function_evaluations=1000)
oc_ins.set_method(method=md)
# set environment
tabfile = os.path.abspath('./oc_tmp4/dakota1.dat')
e = genopt.dakutils.DakotaEnviron(tabfile=tabfile)
oc_ins.set_environ(e)
# set reference orbit
bpms_size = len(oc_ins.bpms)
ref_x0 = np.ones(bpms_size)*0.0
ref_y0 = np.ones(bpms_size)*0.0
oc_ins.set_ref_x0(ref_x0)
oc_ins.set_ref_y0(ref_y0)
# set objective function
oc_ins.ref_flag = "xy"
# generate input
oc_ins.gen_dakota_input()
oc_ins.run(mpi=True, np=4)
#print oc_ins.get_opt_results()
oc_ins.get_orbit((oc_ins.hcor, oc_ins.vcor), oc_ins.get_opt_results(),
                 outfile='orbit.dat')
# plot
#oc_ins.plot()
```

The following figure shows correct the orbit to different reference orbits.



After optimization

Suppose all the optimized results have been generated, here are the possible post-operations:

- 1. Operations on the optimized Machine object;
- 2. Generate new lattice file with optimized results for other programs.

Optimization snippet:

```
latfile = 'test_392.lat'
oc = genopt.DakotaOC(lat_file=latfile)
oc.simple_run(iternum=20)
```

Get optimized results

Optimized results could be retrieved by get_opt_results() method of DakotaOC class:

• return type: list

```
>>> r = oc.get_opt_results(rtype='list')
>>> print(r)
```

```
[0.00013981587907,
7.5578423135e-05,
-5.3982438406e-05,
-1.9620020032e-06,
0.000179420779806,
...
2.0182502319e-05,
0.0001733634281,
8.685656753e-05,
7.3950720611e-05,
8.2924283647e-05]
```

The returned list is alphabetically sorted according to the variables' names.

• return type: dictionary, label format: plain

```
>>> r = oc.get_opt_results()
>>> # or
>>> r = oc.get_opt_results(rtype='dict', label='plain')
>>> print(r)
{'x001': 0.00013981587907,
'x002': 7.5578423135e-05,
'x003': -5.3982438406e-05,
'x004': -1.9620020032e-06,
'x005': 0.00017942079806,
...
'y056': 2.0182502319e-05,
'y057': 0.0001173634281,
'y058': 8.68565675a-05,
'y059': 7.3950720611e-05,
'y060': 8.2924283647e-05}
```

• return type: dictionary, label format: fancy

```
>>> r = oc.get_opt_results(label='fancy')
>>> print(r)
{'FS1_BBS:DCH_D2412': {'config': {'theta_x': 0.00021066533055}, 'id': 1048},
'FS1_BBS:DCH_D2476': {'config': {'theta_x': 0.00025833402592}, 'id': 1098},
...
```

This is the more comprehensive way to represent the results, one of the advantages is results with this format could be easily to apply on to reconfigure method of Machine object, for instance:

```
>>> for k,v in r.items():
>>> m.reconfigure(v['id'], v['config'])
```

Note: get_opt_results has outfile optional parameter, if not defined, use output file that generated by current optimization instance, or use the defined dakota output file, but only valid for cases of label='plain'; label='fancy' is only valid for the case of rtype='dict'.

Get orbit data

get_orbit() could be used to apply all the optimized results, then new Machine could be get in the following way:

```
>>> z,x,y,m = oc.get_orbit()
>>> print(m.conf(1224)['theta_x'])
8.5216269467e-05
```

Or in another way:

```
>>> oc.get_orbit()
>>> m = oc.get_machine()
```

New machine m could be used for the next operations.

Note: get_orbit() could be assigned a optional parameter: outfile, into which the plain ASCII data of zpos, x, and y would be saved.

Get new optimized lattice file

get_opt_latfile() is created to generate new lattice file with optimized results, for the sake of possible next usage of asking for lattice file, this is kind of more general interface.

```
>>> oc.get_opt_latfile(outfile='opt1.lat')
```

Here is the links to the lattice files of original and optimized ones, both could be used as the input lattice file of flame program.

CHAPTER

THREE

API

genopt package

General multi-dimensional optimization package built by Python, incorporating optimization algorithms provided by DAKOTA.

```
version 0.0.2

author Tong Zhang <zhangt@frib.msu.edu>
```

Example

```
>>> # This is a ordinary example to do orbit correction by
>>> # multi-dimensional optimization approach.
>>> # import package
>>> import genopt
>>> # lattice file name
>>> latfile = './contrib/test_392.lat'
>>> # create optimization object
>>> oc_ins = genopt.DakotaOC(lat_file=latfile)
>>> # get indices of BPMs and correctors
>>> bpms = oc_ins.get_elem_by_type('bpm')
>>> cors = oc_ins.get_all_cors()[45:61]
>>> # set BPMs and correctors
>>> oc_ins.set_bpms(bpm=bpms)
>>> oc_ins.set_cors(cor=cors)
>>> # run optimization, enable MPI,
>>> # with optimization of CG, 20 iterations
>>> oc_ins.simple_run(method='cg', mpi=True, np=4, iternum=20)
>>> # get optimized results:
>>> opt_vars = oc_ins.get_opt_results()
>>> # or show orbit after correction
>>> oc_ins.plot()
>>> # or save the orbit data (to file)
>>> oc_ins.get_orbit(outfile='orbit.dat')
```

class DakotaInput(**kws)

Bases: object

template of dakota input file, field could be overriden by providing additional keyword arguments,

Parameters kws – keyword arguments, valid keys are dakota directives

Example

```
>>> dak_inp = DakotaInput(method=["max_iterations = 500",
                                 "convergence_tolerance = 1e-7",
                                 "conmin_frcg",])
     set_template(name='oc')
     write(infile=None)
          write all the input into file, as dakota input file
               Parameters infile - fullname of input file, if not defined, infile will be assigned as
                   'dakota.in' in current working directory
class DakotaParam(label, initial=0.0, lower=-10000000000.0, upper=10000000000.0)
     Bases: object
     create dakota variable for variables block
          Parameters
                • label – string to represent itself, e.g. x001, it is recommended to annotate the num-
                  ber with the format of %03d, i.e. 1 --> 001, 10 --> 010, 100 --> 100, etc.
                • initial – initial value, 0.0 by default
                • lower – lower bound, -1.0e10 by default
                • upper – upper bound, 1.0e10 by default
     initial
     label
     lower
     upper
class DakotaBase(**kws)
     Bases: object
     Base class for general optimization, initialized parameters: valid keyword parameters:

    workdir: root dir for dakota input/output files, the defualt one should be created in /tmp, or

          define some dir path
         •dakexec: full path of dakota executable, the default one should be dakota, or define the full path
         •dakhead: prefixed name for input/output files of dakota, the default one is dakota
         • keep: if keep the working directory (i.e. defined by workdir), default is False
     dakexec
     dakhead
     keep
     workdir
class DakotaOC(lat_file=None, elem_bpm=None, elem_cor=None, elem_hcor=None, elem_vcor=None,
                ref_x0=None, ref_y0=None, ref_flag=None, model=None, optdriver=None, **kws)
     Bases: genopt.dakopt.DakotaBase
```

Dakota optimization class with orbit correction driver

Parameters

• lat_file – lattice file

- **elem_bpm** list of element indice of BPMs
- elem_cor list of element indice of correctors, always folders of 2
- elem_hcor list of element indice of horizontal correctors
- elem_vcor list of element indice of vertical correctors
- ref_x0 reference orbit in x, list of BPM readings
- ref_y0 reference orbit in y, list of BPM readings
- ref_flag string flag for objective functions:

1. "x":
$$\sum \Delta x^2$$
, $\Delta x = x - x_0$;

2. "y":
$$\sum \Delta y^2$$
, $\Delta y = y - y_0$;

3. "xy":
$$\sum \Delta x^2 + \sum \Delta y^2$$
.

- model simulation model, 'flame' or 'impact'
- optdriver analysis driver for optimization, 'flamedriver_oc' by default
- kws -

keywords parameters for additional usage, defined in DakotaBase class valid keys:

- workdir: root dir for dakota input/output files, the defualt one should be created in /tmp, or define some dir path
- dakexec: full path of dakota executable, the default one should be dakota, or define the full path
- dakhead: prefixed name for input/output files of dakota, the default one is dakota
- keep: if keep the working directory (i.e. defined by workdir), default is False

bpms

create_machine(lat_file)

create machine instance with model configuration

- setup _machine
- setup _elem_bpm, _elem_cor or (_elem_hcor and _elem_vcor)

gen_dakota_input(infile=None, debug=False)

generate dakota input file

Parameters

- **infile** dakota input filename
- **debug** if True, generate a simple test input file

get_all_bpms()

get list of all valid bpms indices

Returns a list of bpm indices

Example

```
>>> dakoc = DakotaOC('test/test.lat')
>>> print(dakoc.get_all_bpms())
```

```
get_all_cors(type=None)
```

get list of all valid correctors indices

Parameters type – define corrector type, 'h': horizontal, 'v': vertical, if not defined, return all correctors

Returns a list of corrector indices

Example

```
>>> dakoc = DakotaOC('test/test.lat')
>>> print(dakoc.get_all_cors())
```

get_elem_by_name(name)

get list of element(s) by name(s)

Parameters name – tuple or list of name(s)

Returns list of element indices

Example

```
>>> dakoc = DakotaOC('test/test.lat')
>>> names = ('LS1_CA01:BPM_D1144', 'LS1_WA01:BPM_D1155')
>>> idx = dakoc.get_elem_by_name(names)
>>> print(idx)
[18, 31]
```

get_elem_by_type(type)

get list of element(s) by type

Parameters type – string name of element type

Returns list of element indices

Example

```
>>> dakoc = DakotaOC('test/test.lat')
>>> type = 'bpm'
>>> idx = dakoc.get_elem_by_type(type)
>>> print(idx)
```

get_machine()

get flame machine object for potential usage

Returns flame machine object or None

```
get_opt_latfile(outfile='out.lat')
```

get optimized lattice file for potential next usage, run() or simple_run() should be evoked first to get the optimized results.

Parameters outfile – file name for generated lattice file

Returns lattice file name or None if failed

```
get_opt_results(outfile=None, rtype='dict', label='plain')
```

extract optimized results from dakota output

Parameters

- outfile file name of dakota output file, 'dakota.out' by default
- rtype type of returned results, 'dict' or 'list', 'dict' by default
- label label types for returned variables, only valid when rtype 'dict', 'plain' by default:

- 'plain': variable labels with format of x1, x2, y1, y2, etc. e.g. {'x1': v1, 'y1': v2}

Note: The fancy option will make re-configuring flame machine in a more convenient way, such as:

```
>>> opt_cors = get_opt_results(label='fancy')
>>> for k,v in opt_cors.items():
>>> m.reconfigure(v['id'], v['config'])
>>> # here m is an instance of flame.Machine class
>>>
```

Returns by default return a dict of optimized results with each item of the format like "x1":0.1 or more fancy format by set label with 'fancy', etc., if rtype='list', return a list of values, when the keys are ascend sorted.

Example

```
>>> opt_vars = get_optresults(outfile='flame_oc.out', rtype='dict'):
>>> print(opt_vars)
{'x2': 0.0020782814353, 'x1': -0.0017913264033}
>>> opt_vars = get_optresults(outfile='flame_oc.out', rtype='list'):
>>> print(opt_vars)
[-0.0017913264033, 0.0020782814353]
```

get_orbit(idx=None, val=None, outfile=None)

calculate the orbit with given configurations

Parameters

- idx (idx_hcor, idx_vcor), tuple of list of indices of h/v cors
- val values for each correctos, h/v
- outfile filename to save the data

Returns tuple of zpos, env_x, env_y, machine

hcors

latfile

optdriver

Parameters

- outfile output file of dakota
- figsize figure size, (h, w)
- **dpi** figure dpi

```
ref_flag
```

ref_x0

ref_y0

```
run(mpi=False, np=None)
    run optimization
```

- mpi if True, run DAKOTA in parallel mode, False by default
- **np** number of processes to use, only valid when mpi is True

set_bpms(bpm=None)
set BPMs

Parameters bpm – list of bpm indices, if None, use all BPMs

set_cors(cor=None, hcor=None, vcor=None)

set correctors, if cor, hcor and vcor are None, use all correctors if cor is not None, use cor, ignore hcor and vcor

Parameters

- cor list of corrector indices, hcor, vcor,...
- hcor list of horizontal corrector indices
- vcor list of vertical corrector indices

set_environ(environ=None)

setup environment block, that is setup oc_environ

Parameters environ - DakotaEnviron object, automatically setup if not defined

set_interface(interface=None, **kws)

setup interface block, that is setup oc_interface should be ready to invoke after set_cors and set_bpms

Parameters interface - DakotaInterface object, automatically setup if not defined

set_method(method=None)

setup method block, that is setup oc_method

Parameters method – DakotaMethod object, automatically setup if not defined

set_model(model=None, **kws)

setup model block, that is setup oc_model

Parameters model - DakotaModel object, automatically setup if not defined

set_ref_x0(ref_arr=None)

set reference orbit in x, if not set, use 0s

Parameters ref_arr – array of reference orbit values size should be the same number as selected BPMs

set_ref_y0(ref_arr=None)

set reference orbit in y, if not set, use 0s

Parameters ref_arr – array of reference orbit values size should be the same number as selected BPMs

set_responses(responses=None, **kws)

setup responses block, that is setup oc_responses

Parameters responses – DakotaResponses object, automatically setup if not defined

set_variables(plist=None, initial=0.0001, lower=-0.01, upper=0.01)

setup variables block, that is setup oc_variables should be ready to invoke after set_cors()

- plist list of defined parameters (DakotaParam object), automatically setup if not defined
- initial initial values for all variables, only valid when plist is None
- lower lower bound for all variables, only valid when plist is None
- upper upper bound for all variables, only valid when plist is None

```
simple_run(method='cg', mpi=None, np=None, **kws)
```

run optimization after set_bpms() and set_cors(), by using default configuration and make full use of computing resources.

Parameters

- **method** optimization method, 'cg', 'ps', 'cg' by default
- mpi if True, run DAKOTA in parallel mode, False by default
- np number of processes to use, only valid when mpi is True
- **kws** keyword parameters valid keys:
 - step: gradient step, 1e-6 by default
 - iternum: max iteration number, 20 by default
 - evalnum: max function evaulation number, 1000 by default

vcors

class DakotaEnviron(tabfile=None, **kws)

Bases: object

create datako environment for environment block

Parameters

- tabfile tabular file name, by default not save tabular data
- kws other keyword parameters

Example

```
>>> # default
>>> oc_environ = DakotaEnviron()
>>> print oc_environ.get_config()
[]
>>> # define name of tabular file
>>> oc_environ = DakotaEnviron(tabfile='tmp.dat')
>>> print oc_environ.get_config()
['tabular_data', " tabular_data_file 'tmp.dat'"]
>>>
```

```
get_config(rtype='list')
```

get responses configuration for dakota input block

Parameters rtype - 'list' or 'string'

Returns dakota responses input

class DakotaInterface(mode='fork', driver='flamedriver_oc', latfile=None, bpms=None, hcors=None, vcors=None, ref_x0=None, ref_y0=None, ref_flag=None, **kws)

Bases: object

create dakota interface for interface block

Parameters

- mode 'fork' or 'direct' (future usage)
- driver analysis driver, external ('fork') executable file internal ('direct') executable file
- latfile file name of (flame) lattice file
- **bpms** array of selected BPMs' id
- hcors array of selected horizontal (x) correctors' id
- vcors array of selected vertical (y) correctors' id
- ref_x0 array of BPM readings for reference orbit in x, if not defined, use 0s
- ref_y0 array of BPM readings for reference orbit in y, if not defined, use 0s
- ref_flag string flag for objective functions:
 - 1. "x": $\sum \Delta x^2$, $\Delta x = x x_0$;
 - 2. "y": $\sum \Delta y^2$, $\Delta y = y y_0$;
 - 3. "xy": $\sum \Delta x^2 + \sum \Delta y^2$.
- kws keyword parameters, valid keys: e.g.: * deactivate, possible value: 'active_set_vector'

Note: mode should be set to be 'direct' when the analysis drivers are built with dakota library, presently, 'fork' is used.

Example

```
>>> # for orbit correction
>>> bpms = [1,2,3] # just for demonstration
>>> hcors, vcors = [1,3,5], [2,4,6]
>>> latfile = 'test.lat'
>>> oc_interface = DakotaInterface(mode='fork',
                                      driver='flamedriver_oc',
>>>
                                      latfile=latfile
                                      bpms=bpms, hcors=hcors, vcors=vcors,
>>>
                                      ref_x0=None, ref_y0=None,
>>>
                                      ref_flag=None,
                                      deactivate='active_set_vector')
>>> # add extra configurations
>>> oc_interface.set_extra(p1='v1', p2='v2')
>>> # get configuration
>>> config_str = oc_interface.get_config()
```

bpms

driver

```
get_config(rtype='list')
```

get interface configuration for dakota input block

Parameters rtype - 'list' or 'string'

Returns dakota interface input

hcors

latfile

mode

```
ref_flag
ref_x0
ref_y0
set_extra(**kws)
    add extra configurations
vcors
class DakotaMethod(method='cg', iternum=20, tolerance=0.0001, **kws)
    Bases: object
    create dakota method for method block
```

- method method name, 'cg' by default, all possible choices: 'cg', 'ps'
- iternum max iteration number, 20 by default
- tolerance convergence tolerance, 1e-4 by default
- **kws** other keyword parameters

Example

```
>>> oc_method = DakotaMethod()
>>> print oc_method.get_config()
['conmin_freg', ' convergence_tolerance 0.0001', ' max_iterations 20']
>>> # define method with pattern search
>>> oc_method = DakotaMethod(method='ps')
>>> print oc_method.get_config()
['coliny_pattern_search', ' contraction_factor 0.75', ' max_function_evaluations 500', ' solution_accuracy 0.0001', ' exploratory_moves basic_pattern', ' threshold_delta 0.0001', ' initial_delta 0.5', ' max_iterations 100']
>>> # modify options of pattern search method
>>> oc_method = DakotaMethod(method='ps', max_iterations=200, contraction_factor=0.8)
 >>> print oc_method.get_config()
 \hbox{['coliny\_pattern\_search', 'contraction\_factor 0.8', 'max\_function\_evaluations 500', } \\
 'solution_accuracy 0.0001', 'exploratory_moves basic_pattern', 'threshold_delta 0.0001', 'initial_delta 0.5', 'max_iterations 200']
>>> # conmin_frcg method
>>> oc_method = DakotaMethod(method='cg')
 >>> print oc_method.get_config()
['conmin_frcg', ' convergence_tolerance 0.0001', ' max_iterations 20']
>>> # modify options
>>> oc_method = DakotaMethod(method='cg', max_iterations=100)
>>> print oc_method.get_config()
['conmin_frcg', ' convergence_tolerance 0.0001', ' max_iterations 100']
```

```
get_config(rtype='list')
```

get method configuration for dakota input block

Parameters rtype – 'list' or 'string'

Returns dakota method input

get_default_method(method)

get default configuration of some method

Parameters method – method name, 'cg' or 'ps'

Returns dict of configuration

method(method)

return method configuration

Parameters method - method stirng name, 'cg' or 'ps'

```
Returns list of method configuration
```

```
class DakotaModel(**kws)
    Bases: object
    create dakota model for model block
    get_config(rtype='list')
        get model configuration for dakota input block
        Parameters rtype - 'list' or 'string'
        Returns dakota model input
class DakotaResponses(nfunc=1, gradient=None, hessian=None, **kws)
        Bases: object
    create dakota responses for responses block
```

- **nfunc** num of objective functions
- gradient gradient type: 'analytic' or 'numerical'
- hessian hessian configuration
- kws keyword parameters for gradients and hessians valid keys: any available for responses among which key name of 'grad' is for gradients configuration, the value should be a dict (future)

Example

```
>>> # default responses:
>>> response = DakotaResponses()
 >>> print response.get_config()
['num_objective_functions = 1', 'no_gradients', 'no_hessians']
>>> # responses with analytic gradients:
>>> response = DakotaResponses(gradient='analytic')
>>> print response.get_config()
['num_objective_functions = 1', 'analytic_gradients', 'no_hessians']
>>> # responses with numerical gradients, default configuration:
>>> oc_responses = DakotaResponses(gradient='numerical')
>>> print oc_responses.get_config()
['num_objective_functions = 1', 'numerical_gradients', ' method_source dakota', ' interval_type forward', ' fd_gradient_step_size 1e-06', 'no_hessians']
>>> # responses with numerical gradients, define step:
>>> oc_responses = DakotaResponses(gradient='numerical', step=2.0e-7)
>>> print oc_responses.get_config()
['num_objective_functions = 1', 'numerical_gradients', ' method_source dakota', ' interval_type forward', ' fd_gradient_step_size 2e-07', 'no_hessians']
>>> # given other keyword parameters:
>>> oc_responses = DakotaResponses(gradient='numerical', step=2.0e-7, k1='v1', k2='v2')
>>> print oc_responses.get_config()
['num_objective_functions = 1', 'numerical_gradients', ' method_source dakota',
' interval_type forward', ' fd_gradient_step_size 2e-07', 'no_hessians', 'k2 = v2', 'k1 = v1']
```

```
Parameters rtype – 'list' or 'string'
Returns dakota responses input
gradients(type=None, step=1e-06, **kws)
generate gradients configuration
```

get responses configuration for dakota input block

get_config(rtype='list')

- type 'numerical' or 'analytic' (default)
- step gradient step size, only valid when type is numerical
- kws other keyword parameters

Returns list of configuration

```
get_opt_results(outfile='dakota.out', rtype='dict')
    extract optimized results from dakota output
```

Parameters

- outfile file name of dakota output file, 'dakota.out' by default
- rtype type of returned results, 'dict' or 'list', 'dict' by default

Returns by default return a dict of optimized results with each item of the format like "x1":0.1, etc., or if rtype='list', return a list of values, when the keys are ascend sorted.

Example

```
>>> opt_vars = get_opt_results(outfile='flame_oc.out', rtype='dict'):
>>> print(opt_vars)
{'x2': 0.0020782814353, 'x1': -0.0017913264033}
>>> opt_vars = get_opt_results(outfile='flame_oc.out', rtype='list'):
>>> print(opt_vars)
[-0.0017913264033, 0.0020782814353]
```

Submodules

genopt.dakopt module

General optimization module by utilizing DAKOTA

• orbit correction: DakotaOC

Tong Zhang <zhangt@frib.msu.edu>

2016-10-23 14:26:13 PM EDT

class DakotaBase(**kws)

Bases: object

Base class for general optimization, initialized parameters: valid keyword parameters:

- •workdir: root dir for dakota input/output files, the defualt one should be created in /tmp, or define some dir path
- •dakexec: full path of dakota executable, the default one should be dakota, or define the full path
- •dakhead: prefixed name for input/output files of dakota, the default one is dakota
- keep: if keep the working directory (i.e. defined by workdir), default is False

dakexec

dakhead

keep

workdir

 $\begin{array}{lll} \textbf{class DakotaOC}(lat_file=None, & elem_bpm=None, & elem_cor=None, & elem_hcor=None, & elem_vcor=None, \\ & ref_x0=None, ref_y0=None, ref_flag=None, model=None, optdriver=None, **kws) \\ & \textbf{Bases: genopt.dakopt.DakotaBase} \end{array}$

Dakota optimization class with orbit correction driver

Parameters

- lat_file lattice file
- **elem_bpm** list of element indice of BPMs
- **elem_cor** list of element indice of correctors, always folders of 2
- elem_hcor list of element indice of horizontal correctors
- elem_vcor list of element indice of vertical correctors
- ref_x0 reference orbit in x, list of BPM readings
- ref_y0 reference orbit in y, list of BPM readings
- ref_flag string flag for objective functions:

1. "x":
$$\sum \Delta x^2$$
, $\Delta x = x - x_0$;

2. "y":
$$\sum \Delta y^2$$
, $\Delta y = y - y_0$;

3. "xy":
$$\sum \Delta x^2 + \sum \Delta y^2$$
.

- model simulation model, 'flame' or 'impact'
- optdriver analysis driver for optimization, 'flamedriver_oc' by default
- kws -

keywords parameters for additional usage, defined in DakotaBase class valid keys:

- workdir: root dir for dakota input/output files, the defualt one should be created in /tmp, or define some dir path
- dakexec: full path of dakota executable, the default one should be dakota, or define the full path
- dakhead: prefixed name for input/output files of dakota, the default one is dakota
- keep: if keep the working directory (i.e. defined by workdir), default is False

bpms

create_machine(lat_file)

create machine instance with model configuration

- •setup _machine
- setup _elem_bpm, _elem_cor or (_elem_hcor and _elem_vcor)

gen_dakota_input(infile=None, debug=False)

generate dakota input file

Parameters

- infile dakota input filename
- **debug** if True, generate a simple test input file

get_all_bpms()

get list of all valid bpms indices

Returns a list of bpm indices

Example

```
>>> dakoc = Dakota0C('test/test.lat')
>>> print(dakoc.get_all_bpms())
```

get_all_cors(type=None)

get list of all valid correctors indices

Parameters type – define corrector type, 'h': horizontal, 'v': vertical, if not defined, return all correctors

Returns a list of corrector indices

Example

```
>>> dakoc = DakotaOC('test/test.lat')
>>> print(dakoc.get_all_cors())
```

get_elem_by_name(name)

get list of element(s) by name(s)

Parameters name – tuple or list of name(s)

Returns list of element indices

Example

```
>>> dakoc = DakotaOC('test/test.lat')
>>> names = ('LS1_CA01:BPM_D1144', 'LS1_WA01:BPM_D1155')
>>> idx = dakoc.get_elem_by_name(names)
>>> print(idx)
[18, 31]
```

get_elem_by_type(type)

get list of element(s) by type

Parameters type – string name of element type

Returns list of element indices

Example

```
>>> dakoc = DakotaOC('test/test.lat')
>>> type = 'bpm'
>>> idx = dakoc.get_elem_by_type(type)
>>> print(idx)
```

get_machine()

get flame machine object for potential usage

Returns flame machine object or None

get_opt_latfile(outfile='out.lat')

get optimized lattice file for potential next usage, run() or simple_run() should be evoked first to get the optimized results.

Parameters outfile – file name for generated lattice file

Returns lattice file name or None if failed

```
get_opt_results(outfile=None, rtype='dict', label='plain')
```

extract optimized results from dakota output

- outfile file name of dakota output file, 'dakota.out' by default
- rtype type of returned results, 'dict' or 'list', 'dict' by default
- label label types for returned variables, only valid when rtype 'dict', 'plain' by default:
 - 'plain': variable labels with format of x1, x2, y1, y2, etc. e.g. {'x1': v1,'y1': v2}

Note: The fancy option will make re-configuring flame machine in a more convenient way, such as:

```
>>> opt_cors = get_opt_results(label='fancy')
>>> for k,v in opt_cors.items():
>>> m.reconfigure(v['id'], v['config'])
>>> # here m is an instance of flame.Machine class
>>>
```

Returns by default return a dict of optimized results with each item of the format like "x1":0.1 or more fancy format by set label with 'fancy', etc., if rtype='list', return a list of values, when the keys are ascend sorted.

Example

```
>>> opt_vars = get_optresults(outfile='flame_oc.out', rtype='dict'):
>>> print(opt_vars)
{'x2': 0.0020782814353, 'x1': -0.0017913264033}
>>> opt_vars = get_optresults(outfile='flame_oc.out', rtype='list'):
>>> print(opt_vars)
[-0.0017913264033, 0.0020782814353]
```

get_orbit(idx=None, val=None, outfile=None)

calculate the orbit with given configurations

Parameters

- idx (idx_hcor, idx_vcor), tuple of list of indices of h/v cors
- val values for each correctos, h/v
- outfile filename to save the data

Returns tuple of zpos, env_x, env_y, machine

hcors

latfile

optdriver

Parameters

• outfile - output file of dakota

```
• figsize – figure size, (h, w)
```

• **dpi** – figure dpi

ref_flag

ref_x0

ref_y0

run(mpi=False, np=None)

run optimization

Parameters

- mpi if True, run DAKOTA in parallel mode, False by default
- np number of processes to use, only valid when mpi is True

set_bpms(bpm=None)

set BPMs

Parameters bpm – list of bpm indices, if None, use all BPMs

```
set_cors(cor=None, hcor=None, vcor=None)
```

set correctors, if cor, hcor and vcor are None, use all correctors if cor is not None, use cor, ignore hcor and vcor

Parameters

- **cor** list of corrector indices, hcor, vcor,...
- hcor list of horizontal corrector indices
- vcor list of vertical corrector indices

set_environ(environ=None)

setup environment block, that is setup oc_environ

Parameters environ – DakotaEnviron object, automatically setup if not defined

```
set_interface(interface=None, **kws)
```

setup interface block, that is setup oc_interface should be ready to invoke after set_cors and set_bpms

Parameters interface – DakotaInterface object, automatically setup if not defined

```
set_method(method=None)
```

setup method block, that is setup oc_method

Parameters method - DakotaMethod object, automatically setup if not defined

```
set_model(model=None, **kws)
```

setup model block, that is setup oc_model

Parameters model – DakotaModel object, automatically setup if not defined

```
set_ref_x0(ref_arr=None)
```

set reference orbit in x, if not set, use 0s

Parameters ref_arr – array of reference orbit values size should be the same number as selected BPMs

```
set_ref_y0(ref_arr=None)
```

set reference orbit in y, if not set, use 0s

Parameters ref_arr – array of reference orbit values size should be the same number as selected BPMs

set_responses(responses=None, **kws)
setup responses block, that is setup oc_responses

Parameters responses - DakotaResponses object, automatically setup if not defined

set_variables(plist=None, initial=0.0001, lower=-0.01, upper=0.01)

setup variables block, that is setup oc_variables should be ready to invoke after set_cors()

Parameters

- plist list of defined parameters (DakotaParam object), automatically setup if not defined
- initial initial values for all variables, only valid when plist is None
- lower lower bound for all variables, only valid when plist is None
- upper upper bound for all variables, only valid when plist is None

simple_run(method='cg', mpi=None, np=None, **kws)

run optimization after set_bpms() and set_cors(), by using default configuration and make full use of computing resources.

Parameters

- method optimization method, 'cg', 'ps', 'cg' by default
- mpi if True, run DAKOTA in parallel mode, False by default
- np number of processes to use, only valid when mpi is True
- **kws** keyword parameters valid keys:
 - step: gradient step, 1e-6 by default
 - iternum: max iteration number, 20 by default
 - evalnum: max function evaulation number, 1000 by default

vcors

test_dakotaoc1()

test_dakotaoc2()

genopt.dakutils module

module contains utilities:

- generate dakota input files
- extract data from output files

Tong Zhang <zhangt@frib.msu.edu>

2016-10-17 09:19:25 AM EDT

class DakotaEnviron(tabfile=None, **kws)

Bases: object

create datako environment for environment block

Parameters

- tabfile tabular file name, by default not save tabular data
- kws other keyword parameters

Example

```
>>> # default
>>> oc_environ = DakotaEnviron()
>>> print oc_environ.get_config()
[]
>>> # define name of tabular file
>>> oc_environ = DakotaEnviron(tabfile='tmp.dat')
>>> print oc_environ.get_config()
['tabular_data', " tabular_data_file 'tmp.dat'"]
>>>
```

get_config(rtype='list')

get responses configuration for dakota input block

Parameters rtype – 'list' or 'string'

Returns dakota responses input

class DakotaInput(**kws)

Bases: object

template of dakota input file, field could be overriden by providing additional keyword arguments,

Parameters kws – keyword arguments, valid keys are dakota directives

Example

set_template(name='oc')

write(infile=None)

write all the input into file, as dakota input file

Parameters infile – fullname of input file, if not defined, infile will be assigned as 'dakota.in' in current working directory

class DakotaInterface(mode='fork', driver='flamedriver_oc', latfile=None, bpms=None, hcors=None, vcors=None, ref_x0=None, ref_y0=None, ref_flag=None, **kws)

Bases: object

create dakota interface for interface block

Parameters

- mode 'fork' or 'direct' (future usage)
- driver analysis driver, external ('fork') executable file internal ('direct') executable file
- latfile file name of (flame) lattice file
- **bpms** array of selected BPMs' id
- hcors array of selected horizontal (x) correctors' id
- vcors array of selected vertical (y) correctors' id
- ref_x0 array of BPM readings for reference orbit in x, if not defined, use 0s
- ref_y0 array of BPM readings for reference orbit in y, if not defined, use 0s

• ref_flag – string flag for objective functions:

```
1. "x": \sum \Delta x^2, \Delta x = x - x_0;
```

2. "y":
$$\sum \Delta y^2$$
, $\Delta y = y - y_0$;

3. "xy":
$$\sum \Delta x^2 + \sum \Delta y^2$$
.

• kws – keyword parameters, valid keys: e.g.: * deactivate, possible value: 'active_set_vector'

Note: mode should be set to be 'direct' when the analysis drivers are built with dakota library, presently, 'fork' is used.

Example

```
>>> # for orbit correction
>>> bpms = [1,2,3] # just for demonstration
>>> hcors, vcors = [1,3,5], [2,4,6]
>>> latfile = 'test.lat'
>>> oc_interface = DakotaInterface(mode='fork',
                                       driver='flamedriver_oc',
                                       latfile=latfile
>>>
                                       bpms=bpms, hcors=hcors, vcors=vcors,
>>>
                                       ref_x0=None, ref_y0=None,
>>>
                                       ref_flag=None,
                                       deactivate='active_set_vector')
>>> # add extra configurations
>>> oc_interface.set_extra(p1='v1', p2='v2')
>>> # get configuration
>>> config_str = oc_interface.get_config()
```

bpms

```
driver
```

```
get_config(rtype='list')
```

get interface configuration for dakota input block

Parameters rtype - 'list' or 'string'

Returns dakota interface input

hcors

latfile

mode

ref_flag

ref_x0

ref_y0

set_extra(**kws)

add extra configurations

vcors

class DakotaMethod(*method*='*cg*', *iternum*=20, *tolerance*=0.0001, **kws)

Bases: object

create dakota method for method block

Parameters

- method method name, 'cg' by default, all possible choices: 'cg', 'ps'
- iternum max iteration number, 20 by default
- tolerance convergence tolerance, 1e-4 by default
- kws other keyword parameters

```
Example
>>> oc_method = DakotaMethod()
 >>> print oc_method.get_config()
['conmin_frcg', ' convergence_tolerance 0.0001', ' max_iterations 20']
>>> # define method with pattern search
>>> oc_method = DakotaMethod(method='ps')
 >>> print oc_method.get_config()
['coliny_pattern_search', 'contraction_factor 0.75', 'max_function_evaluations 500', 'solution_accuracy 0.0001', 'exploratory_moves basic_pattern', 'threshold_delta 0.0001', 'initial_delta 0.5', 'max_iterations 100']
>>> # modify options of pattern search method
>>> oc_method = DakotaMethod(method='ps', max_iterations=200, contraction_factor=0.8)
 >>> print oc_method.get_config()
['coliny_pattern_search', 'contraction_factor 0.8', 'max_function_evaluations 500', 'solution_accuracy 0.0001', 'exploratory_moves basic_pattern', 'threshold_delta 0.0001', 'initial_delta 0.5', 'max_iterations 200']
>>> # conmin_frcg method
>>> oc_method = DakotaMethod(method='cg')
>>> print oc_method.get_config()
['conmin_frcg', ' convergence_tolerance 0.0001', ' max_iterations 20']
>>> # modify options
>>> oc_method = DakotaMethod(method='cg', max_iterations=100)
>>> print oc_method.get_config()
['conmin_frcg', ' convergence_tolerance 0.0001', ' max_iterations 100']
get_config(rtype='list')
        get method configuration for dakota input block
              Parameters rtype – 'list' or 'string'
              Returns dakota method input
get_default_method(method)
```

get default configuration of some method

Parameters method - method name, 'cg' or 'ps'

Returns dict of configuration

method(method)

return method configuration

Parameters method - method stirng name, 'cg' or 'ps'

Returns list of method configuration

```
class DakotaModel(**kws)
```

Bases: object

create dakota model for model block

get_config(rtype='list')

get model configuration for dakota input block

Parameters rtype – 'list' or 'string'

Returns dakota model input

class DakotaParam(label, initial=0.0, lower=-10000000000.0, upper=10000000000.0)

Bases: object

create dakota variable for variables block

Parameters

- label string to represent itself, e.g. x001, it is recommended to annotate the number with the format of %03d, i.e. 1 --> 001, 10 --> 010, 100 --> 100, etc.
- initial initial value, 0.0 by default
- lower lower bound, -1.0e10 by default
- upper upper bound, 1.0e10 by default

initial

label

lower

upper

class DakotaResponses(nfunc=1, gradient=None, hessian=None, **kws)

Bases: object

create dakota responses for responses block

Parameters

- **nfunc** num of objective functions
- gradient gradient type: 'analytic' or 'numerical'
- **hessian** hessian configuration
- **kws** keyword parameters for gradients and hessians valid keys: any available for responses among which key name of 'grad' is for gradients configuration, the value should be a dict (future)

Example

```
>>> # default responses
>>> response = DakotaResponses()
>>> print response.get_config()
['num_objective_functions = 1', 'no_gradients', 'no_hessians']
>>> # responses with analytic gradients:
>>> response = DakotaResponses(gradient='analytic')
>>> print response.get_config()
['num_objective_functions = 1', 'analytic_gradients', 'no_hessians']
>>> # responses with numerical gradients, default configuration:
>>> oc_responses = DakotaResponses(gradient='numerical')
>>> print oc_responses.get_config()
['num_objective_functions = 1', 'numerical_gradients', ' method_source dakota',
' interval_type forward', ' fd_gradient_step_size 1e-06', 'no_hessians']
>>> # responses with numerical gradients, define step:
>>> oc_responses = DakotaResponses(gradient='numerical', step=2.0e-7)
>>> print oc_responses.get_config()
['num_objective_functions = 1', 'numerical_gradients', ' method_source dakota', ' interval_type forward', ' fd_gradient_step_size 2e-07', 'no_hessians']
>>> # given other keyword parameters:
>>> oc_responses = DakotaResponses(gradient='numerical', step=2.0e-7, k1='v1', k2='v2')
>>> print oc_responses.get_config()
['num_objective_functions = 1', 'numerical_gradients', ' method_source dakota',
' interval_type forward', ' fd_gradient_step_size 2e-07', 'no_hessians', 'k2 = v2', 'k1 = v1']
```

get_config(rtype='list')

get responses configuration for dakota input block

Parameters rtype - 'list' or 'string'

Returns dakota responses input

```
gradients(type=None, step=1e-06, **kws)
generate gradients configuration
```

Parameters

- type 'numerical' or 'analytic' (default)
- **step** gradient step size, only valid when type is numerical
- kws other keyword parameters

Returns list of configuration

```
generate_latfile(machine, latfile='out.lat')
```

Generate lattice file for the usage of FLAME code

Parameters

- machine flame machine object
- latfile file name for generated lattice file, 'out.lat' by default

Returns None if failed to generate lattice file, or the out file name

Example

```
>>> from flame import Machine
>>> latfile = 'test.lat'
>>> m = Machine(open(latfile))
>>> outfile1 = generate_latfile(m, 'out1.lat')
>>> m.reconfigure(80, {'theta_x': 0.1})
>>> outfile2 = generate_latfile(m, 'out2.lat')
>>>
```

Warning: To get element configuration only by m.conf(i) method, where m is flame. Machine object, i is element index, when some re-configuring operation is done, m.conf(i) will be update, but m.conf()["elements"] remains with the initial value.

```
get_opt_results(outfile='dakota.out', rtype='dict')
    extract optimized results from dakota output
```

Parameters

- outfile file name of dakota output file, 'dakota.out' by default
- rtype type of returned results, 'dict' or 'list', 'dict' by default

Returns by default return a dict of optimized results with each item of the format like "x1":0.1, etc., or if rtype='list', return a list of values, when the keys are ascend sorted.

Example

```
>>> opt_vars = get_opt_results(outfile='flame_oc.out', rtype='dict'):
>>> print(opt_vars)
{'x2': 0.0020782814353, 'x1': -0.0017913264033}
>>> opt_vars = get_opt_results(outfile='flame_oc.out', rtype='list'):
>>> print(opt_vars)
[-0.0017913264033, 0.0020782814353]
```

random_string(length=8)

generate random string with given length

Parameters length – string length, 8 by default **Returns** random strings with defined length

test_dakotaenviron()
test_dakotainput()
test_dakotainterface()
test_dakotamethod()
test_dakotamodel()
test_dakotaparam()
test_dakotaresponses()
test_get_opt_results()

CHAPTER

FOUR

INDICES AND TABLES

- genindex
- modindex
- search

PYTHON MODULE INDEX

g

genopt, 15
genopt.dakopt, 25
genopt.dakutils, 30

a	end	tac	Do	cu	me	nta	tio	n
---	-----	-----	----	----	----	-----	-----	---

40 Python Module Index

В	get_config() (DakotaResponses method), 24, 34
bpms (DakotaInterface attribute), 22, 32	get_default_method() (DakotaMethod method), 23,
bpms (DakotaOC attribute), 17, 26	33
1 (get_elem_by_name() (DakotaOC method), 18, 27
C	get_elem_by_type() (DakotaOC method), 18, 27
create_machine() (DakotaOC method), 17, 26	get_machine() (DakotaOC method), 18, 27 get_opt_latfile() (DakotaOC method), 18, 27
D	get_opt_results() (DakotaOC method), 18, 27 get_opt_results() (in module genopt), 25
dakexec (DakotaBase attribute), 16, 25	get_opt_results() (in module genopt.dakutils), 35
dakhead (DakotaBase attribute), 16, 25	get_orbit() (DakotaOC method), 19, 28
DakotaBase (class in genopt), 16	gradients() (DakotaResponses method), 24, 35
DakotaBase (class in genopt.dakopt), 25	8
DakotaEnviron (class in genopt), 21	H
DakotaEnviron (class in genopt.dakutils), 30	hears (DakataIntarface attribute) 22 32
DakotaInput (class in genopt), 15	hcors (DakotaInterface attribute), 22, 32
DakotaInput (class in genopt.dakutils), 31	hcors (DakotaOC attribute), 19, 28
DakotaInterface (class in genopt), 21	1
DakotaInterface (class in genopt.dakutils), 31	· · · · 1/D 1 · · D · · · · · · 1 · ·) 1/ 0/
DakotaMethod (class in genopt), 23	initial (DakotaParam attribute), 16, 34
DakotaMethod (class in genopt.dakutils), 32	K
DakotaModel (class in genopt), 24	
DakotaModel (class in genopt.dakutils), 33	keep (DakotaBase attribute), 16, 25
DakotaOC (class in genopt), 16	1
DakotaOC (class in genopt.dakopt), 25	L
DakotaParam (class in genopt), 16	label (DakotaParam attribute), 16, 34
DakotaParam (class in genopt.dakutils), 33	latfile (DakotaInterface attribute), 22, 32
DakotaResponses (class in genopt), 24	latfile (DakotaOC attribute), 19, 28
DakotaResponses (class in genopt.dakutils), 34	lower (DakotaParam attribute), 16, 34
driver (DakotaInterface attribute), 22, 32	A 4
<u> </u>	M
G	method() (DakotaMethod method), 23, 33
gen_dakota_input() (DakotaOC method), 17, 26	mode (DakotaInterface attribute), 22, 32
generate_latfile() (in module genopt.dakutils), 35	
genopt (module), 15	0
genopt.dakopt (module), 25	optdriver (DakotaOC attribute), 19, 28
genopt.dakutils (module), 30	
get_all_bpms() (DakotaOC method), 17, 26	P
get_all_cors() (DakotaOC method), 17, 27	plot() (DakotaOC method), 19, 28
get_config() (DakotaEnviron method), 21, 31	plotty (Dakotaoe litetiou), 17, 20
get_config() (DakotaInterface method), 22, 32	R
get_config() (DakotaMethod method), 23, 33	
get_config() (DakotaModel method), 24, 33	random_string() (in module genopt.dakutils), 35

```
ref flag (DakotaInterface attribute), 22, 32
ref_flag (DakotaOC attribute), 19, 29
ref x0 (DakotaInterface attribute), 23, 32
ref_x0 (DakotaOC attribute), 19, 29
ref_y0 (DakotaInterface attribute), 23, 32
ref_y0 (DakotaOC attribute), 19, 29
run() (DakotaOC method), 19, 29
S
set_bpms() (DakotaOC method), 20, 29
set cors() (DakotaOC method), 20, 29
set_environ() (DakotaOC method), 20, 29
set_extra() (DakotaInterface method), 23, 32
set_interface() (DakotaOC method), 20, 29
set_method() (DakotaOC method), 20, 29
set model() (DakotaOC method), 20, 29
set_ref_x0() (DakotaOC method), 20, 29
set_ref_y0() (DakotaOC method), 20, 29
set_responses() (DakotaOC method), 20, 30
set_template() (DakotaInput method), 16, 31
set_variables() (DakotaOC method), 20, 30
simple_run() (DakotaOC method), 21, 30
test_dakotaenviron() (in module genopt.dakutils), 36
test_dakotainput() (in module genopt.dakutils), 36
test_dakotainterface() (in module genopt.dakutils),
test dakotamethod() (in module genopt.dakutils), 36
test_dakotamodel() (in module genopt.dakutils), 36
test dakotaoc1() (in module genopt.dakopt), 30
test_dakotaoc2() (in module genopt.dakopt), 30
test_dakotaparam() (in module genopt.dakutils), 36
test_dakotaresponses() (in module genopt.dakutils),
test_get_opt_results() (in module genopt.dakutils),
U
upper (DakotaParam attribute), 16, 34
V
vcors (DakotaInterface attribute), 23, 32
vcors (DakotaOC attribute), 21, 30
W
workdir (DakotaBase attribute), 16, 25
write() (DakotaInput method), 16, 31
```

42 Index