
pydaddy
Release 1.0.0

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Apr 28, 2022

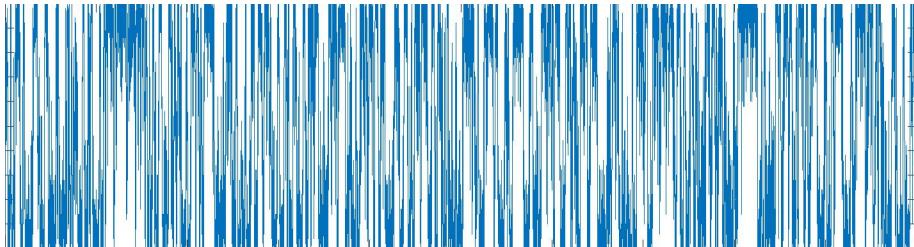
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pydaddy : a package to derive governing stochastic differential equations from timeseries data.

**CHAPTER
ONE**

PYDADDY



1.1 pydaddy

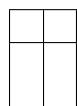
A package to derive an SDE equation form the data.

pydaddy is a python package implementing a data driven SDE method. pydaddy provides an interface which takes time series data as input, runs the analysis and returns an output object through which data and analysed results can be systematically visualized and saved.

1.2 How it works

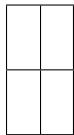
Suppose $m(t)$ is a SDE time series data.

The the package calculates the deterministic (drift) and stochastic (diffusion) component of dm/dt .



This data driven SDE method is based on the assumption that the noise in the time series is uncorrelated and Gaussian in nature, with zero mean and unit variance.

pydaddy extracts the noise form the data and checks if it holds true to its assumptions.



1.3 Features

- Simple one line execution of analysis.
- Produces intuitive visuals of data and obtained result.
- Supports time series data with both scalar and vector order parameters.
- Generates interactive, drift and diffusion sliders for user defined timescale range or list.
- Diagnostics help understand change in order of drift and diffusion with varying timescale.

1.4 Requirements

Python:

- python version >=3.5 , <=3.8

Packages dependencies:

- statsmodels, version 0.11.1
- matplotlib, version 3.2.2
- scipy, version 1.5.2
- numpy, version 1.19.1
- seaborn, version 0.10.1
- tqdm, version 4.48.2
- plotly, version 4.14.3

1.5 Installation

pydaddy is available both on pypi and anaconda cloud, which offers two recommended methods of installations, using pip python package manager or anaconda package manager.

Note: To run the example notebooks on your system after installing the package, please download the notebook files to your PC from the [github](#) repo.

Alternately, its recommended to simply clone or download git the repository.

1.5.1 Installing via conda

Conda is a package manager that handles sourcing of all dependencies in a relatively straight-forward, cross-platform manner. pydaddy is available on the tee-lab channel.

Important: Make sure you have anaconda or miniconda installed and have activated the conda default (base) environment using `conda activate base`, before proceeding.

Install pydaddy package with all its dependencies using conda simply requires executing the command

```
conda install -c tee-lab pydaddy
```

To install pydaddy in a clean virtual environment use

```
conda create --name MY_ENV_NAME -c tee-lab pydaddy
```

Replace `MY_ENV_NAME` with your desired name for environment.

You can now access pydaddy package by activating the newly created environment, `conda activate MY_ENV_NAME`

1.5.2 Installing via pip

To install using pip package manager, run:

```
python -m pip install pydaddy
```

1.5.3 Manual installation

Installing Latest unreleased version

Without git, you will need to download the zip-file of the code, extract it and follow the above instructions.

[Click here to download source code zip file.](#)

This method is not recommended unless you experience problems with conda or pip. To install using setuptools, download the source code manually and run `python setup.py install` from the terminal.

This will install the package in your current environment (if you are working in any environment).

If you have git installed, you can clone the repo and install using the following commands.

```
$ git clone https://github.com/tee-lab/pydaddy.git  
$ cd pydaddy  
$ python setup.py install
```

Important: Without git, you will need to download the zip-file of the code, extract it and follow the above instructions.

[Click here to download source code zip file.](#)

1.6 Usage

The time series data is given as input to the `Characterize` method along with all other optional parameters.

Show `pydaddy.Characterize` documentation

Characterize

```
class pydaddy.Characterize(data, t=1.0, Dt=1, dt=1, bins=None, inc=None, inc_x=None,
                            inc_y=None, slider_timescales=None, n_trials=1,
                            show_summary=True, drift_threshold=None, diff_threshold=None,
                            drift_degree=5, diff_degree=5, drift_alpha=0, diff_alpha=0,
                            fit_functions=False, **kwargs)
```

Bases: `object`

Analyse a time series data and get drift and diffusion plots.

Parameters

- `data` (`list`) – time series data to be analysed, `data = [x]` for scalar data and `data = [x1, x2]` for vector where `x, x1` and `x2` are of `numpy.array` object type
- `t` (`float, array, optional(default=1.0)`) – float if its time increment between observation
`numpy.array` if time stamp of time series
- `Dt` (`int, 'auto', optional(default='auto')`) – time scale for drift
if ‘auto’ time scale is decided based of drift order.
- `dt` (`int, optional(default=1)`) – time scale for diffusion
- `inc` (`float, optional(default=0.01)`) – increment in order parameter for scalar data
- `inc_x` (`float, optional(default=0.1)`) – increment in order parameter for vector data `x1`
- `inc_y` (`float, optional(default=0.1)`) – increment in order parameter for vector data `x2`
- `fft` (`bool, optional(default=True)`) – if true use fft method to calculate autocorrelation else, use standard method
- `slider_timescales` (`list, optional(default=None)`) – List of timescale values to include in slider.
- `n_trials` (`int, optional(default=1)`) – Number of trials, concatenated time-series of multiple trials is used.
- `show_summary` (`bool, optional(default=True)`) – print data summary and show summary chart.
- `**kwargs` – all the parameters for inherited methods.

Returns `output` – object to access the analysed data, parameters, plots and save them.

Return type `pydaddy.output.Output`

1.6.1 Key parameters

data: list time series data to be analysed, data = [x] for scalar data and data = [x1, x2] for vector where x, x1 and x2 are of numpy.array types.

t: array or float float if its time increment between observation numpy.array if time stamp of time series

See doc strings or code documentation for more information.

1.6.2 Example using sample data set

See *Data set description* for more information about the datasets.

```
import pydaddy
#load data
data, t = pydaddy.load_sample_dataset('model-data-vector-ternary')
# Analyse
ddsde = pydaddy.Characterize(data,t)

# Show drift slider plot
ddsde.drift()
# Show diffusion slider plot
ddsde.diffusion()
# Show timeseries plot
ddsde.timeseries()
# Show histograms
ddsde.histograms()
# Show all inputed, calculated and assumed parameters of the analysis
ddsde.parameters()
# Export data to disk
ddsde.export_data()
```

Characterize returns an output object in which all analysed results are stored. Results can be visualised or stored by calling appropriate functions:

Show pydaddy.output.output documentation

pydaddy.output.output

pydaddy.output

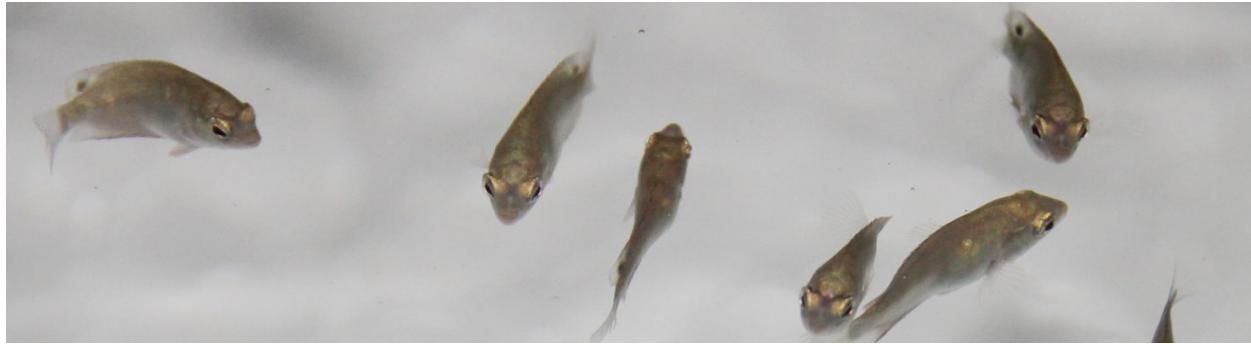
alias of <module ‘pydaddy.output’ from ‘/home/docs/checkouts/readthedocs.org/user_builds/pyddsde/checkouts/latest/pydadd

- **summary()**: show summary
- **drift()** : drift slider plot
- **diffusion()** : diffusion slider plot
- **timeseries()**: time series plot
- **histograms()** : histogram plots
- **noise_characteristics()**: noise characteristics plots
- **visualise(timescale)**: drift and diffusion plots for a timescale
- **diagnostics()**: diagnostics plots

- `data(timescale)`: get drift and diffusion data for a timescale
- `export_data()`: Save data as csv files and mat files
- `plot_data(data)`: plot data on a 3d axis

For more examples see [this](#) notebook.

1.7 Motivation



This project is motivated by the study of group behaviour dynamics of animals, especially schooling fish. The sample data distributed along with this package is from experiments conducted by [TEElab, IISc](#).

1.8 Data set description

pydaddy has six data set included along with the package which can be loaded using `load_sample_dataset(dataset_name)` function

Show `pydaddy.load_sample_dataset` documentation

pydaddy.load_sample_dataset

class pydaddy.load_sample_dataset(name)

Bases:

Load sample data set provided.

Available data sets:

‘fish-data-etroplus’

‘model-data-scalar-pairwise’

‘model-data-scalar-ternary’

‘model-data-vector-pairwise’

‘model-data-vector-ternary’

Parameters `name (str)` – name of the data set

Returns

- `data (list)` – timeseries data

- **t** (*float, array*) – timescale
-

1.8.1 Experiment data (from experimentation or from observations)

fish-data-ectropus: A data from experiment conducted with a group of 30 fish, in which the group polarity in x and y directions are recorded every 0.12 seconds.

Source

- The fish data is a part of the work done in the *Noise-Induced Schooling of Fish*
-

1.8.2 Simulation data

A synthetic data set obtained from the simulation of fish interactions.

model-data-vector-pairwise : Pairwise interaction of fish simulated in two dimension.

model-data-vector-ternary: Ternary interaction of fish simulated in two dimension.

model-data-scalar-pairwise: Pairwise interaction of fish simulated in single dimension.

model-data-scalar-ternary: Ternary interaction of fish simulated in single dimension.

Source

- The simulation method is inspired from the work done in *Noise-induced Effects in Collective Dynamics and Inferring Local Interactions from Data*
-

1.9 Glossary

drift First jump moment

diffusion second jump moment

1.10 Acknowledgement

1.11 Licence

Distributed under **GNU General Public License v3.0**. See `Licence.txt` for more information.

1.12 Reference

- [1] Noise-induced Effects in Collective Dynamics and Inferring Local Interactions from Data [Preprint](#) [Github](#)
- [2] Noise-Induced Schooling of Fish [Preprint](#) [Github](#)

**CHAPTER
TWO**

CODE EXAMPLES

2.1 Code Examples

Example Codes will be here

INSTALLATION GUIDE

3.1 pyddsde Installation Guide

3.1.1 Anaconda

This guide explains creating and installing pyddsde package.

If you don't have Anaconda installed, you can do so from [Anaconda Website](#)

Step 1 : Clone the git repo

Open the terminal in your prefered directory and execute the below command to clone the repo

```
git clone https://github.com/tee-lab/pyFish.git
```

After cloning the terminal should look like this

Step 2 : Change the directory to pyddsde

```
cd pyddsde
```

Typing `ls` should show the following content

Step 3 : Create python environment

```
conda env create -f environment.yml
```

Now, an environment named `pyddsde` should be created

Step 4 : Activate pyddsde environment

```
conda activate pyddsde
```

The (pyddsde) should appear in the terminal.

Step 5 : Install pyddsde

```
python -m pip install .
```

If you see a similar output at the end then the package is successfully installed

You can run the notebook files using jupyter notebook (or jupyter lab)

```
jupyter notebook
```

This should open the notebook application in the browser, click on notebooks folder and open the .ipynb notebook file.

After opening the file, click on the cell and press Shift+Enter to execute that cell and move to the next.

3.1.2 pip

CODE DOCUMENTATION

4.1 Modules

4.1.1 Key Modules

Characterize

```
class pydaddy.Characterize(data, t=1.0, Dt=1, dt=1, bins=None, inc=None, inc_x=None, inc_y=None,
                           slider_timescales=None, n_trials=1, show_summary=True,
                           drift_threshold=None, diff_threshold=None, drift_degree=5, diff_degree=5,
                           drift_alpha=0, diff_alpha=0, fit_functions=False, **kwargs)
```

Bases: object

Analyse a time series data and get drift and diffusion plots.

Parameters

- **data** (*list*) – time series data to be analysed, data = [x] for scalar data and data = [x1, x2] for vector where x, x1 and x2 are of numpy.array object type
- **t** (*float, array, optional(default=1.0)*) – float if its time increment between observation
numpy.array if time stamp of time series
- **Dt** (*int, 'auto', optional(default='auto')*) – time scale for drift
if ‘auto’ time scale is decided based of drift order.
- **dt** (*int, optional(default=1)*) – time scale for difusion
- **inc** (*float, optional(default=0.01)*) – increment in order parameter for scalar data
- **inc_x** (*float, optional(default=0.1)*) – increment in order parameter for vector data x1
- **inc_y** (*float, optional(default=0.1)*) – increment in order parameter for vector data x2
- **fft** (*bool, optional(default=True)*) – if true use fft method to calculate autocorrelation else, use standard method
- **slider_timescales** (*list, optional(default=None)*) – List of timescale values to include in slider.
- **n_trials** (*int, optional(default=1)*) – Number of trials, concatenated timeseries of multiple trials is used.

- **show_summary**(*bool*, *optional (default=True)*) – print data summary and show summary chart.
- ****kwargs** – all the parameters for inherited methods.

Returns **output** – object to access the analysed data, parameters, plots and save them.

Return type *pydaddy.output.Output*

load_sample_dataset

```
class pydaddy.load_sample_dataset(name)
```

Bases:

Load sample data set provided.

Available data sets:

- ‘fish-data-etroplus’
- ‘model-data-scalar-pairwise’
- ‘model-data-scalar-ternary’
- ‘model-data-vector-pairwise’
- ‘model-data-vector-ternary’

Parameters **name** (*str*) – name of the data set

Returns

- **data** (*list*) – timeseries data
- **t** (*float, array*) – timescale

output

```
class pydaddy.output.Output(ddsde, **kwargs)
```

Bases: *pydaddy.preprocessing.Preprocessing*, *pydaddy.visualize.Visualize*

Class to plot and save data and parameters

property A1

property A2

property B11

property B12

property B21

property B22

property F

property G

property _data_avgdiff

```

property _data_avgdiffX
property _data_avgdiffXY
property _data_avgdiffY
property _data_avgdiffYX
property _data_avgdrift
property _data_avgdriftX
property _data_avgdriftY
property _data_diff_ebar
property _data_drift_ebar
_print_function_diagnostics(f, x, y, name, symbol)
_print_function_diagnostics_2d(f, x, y, z, name, symbol)
_update_slider_data(slider_timescales)
autocorrelation()
cross_diffusion(slider_timescales=None, limits=None, polar=False, **plot_text)
    Display diffusion cross correlation slider figure
        Parameters polynomial_order (None or int, default=None) – order of polynomial to fit, if None, no fitting is done.
        Returns opens diffusion slider
        Return type None
data(drift_time_scale=None, diff_time_scale=None)
    Get the drift, diffusion and order parameter data for any timescale the analysis is done.
        Parameters
            • drift_time_scale (int, optional(default=None)) – time-scale of drift data, if None, returns data analysed for given dt
            • diff_time_scale (int, optional(default=None)) – time-scale of diffusion data, if None, returns data analysed for given delta_t
        Returns
            • if vector, [avgdriftX, avgdriftY, avgdiffX, avgdiffY, op_x, op_y]
            • else, [avgdrift, avgdiff, op]
        Return type list
diffusion(slider_timescales=None, limits=None, polar=False, **plot_text)
    Display diffusion slider figure
        Parameters polynomial_order (None or int, default=None) – order of polynomial to fit, if None, no fitting is done.
        Returns opens diffusion slider
        Return type None

```

drift(*limits=None, polar=False, slider_timescales=None, **plot_text*)

Display drift slider figure

Parameters

- **polynomial_order** (*None or int, default=None*) – order of polynomial to fit, if None, no fitting is done.
- ****plot_text** – plots' axis and text label

For scalar analysis *x_lable* : x axis label

y_label : y axis label

For vector analysis *title1* : first plot title

x_label1 : first plot x label

y_label1 : first plot y label

z_label1 : first plot z label

title2 : second plot title

x_label2 : second plot x label

y_label2 : seocnd plot y label

z_label2 : second plot z label

Returns opens drift slider

Return type None

export_data(*filename=None, raw=False*)

Returns a pandas dataframe containing the drift and diffusion values. :param filename: If provided, the data will be saved as a CSV at the given path. Else, a dataframe will be returned. :type filename: str, optional (default=None). :param raw: If True, the raw, the drift and diffusion will be returned as raw unbinned data. Otherwise (default),

drift and diffusion as binwise-average Kramers-Moyal coefficients are returned.

Returns df

Return type Pandas dataframe containing the estimated drift and diffusion coefficients.

fit_diagnostics()

histogram(*kde=False, heatmap=False, dpi=150, title_size=14, label_size=15, tick_size=12, label_pad=8, **plot_text*)

Show histogram polt chart

Parameters

- **kde** (*bool, (default=False)*) – If True, plots kde for histograms
- **dpi** (*int, (default=150)*) – figure resolution
- **title_size** (*int, (default=14)*) – title font size
- **label_size** (*int, (default=15)*) – axis label font size
- **tick_size** (*int, (default=12)*) – axis ticks font size
- **label_pad** (*int, (default=8)*) – axis label padding

- ****plot_text** – plots' axis and title text

For scalar analysis histograms: hist_title : title

hist_xlabel : x label

hist_ylabel : y label

For vector analysis histograms: hist1_title : first histogram title

hist1_xlabel : first histogram x label

hist1_ylabel : first histogram y label

hist2_title : second histogram title

hist2_xlabel : second histogram x label

hist2_ylabel : second histogram y label

hist3_title : third histogram title

hist3_xlabel : third histogram x label

hist3_ylabel : third histogram y label

hist4_title : fourth (3d) histogram title hist4_xlabel : fourth (3d) histogram x label

hist4_ylabel : fourth (3d) histogram y label hist4_zlabel : fourth (3d) histogram z label

Returns histogram chart

Return type matplotlib.pyplot.figure

noise_diagnostics(loc=None)

parameters()

Get all given and assumed parameters used for the analysis

Returns params – all parameters given and assumed used for analysis

Return type dict, json

plot_data(data_in, ax=None, clear=False, title=None, x_label='x', y_label='y', z_label='z', tick_size=12, title_size=16, label_size=14, label_pad=12, legend_label=None, dpi=150)

Plot and visualize vector drift or diffusion data of a 3d axis

Can be used plot multiple data on the same figure and compare by passing the axis of figure.

Parameters

- **data_in** (`numpy.array`) – vector drift or diffusion data to plot
- **ax** (`figure axis, (default=None)`) – If ax is None, a new axis will be created and data will be plotted on it.
- **clear** (`bool, (default=False)`) – if True, clear the figure.
- **title** (`str, (default=None)`) – title of the figure
- **x_label** (`str, (default='x')`) – x-axis label
- **y_label** (`str, (default='y')`) – y-axis label
- **z_label** (`str, (default='z')`) – z-axis label
- **tick_size** (`int, (default=12)`) – axis ticks font size
- **title_size** (`int, (default=16)`) – title font size

- **label_size** (*int, (default=14)*) – axis label font size
- **label_pad** (*int, (default=12)*) – axis label padding
- **legend_label** (*str, (default=None)*) – data legend label
- **dpi** (*int, (default=150)*) – figure resolution

Returns

- **ax** (*3d figure axis*) – axis of the 3d figure.
- **fig** (*matplotlib figure*) – returns figure only if the input ax is None.

release()

Clears the memory, recommended to be used while analysing multiple data files in loop.

Return type None

simulate(*sigma=4, dt=None, T=None, **functions*)

Simulate SDE

Takes drift and diffusion functions as input and simuates the SDE using the analysis parameters.

The drift and diffusion functions given must be callable type functions.

For scalar F and G (drift and diffusion) must take one input and return a number

Parameters

- **sigma** (*float*) – magnitude of the noise eta(t)
- ****functions** – drift and diffusion callable functions

For scalar analysis F : drift function

G : diffusion dunction

For vector analysis A1 : drift X

A2 : drift Y

B11 : diffusion X

B22 : diffusion Y

B12 : diffusion XY

B21 : diffusion YX

Returns

simulated timeseries – [M] if scalar

[Mx, My] is vector

Return type list

Examples

```
# For scalar analysis def drift_function(x):
    return 0.125 * x

def diffusion_function(x): return -(x**2 + 1)

simulated_data = ddsde.simulate(F=drift_function, G=diffusion_function)

# For vector analysis def drift_x(x, y):
    return x*x + y*y * x*y**2

def dirft_y(x, y): return x*y

def diffusion_x(x,y): return x**2 + x*y

def diffusion_y(x,y): return y**2 + x*y

def diffusion_xy(x,y): return 0

def diffusion_yx(x,y): return 0

simulated_data = ddsde.simulate(A1=drift_x, A2=drift_y, B11=diffusion_x, B22=diffusion_y,
    B12=diffusion_xy, B21=diffusion_yx )

summary(start=0, end=1000, kde=True, tick_size=12, title_size=15, label_size=15, label_pad=8, n_ticks=3,
    ret_fig=False, **plot_text)
```

Print summary of data and show summary plots chart

Parameters

- **start** (*int*, (*default=0*)) – starting index, begin plotting timeseries from this point
- **end** (*int*, (*default=1000*)) – end point, plots timeseries till this index
- **kde** (*bool*, (*default=False*)) – if True, plot kde for histograms
- **title_size** (*int*, (*default=15*)) – title font size
- **tick_size** (*int*, (*default=12*)) – axis tick size
- **label_size** (*int*, (*default=15*)) – label font size
- **label_pad** (*int*, (*default=8*)) – axis label padding
- **n_ticks** (*int*, (*default=3*)) – number of axis ticks
- **ret_fig** (*bool*, (*default=True*)) – if True return figure object
- ****plot_text** – plots' title and axis texts

For scalar analysis summary plot: timeseries_title : title of timeseries plot

timeseries_xlabel : x label of timeseries

timeseries_ylabel : y label of timeseries

drift_title : drift plot title

drift_xlabel : drift plot x label

drift_ylabel : drift plot ylabel

diffusion_title : diffusion plot title

```
diffusion_xlabel : diffusion plot x label
diffusion_ylabel : diffusion plot y label

For vector analysis summary plot: timeseries1_title : first timeseries plot title
timeseries1_ylabel : first timeseries plot ylabel
timeseries1_xlabel : first timeseries plot xlabel
timeseries1_legend1 : first timeseries (Mx) legend label
timeseries1_legend2 : first timeseries (My) legend label
timeseries2_title : second timeseries plot title
timeseries2_xlabel : second timeseries plot x label
timeseries2_ylabel : second timeseries plot y label
2dhist1_title : Mx 2d histogram title
2dhist1_xlabel : Mx 2d histogram x label
2dhist1_ylabel : Mx 2d histogram y label
2dhist2_title : My 2d histogram title
2dhist2_xlabel : My 2d histogram x label
2dhist2_ylabel : My 2d histogram y label
2dhist3_title : M 3d histogram title
2dhist3_xlabel : M 2d histogram x label
2dhist3_ylabel : M 2d histogram y label
3dhist_title : 3d histogram title
3dhist_xlabel : 3d histogram x label
3dhist_ylabel : 3d histogram y label
3dhist_zlabel : 3d histogram z label
driftx_title : drift x plot title
driftx_xlabel : drift x plot x label
driftx_ylabel : drift x plot y label
driftx_zlabel : drift x plot z label
drifty_title : drift y plot title
drifty_xlabel : drift y plot x label
drifty_ylabel : drift y plot y label
drifty_zlabel : drift y plot z label
diffusionx_title : diffusion x plot title
diffusionx_xlabel : diffusion x plot x label
diffusionx_ylabel : diffusion x plot y label
diffusionx_zlabel : diffusion x plot z label
diffusiony_title : diffusion y plot title
```

```
diffusiony_xlabel : diffusion y plot x label  
diffusiony_ylabel : diffusion y plot y label  
diffusiony_zlabel : diffusion y plot z label
```

Return type None, or figure

Raises ValueError – If start is greater than end

timeseries(*start=0, end=1000, n_ticks=3, dpi=150, tick_size=12, title_size=14, label_size=14, label_pad=0, **plot_text*)

Show plot of input data

Parameters

- **start** (*int, (default=0)*) – starting index, begin plotting timeseries from this point
- **end** (*int, default=1000*) – end point, plots timeseries till this index
- **n_ticks** (*int, (default=3)*) – number of axis ticks
- **dpi** (*int, (default=150)*) – dpi of the figure
- **title_size** (*int, (default=15)*) – title font size
- **tick_size** (*int, (default=12)*) – axis tick size
- **label_size** (*int, (default=15)*) – label font size
- **label_pad** (*int, (default=8)*) – axis label padding
- ****plot_text** – plots' title and axis texts

For scalar analysis plot: timeseries_title : title

```
timeseries_xlabel : x label  
timeseries_ylabel : y label
```

For vector analysis plot: timeseries1_title : first timeseries plot title

```
timeseries1_xlabel : first timeseries plot x label  
timeseries1_ylabel : first timeseries plot y label  
timeseries2_title : second timeseries plot title  
timeseries2_xlabel : second timeseries plot x label  
timeseries2_ylabel : second timeseries plot y label  
timeseries3_title : third timeseries plot title  
timeseries3_xlabel : third timeseries plot x label  
timeseries3_ylabel : third timeseries plot y label
```

Returns time series plot

Return type matplotlib.pyplot.figure

Raises ValueError – If start is greater than end

visualize(*drift_time_scale=None, diff_time_scale=None*)

Display drift and diffusion plots for a time scale.

Parameters `time_scale` (*int, optional(default=None)*) – timescale for which drift and diffusion plots need to be shown. If None, displays the plots for inputed timescale.

Returns `displays plots`

Return type None

4.1.2 Additional Modules

SDE

`class pydaddy.sde.SDE(**kwargs)`

Bases: `object`

A class to form a basic SDE from data

Parameters

- `X (array_like)` – time series data
- `t_int (float)` – time step in time series
- `Dt (int)` – analysis time step
- `inc (float)` – max increment for binning the data

Returns

- `diff (array_like)` – diffusion in time series
- `drift (array_like)` – drift in time series
- `avgdiff (array_like)` – average diffusion
- `avgdrift (array_like)` – average drift
- *meta private:*

`_diffusion(X, t_int, dt=1)`

Get Diffusion coefficient vector of data

Parameters

- `X (array_like)` – time series data
- `t_int (float)` – time step in time series
- `dt (int)` – diffusion calculation timescale

Returns

`diff` – Diffusion

Return type array.

`_diffusion_from_residual(X, F, t_int, dt=1)`

Get diffusion using residuals about drift function.

Parameters

- `(np.array) (X)` –
- `(float) (t_int)` –
- `(Callable) (F)` –

`_diffusion_x_from_residual(x, y, A1, t_int, dt)`

_diffusion_xy(*x, y, t_int, dt*)

Get cross-correlation coefficients between x and y arrays.

Parameters

- **x** (`numpy.array`) – x data
- **y** (`numpy.array`) – y data
- **t_int** (`float`) – time difference between consecutive observations
- **dt** (`diffusion calculation timescale`) –

Returns `diffusion_xy` – cross-correlation coefficients between x and y data

Return type `numpy.array`

_diffusion_xy_from_residual(*x, y, A1, A2, t_int, dt*)**_diffusion_y_from_residual**(*x, y, A2, t_int, dt*)**_diffusion_yx**(*x, y, t_int, dt*)

Get cross-correlation coefficients between x and y arrays.

Parameters

- **x** (`numpy.array`) – x data
- **y** (`numpy.array`) – y data
- **t_int** (`float`) – time difference between consecutive observations
- **dt** (`diffusion calculation timescale`) –

Returns `diffusion_xy` – cross-correlation coefficients between y and x data

Return type `numpy.array`

_drift(*X, t_int, Dt*)

Get Drift coefficient vector of data.

Parameters

- **X** (`array_like`) – Time Series data
- **t_int** (`float`) – time difference between consecutive observations
- **Dt** (`float`) – drift calculation timescale

Returns `diff` – Diffusion in time series

Return type `array_like`

Notes

Drift is calculated as follows

$$\text{drift} = \frac{x(i + Dt) - x(i)}{tint * Dt}$$

_drift_and_diffusion(*X*, *t_int*, *Dt*, *dt*, *inc*, *drift_threshold*, *drift_degree*, *drift_alpha*, *diff_threshold*, *diff_degree*, *diff_alpha*, *fast_mode*)

Get drift and diffusion coefficients for a given timeseries data

Parameters

- **X** (`numpy.array`) – time series data
- **t_int** (`float`) – time difference between consecutive observations
- **Dt** (`int`) – timescale to calculate drift
- **dt** (`int`) – timescale to calculate diffusion
- **inc** (`float`) – step increments in order parameter
- **drift_threshold** (`float or None`) – threshold to use for fitting drift function. If None, automatic model selection will be used.
- **diff_threshold** (`float or None`) – threshold to use for fitting diffusion function. If None, automatic model selection will be used.

Returns

- —
- **diff** (`array`) – diffusion of the data
- **drift** (`array`) – drift, of the data
- **avgdiff** (`array`) – average diffusion
- **avgdrift** (`array`) – average drift
- **op** (`array`) – order parameter

_isValidRange(*r*)

Checks if the specified range of order parameter is valid range

Parameters **r** (`tuple, list`) – range of order parameter

Returns True if valid, False if not.

Return type bool

_km_coefficient(*order*, *X*, *t_int*)

_order_parameter(*X*, *inc*, *r*)

Get order parameter array for a given range and increments

If range is None or not valid, order parameter array will be generated considering maximum and minimum limits of the data as the range

Parameters

- **X** (`numpy.array`) – data
- **inc** (`float`) – step increments in order parameter
- **r** (`tuple, list`) – range of the order parameter

Returns first element will be the order parameter array second element is the range used

Return type tuple

_residual(*X*, *t_int*, *Dt*, *dt=1*)

Get the residual.

_vector_drift_diff(*x, y, inc_x, inc_y, t_int, Dt, dt, drift_threshold, drift_degree, drift_alpha, diff_threshold, diff_degree, diff_alpha, fast_mode*)

Get average binned drift and diffusion coefficients for given x and y data

Parameters

- **x** (*array_like*) – timeseries x data
- **y** (*array_like*) – timeseries y data
- **inc_x** (*float*) – step increment of order parameter for x
- **inc_y** (*float*) – step increment of order parameter for y
- **Dt** (*int*) – timescale to calculate drift
- **dt** (*int*) – timescale to calculate diffusion

Returns [avgdriftX, avgdriftY, avgdiffX, avgdiffY, avgdiffXY, op_x, op_y]

Return type list

metrics

class pydaddy.metrics.**Metrics**(**kwargs)

Bases: object

Helper/utility module

_R2(*data, op, poly, k, adj=False*)

R-square value between the predicted and expected values

Parameters

- **data** (*array*) – depended variable values, expected values, data
- **op** (*array*) – independent variable values
- **poly** (*numpy.poly1d*) – numpy polynomial fitted object
- **k** (*int*) – degree of the polynomial poly
- **adj** (*bool*) – if True, use R2-adjusted method instead of R2

Returns R2 – R2 or R2-adjusted depending upon ‘adj’ value

Return type float

_R2_adj(*data, op, poly, k*)

Get R-squared adjusted parameter between data and fitted polynomial

Parameters

- **data** (*array*) – depended variable values, expected values, data
- **op** (*array*) – independent variable for which the data is defined
- **poly** (*numpy.poly1d*) – numpy polynomial fitted object
- **k** (*int*) – degree of polynomial

Returns R2-adjusted – R2 adjusted parameter between data and fitted polynomial

Return type float

_closest_time_scale(*time_scale, slider*)

Gives closest matching time scale available from the slider keys.

_combined_data_dict()

Get all drift and diffusion data in dictionary format.

_csv_header(*prefix, file_name*)

Generate headers for CSV file.

_divergence(*a, b*)

Get the divergence between two timeseries data, the divergence returned here is defined as follows: divergence = $0.5 * (\text{KL_divergence}(p,q) + \text{KL_divergence}(q,p))$

The probability density of a and b input timeseries is calculated before finding the divergence.

Parameters

- **a** (array) – observed timeseries data
- **b** (array) – simulated timeseries data

Returns divergence

Return type float

_fit_plane(*x, y, z, order=2*)

Fits n-th order plane to data in the form $z = f(x,y)$ where $f(x,y)$ the best fit equation of plane for the data computed using least square method.

Parameters

- **x** (2D array) – order parameter x
- **y** (2D array) – order parameter y
- **z** (2D array) – derived drift or diffusion data
- **order** (int) – order of the 2D plane to fit

Returns A callable object takes in x and y as inputs and returns $z = f(x,y)$, where $f(x,y)$ is the fitted function of the plane.

Return type `pydaddy.metrics.Plane`

_fit_poly(*x, y, deg*)

Fits polynomial of degree *deg*

Parameters

- **x** (array) – independent variable
- **y** (array) – depended variable
- **deg** (int) – degree of the polynomial

Returns

- **poly** (`numpy.poly1d`) – polynomial object
- **x** (array) – values of x for where y is defined

Notes

The nan values in the input x and y (if any) will be ignored.

_fit_poly_sparse(x, y, deg, threshold=0.05, alpha=0, weights=None)

Fit a polynomial using sparse regression using STLSQ (Sequentially thresholded least-squares) :param x: (np.array) Independent and dependent variables :param y: (np.array) Independent and dependent variables :param deg: (int) Maximum degree of the polynomial :param threshold: (float) Threshold for sparse fit.

_get_data_from_slider(drift_time_scale=None, diff_time_scale=None)

Get drift and diffusion data from slider data dictionary, if key not valid, returns the data corresponding to closest matching one.

_get_data_range(x)

Get range of the values in x, (min(x), max(x)), rounded to 3 decimal places.

_get_num_points(drift_time_scale, diff_time_scale)

_get_stacked_data()

Get a dictionary of all (op_x, op_y, driftX, driftY, diffX, diffY) slider data stacked into numpy arrays.

_interpolate_missing(y, copy=True)

Interpolate missing data

Parameters

- **y** (array) – data with missing (nan) values
- **copy** (bool, optional (default=True)) – if True makes a copy of the input array object

Returns y – interpolated data

Return type array

_is_valid_slider_timescale_list(slider_list)

Checks if the given slider timescale lists contains valid entries

Parameters **slider_list** (list, tuple) – timescales to include in the slider

Returns True if all values are valid, else False

Return type bool

_isnotebook()

_kl_divergence(p, q)

Calculates KL divergence between two probability distributions p and q

Parameters

- **p** (array) – distribution p
- **q** (array) – distribution q

Returns kl_divergence – kl divergence between p and q

Return type float

_make_directory(p, i=1)

Recursively create directory for a given path

Parameters **path** (str) – destination path

Returns `path` – path of created directory, same as input path.

Return type str

_nan_helper(*x*)

Helper function used to handle missing data

Parameters `x` (array) – data

Return type callable function

_remove_nan(*x, y*)

Removes NaN's by deleting the indices where both *x* and *y* have NaN's

Parameters

- `x` (array) – first input
- `y` (array) – second input

Returns *x, y* - with all nan's removed

Return type array

_rms(*x*)

Calculates root mean square error of *x*

Parameters `x` (array) – input

Returns rms – rms error

Return type float

_save_csv(*dir_path, file_name, data, fmt='%.4f, add_headers=True*)

Save data to CSV file.

_stack_slider_data(*d, slider_data, index*)

Stack data from slider dictionary, corresponding to the given index, into columns of numpy array.

_zip_dir(*dir_path*)

Make ZIP file of the exported result.

class pydaddy.metrics.Plane(*coefficients, order*)

Bases: object

Create first or second order plane surfaces.

expr()

fitters

Code for fitting polynomials to data.

class pydaddy.fitters.Poly(*coeffs, degree, stderr*)

Bases: object

class pydaddy.fitters.Poly1D(*coeffs, degree, stderr=None*)

Bases: `pydaddy.fitters.Poly`

A rudimentary 2D polynomial class for polynomials with optional error intervals for coefficients. Returns polynomial objects that can be called or pretty-printed.

```
class pydaddy.fitters.Poly2D(coeffs, degree, stderr=None)
```

Bases: *pydaddy.fitters.Poly*

A rudimentary 2D polynomial class for polynomials with optional error intervals for coefficients. Returns polynomial objects that can be called or pretty-printed.

```
class pydaddy.fitters.PolyFit1D(**kwargs)
```

Bases: *pydaddy.fitters.PolyFitBase*

```
_evaluate_poly(c, x)
```

```
_get_callable_poly(coeffs, stderr)
```

Construct a callable polynomial from a given coefficient array.

```
_get_coeffs()
```

```
_get_poly_dictionary(x)
```

```
class pydaddy.fitters.PolyFit2D(**kwargs)
```

Bases: *pydaddy.fitters.PolyFitBase*

```
_evaluate_poly(c, x)
```

```
_get_callable_poly(coeffs, stderr)
```

```
_get_coeffs()
```

```
_get_poly_dictionary(x)
```

```
class pydaddy.fitters.PolyFitBase(threshold=0, max_degree=5, alpha=0, library=None)
```

Bases: *object*

Fits polynomial to estimated drift and diffusion functions with sparse regression.

```
_evaluate(c, x)
```

```
_evaluate_poly(c, x)
```

```
_get_bic(p, x, y)
```

Compute the BIC for a fitted polynomial with given data *x*, *y*.

```
_get_callable_poly(coeffs, stderr)
```

```
_get_coeffs()
```

```
_get_cv_error(x, y, folds)
```

```
_get_poly_dictionary(x)
```

```
fit(x, y, weights=None)
```

Fit a polynomial using sparse regression using STLSQ (Sequentially thresholded least-squares) :param *x*: Independent variable. Could either be an array (for 1D case) or

a list of two arrays (for 2D case).

Parameters

- **y** (*np.array*) – Dependent variable
- **weights** (*np.array*) – Sample weights for regression. If None (default), simple unweighted ridge regression will be performed.

Returns np.poly1d object for 1D case, Poly2D object for 2D case.

model_selection(*thresholds*, *x*, *y*, *weights=None*, *method='cv'*, *plot=False*)

Automatically choose the best threshold using BIC. :param thresholds: List of thresholds to search over. :param x: Data to be used for parameter tuning. :param y: Data to be used for parameter tuning. :param weights: (Optional) weights for fitting. :param method: {‘bic’, ‘cv’} The metric used for model selection :param plot: If true, plot the model selection curves

tune_and_fit(*x*, *y*, *thresholds=None*, *steps=20*, *plot=False*)

Parameters

- **x** – Data to fit
- **y** – Data to fit
- **thresholds** – List of thresholds to try, will be automatically chosen if None
- **steps** – When auto-choosing thresholds, the number of steps to take in the threshold range.
- **plot** – Whether to plot the cross-validation error curves.

analysis

class pydaddy.analysis.AutoCorrelation(***kwargs*)

Bases: object

This class defines methods to calculate the _autocorrelation function of time series, fit an exponential curve to it and calculate the _autocorrealtion time.

Parameters: fft : bool If True, use fft method (wiener khinchin theorem) to calculate acf.

_acf(*data*, *t_lag*)

Get auto correaltion function for given *data* and lag *t_lag*

Parameters

- **data** (array) – timeseries data
- **t_lag** (int) – maxmum lag

Returns

- **x** (array) – lags
- **c** (array) – correlation values

Notes

If fft flag is set True and no valid fft points are found, the method uses standard formula method to calculate the autocorrealtion function

_acf_fft(*data*, *t_lag*)

Calculates autocorrelation using wiener khinchin theorem.

_act(*X*, *t_lag=1000*)

Get autocorrelation time of X.

```
_ccf(x, y, t_lag)
    " Returns the cross-correlation function between x and y.

_fit_exp(x, y)
    Fits an exponential function of the form a*exp((-1/b)*t) + c
```

Parameters

- **x** (*array*) – x data
- **y** (*array*) – y data

Returns

- **params** (*Tuple (a, b, c) containing the fitted parameters.*)
- **cov** (*Covariance matrix of errors*)

Notes

Reference : `scipy.optimize.curve_fit`

```
_get_autocorr_time(X, t_lag=1000, update=True)
```

Get the autocorrelation time of data X, for the analysis.

```
_nan_acf(data, t_lag)
```

Calculates autocorrelation using the correlation formula, ignoring all points with nan's

```
_nan_ccf(data_x, data_y, t_lag)
```

Calculates cross-correlation using the correlation formula, ignoring all points with nan's

```
class pydaddy.analysis.GaussianTest(**kwargs)
```

Bases: `pydaddy.analysis.UnderlyingNoise`, `pydaddy.metrics.Metrics`, `pydaddy.analysis.AutoCorrelation`

Used to check if the noise is gaussian in nature

```
_get_critical_values(kl_dist)
```

Get critical values of null hypothesis, i.e values at the boundaries of 2.5% and 97.5% of null hypothesis

```
_noise_analysis(X, Dt, dt, t_int, inc, point=0, **kwargs)
```

Check if noise is gaussian

Parameters

- **X** (*array*) – timeseries data
- **Dt** (*int*) – drift timescale
- **inc** (*float*) – increment in order parameter of X
- **point** (*int*) – point at which noise is to be extracted

Returns

- gaussian_noise (*bool*) : True if the noise is gaussian
- noise (*array*) : extracted noise
- kl_dist (*array*) : null hypothesis uses

- **k** (float) : test statistics used for test of hypothesis
- **l_lim** (float) : lower critical limit
- **h_lim** (float) : upper critical limit
- **noise_correlation** (array) : noise autocorrelation

Return type tuple

class pydaddy.analysis.UnderlyingNoise(**kwargs)

Bases: [pydaddy.sde.SDE](#)

Extract noise from time series

_noise(X, bins, avg_drift, inc, t_int, point=0)

Get noise from X at a paticular point

Parameters

- **X** (array) – time series
- **inc** (float) – binning increments
- **point** (float) – point at which noise is to be extracted
- **Dt** (int) – drift time scale
- **t_int** (int) – time difference between consecutive observations

Returns noise extracted from data at given point

Return type array

_noise_vector(X, Y, bins_x, bins_y, avg_drift_x, avg_drift_y, inc_x, inc_y, t_int, point_x=0, point_y=0)

_residual_timeseries(X, bins, avg_drift, t_int)

_residual_timeseries_vector(X, Y, bins_x, bins_y, avg_drift_x, avg_drift_y, t_int)

preprocessing

exception pydaddy.preprocessing.Error

Bases: Exception

Base class for exceptions in this module.

exception pydaddy.preprocessing.InputError(expression, message)

Bases: [pydaddy.preprocessing.Error](#)

Exception raised for errors in the input.

Attributes: expression – input expression in which the error occurred message – explanation of the error

```
class pydaddy.preprocessing.Preprocessing(**kwargs)
Bases: pydaddy.analysis.GaussianTest

    pass


    find_order(x)
        Get expected order by elimination least likely to be values from all possible values. Then decides the order by looking at the R2 values.

    get_o1_o2(x)
        Get o1 and o2 values for r2_adjusted multiple Dt

    o1(x, i=0)
        All possible values of order

    o2(x)
        Least likely values of order

    optimum_timescale(X, M_square, t_int, Dt='auto', max_order=10, t_lag=1000, inc=0.01)
        Get timescale based on observed order of drift

    order(X, M_square, t_int, Dt='auto', dt=1, max_order=10, inc=0.01)
        Find the order of drift and diffusion, and timescale based on drift order.
```

Notes

Time scale = autocorrelation time if drift order is 1, else its auto correlation time.

```
preprocess()

r2_vs_order(op1, op2, avgDrift, avgDiff, max_order)
    Get R2 for different order

r2_vs_order_multi_dt(X, M_square, t_int, dt=1, max_order=10, inc=0.01)
    Get R2 vs order for different Dt

remove_nan(x, y, sample_size=10)
    Removes NaN's by deleting the indices where both x and y have NaN's
```

Parameters

- **x** (array) – first input
- **y** (array) – second input

Returns x, y - with all nan's removed

Return type array

```
remove_outliers(xs, y, quantile=0.01)
```

Remove points corresponding to outliers in y. xs is a list of one or more arrays, indices corresponding to outliers in y will be removed from each array in xs as well.

```
rms_variation(x)
```

Get rms variation of array

```
_timestep(t)
_validate_inputs()
Initialize and validate all inputs.
```

visualize

```
class pydaddy.visualize.Visualize(op_x, op_y, op, autocorrelation_time, **kwargs)
```

Bases: *pydaddy.metrics.Metrics*

Module to visualize and plot analysed data

```
_acf_plot(ax, acf, lags, a, b, c, act, title)
```

```
_acf_plot_multi(ax, acf1, acf2, lags, act1, act2, title=None)
```

```
_histogram3d(x, bins=20, normed=False, color='blue', alpha=1, hold=False, plot_hist=False)
```

Plotting a 3D histogram

Parameters

- **sample** (*array_like*.*）* – The data to be histogrammed. It must be an (N,2) array or data that can be converted to such. The rows of the resulting array are the coordinates of points in a 2 dimensional polytope.
- **bins** (*sequence or int, optional, default: 10.*) – The bin specification:
 - A sequence of arrays describing the bin edges along each dimension.
 - The number of bins for each dimension (*bins = [binx,biny]*)
 - The number of bins for all dimensions (*bins = bins*).
- **normed** (*bool, optional, default: False.*) – If False, returns the number of samples in each bin. If True, returns the bin density *bin_count / sample_count / bin_volume*.
- **color** (*string, matplotlib color arg, default = 'blue'*) –
- **alpha** (*float, optional, default: 1.*) – 0.0 transparent through 1.0 opaque
- **hold** (*boolean, optional, default: False*) –

Returns

- **H** (*ndarray*.*）* – The bidimensional histogram of sample x.
- **edges** (*list*.*）* – A list of 2 arrays describing the bin edges for each dimension.

Examples

```
>>> r = np.random.randn(1000,2)
>>> H, edges = np._histogram3d(r,bins=[10,15])
```

```
_km_plot(ax, km_2, km_4, title)
```

```
_matrix_plot(ax, mat)
```

```

_noise_plot(ax, residual, title)

_noise_plot_2d(ax, res_x, res_y, title)

_plot_3d_histogam(Mx, My, ax=None, title='PDF', xlabel='$M_{\{x\}}$', ylabel='$M_{\{y\}}$',
                   zlabel='Frequency', tick_size=12, title_size=14, label_size=10, label_pad=12,
                   r_fig=False, dpi=150)
    Plot 3d bar plot

_plot_autocorrelation_1d(lags, acf)

_plot_autocorrelation_2d(lags, acfx, acfy, acfm, ccf)

_plot_data(data_in, title='title', x_label='$m_x$', y_label='$m_y$', z_label='z', zlim=None, ax=None,
            clear=True, legend=False, plot_plane=False, tick_size=12, title_size=16, label_size=14,
            label_pad=12, label=None, order=3, m=False, m_th=2, dpi=150, heatmap=False)
    Plot data on a 3d axis

_plot_heatmap(data, title='title', num_ticks=5)
    Plots heatmap of data

_plot_histograms(timeseries, vector, heatmap=False, dpi=150, kde=False, title_size=14, label_size=15,
                  tick_size=12, label_pad=8, **plot_text)
    Plot histogram figures

_plot_noise_characterstics(data, dpi=150, kde=True, title_size=14, tick_size=15, label_size=15,
                             label_pad=8)
    Plot noise charactersitic figure

_plot_summary(data, vector=True, kde=False, tick_size=12, title_size=15, label_size=15, label_pad=8,
               n_ticks=3, timeseries_start=0, timeseries_end=1000, **plot_text)
    Plots the summary chart

_plot_timeseries(timeseries, vector, start=0, stop=1000, n_ticks=3, dpi=150, tick_size=12, title_size=14,
                  label_size=14, label_pad=0, **plot_text)
    Plots timeseries figure

_qq_plot(ax, residual, title)

_remove_nans(Mx, My)
    Remove nan's from data

_set_zaxis_to_left(ax)
    Sets the z-axis of 3d figure to left

_slider_2d(slider_data, init_pos=0, limits=None, prefix='Dt', **plot_text)
    Get slider for analysed scalar data

_slider_3d(slider_data, init_pos=0, prefix='dt', zlim=None, order=None, polar=False, **plot_text)
    Get slider for analysed vector data.

_stylize_axes(ax, x_label=None, y_label=None, title=None, tick_size=20, title_size=20, label_size=20,
               label_pad=12)
    Beautify the plot axis

_trace_pane(data)
    Thrace an arbetary surface that covers the data points.

```

Notes

To be used only to get a better visual of the shape of the surface.

`_update_axis_range(ax, x, both=True)`

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FIVE**

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