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**MICS**

*Release 0.2.0*

**May 09, 2018**



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Mixtures of Independently Collected Samples

- Free software: MIT license

## 1.1 Installation

```
pip install mics
```

## 1.2 Documentation

<https://mics.readthedocs.io/>

## 1.3 Development

To run the all tests run:

```
tox
```

Note, to combine the coverage data from all the tox environments run:

Windows	<pre>set PYTEST_ADDOPTS=--cov-append tox</pre>
Other	<pre>PYTEST_ADDOPTS=--cov-append tox</pre>



## CHAPTER 2

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### Installation

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At the command line:

```
pip install mics
```





## CHAPTER 3

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### Usage

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To use MICS in a project:

```
import mics
```



## 4.1 mics

## 4.2 mixtures

**class** `mics.mixtures.mixture`

A mixture of independently collected samples (MICS)

### Parameters

- **samples** (*list or tuple*) – a list of samples.
- **title** (*str, optional*) – a title.
- **verbose** (*bool, optional*) – a verbosity tag.
- **tol** (*float, optional*) – a tolerance.

**free\_energies** (*reference=0*)

Returns a data frame containing the relative free energies of the datasetd samples of a *mixture*, as well as their standard errors.

**reweighting** (*potential, properties={}, derivatives={}, combinations={}, conditions=Empty DataFrame Columns: [] Index: [], reference=0, \*\*kwargs*)

Performs reweighting of the properties computed by *functions* from the mixture to the samples determined by the provided *potential* with all *parameter* values.

### Parameters

- **potential** (*string*)
- **properties** (*dict of strings*)
- **combinations** (*dict of strings*)
- **derivatives** (*dict of tuples*)
- **conditions** (*pandas.DataFrame*)

- **verbose** (*boolean*)
- **\*\*kwargs**

## 4.3 samples

**class** `mics.samples.pool` (*label=""*, *verbose=False*)

A pool of independently collected samples.

**class** `mics.samples.sample` (*dataset*, *potential*, *autocorr=None*, *label=None*, *batchsize=None*, *verbose=False*, *\*\*kwargs*)

A sample of configurations collected at a specific equilibrium state, aimed to be part of a mixture of independently collected samples (MICS).

### Args:

**dataset** (**pandas.DataFrame**): a data frame whose rows represent configurations datasetd according to a given probability distribution and whose columns contain a number of properties evaluated for such configurations.

**potential** (**function**): the reduced potential that defines the equilibrium sample. This function might for instance receive **x** and return the result of an element-wise calculation involving **x["a"]**, **x["b"]**, etc, with "a", "b", etc being names of properties in **dataset**.

**autocorr** (**function, optional**): a function similar to **potential**, but whose result is an autocorrelated property to be used for determining the effective dataset size. If omitted, **potential** will be used to for this purpose.

**Note:** Formally, functions **potential** and **autocorr** must receive **x** and return **y**, where  $length(y) == nrow(x)$ .

## 4.4 utils

`mics.utils.covariance` (*y, ym, b*)

Computes the covariance matrix of the rows of matrix *y* among themselves. The method of Overlap Batch Mean (OBM) is employed with blocks of size *b*.

`mics.utils.cross_covariance` (*y, ym, z, zm, b*)

Computes the cross-covariance matrix between the rows of matrix *y* with those of matrix *z*. The method of Overlap Batch Mean (OBM) is employed with blocks of size *b*.

`mics.utils.genfunc` (*function, variables, constants*)

Returns a function based on the passed argument.

`mics.utils.multimap` (*functions, sample*)

Applies a list of *functions* to **DataFrame** *sample* and returns a numpy matrix whose number of rows is equal to the length of list *functions* and whose number of columns is equal to the number of rows in *sample*.

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**Note:** Each function of the array might for instance receive *x* and return the result of an element-wise calculation involving  $x["A"]$ ,  $x["B"]$ , etc, with "A", "B", etc being names of properties in **DataFrame** *sample*.

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`mics.utils.overlapSampling` (*u*)

Computes the relative free energies of all sampled states using the Overlap Sampling method of Lee and Scott (1980).

`mics.utils.pinv(A)`

Computes the Moore-Penrose pseudoinverse of a symmetric matrix using eigenvalue decomposition.



Contributions are welcome, and they are greatly appreciated! Every little bit helps, and credit will always be given.

### 5.1 Bug reports

When [reporting a bug](#) please include:

- Your operating system name and version.
- Any details about your local setup that might be helpful in troubleshooting.
- Detailed steps to reproduce the bug.

### 5.2 Documentation improvements

MICS could always use more documentation, whether as part of the official MICS docs, in docstrings, or even on the web in blog posts, articles, and such.

### 5.3 Feature requests and feedback

The best way to send feedback is to file an issue at <https://github.com/craabreu/mics/issues>.

If you are proposing a feature:

- Explain in detail how it would work.
- Keep the scope as narrow as possible, to make it easier to implement.
- Remember that this is a volunteer-driven project, and that code contributions are welcome :)

## 5.4 Development

To set up *mics* for local development:

1. Fork *mics* (look for the “Fork” button).
2. Clone your fork locally:

```
git clone git@github.com:your_name_here/mics.git
```

3. Create a branch for local development:

```
git checkout -b name-of-your-bugfix-or-feature
```

Now you can make your changes locally.

4. When you’re done making changes, run all the checks, doc builder and spell checker with *tox* one command:

```
tox
```

5. Commit your changes and push your branch to GitHub:

```
git add .  
git commit -m "Your detailed description of your changes."  
git push origin name-of-your-bugfix-or-feature
```

6. Submit a pull request through the GitHub website.

### 5.4.1 Pull Request Guidelines

If you need some code review or feedback while you’re developing the code just make the pull request.

For merging, you should:

1. Include passing tests (run *tox*)<sup>1</sup>.
2. Update documentation when there’s new API, functionality etc.
3. Add a note to *CHANGELOG.rst* about the changes.
4. Add yourself to *AUTHORS.rst*.

### 5.4.2 Tips

To run a subset of tests:

```
tox -e envname -- py.test -k test_myfeature
```

To run all the test environments in *parallel* (you need to `pip install detox`):

```
detox
```

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<sup>1</sup> If you don’t have all the necessary python versions available locally you can rely on Travis - it will run the tests for each change you add in the pull request.

It will be slower though ...



## CHAPTER 6

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### Authors

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- Charles R. A. Abreu - <http://atoms.peq.coppe.ufrj.br>



### 7.1 0.2.0 (2018-05-09)

- Implementation of classes `sample`, `pool`, `mixture`, `MICS` and `MBAR`.

### 7.2 0.1.0 (2017-10-11)

- First release on PyPI.



## CHAPTER 8

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### Indices and tables

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