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# alchemtest Documentation

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## Overview

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**Warning:** This library is in an **alpha** state. The library and the documentation is incomplete. Use in production at your own risk.

**alchemtest** is a collection of test datasets for alchemical free energy calculations. The datasets come from a variety of software packages, primarily molecular dynamics engines, and are used as the test set for [alchemlyb](#). The package is standalone, however, and can be used for any purpose.



# CHAPTER 1

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## Installing alchemtest

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**alchemtest** is pure-Python, so it can be installed easily via pip:

```
pip install alchemtest
```

If you wish to install this in your user site-packages, use the --user flag:

```
pip install --user alchemtest
```

### 1.1 Installing from source

from source. Clone the source from GitHub with:

```
git clone https://github.com/alchemistry/alchemtest.git
```

then do:

```
cd alchemtest  
pip install .
```

If you wish to install this in your user site-packages, use the --user flag:

```
pip install --user .
```



# CHAPTER 2

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## Basic usage

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All datasets in `alchemtest` are accessible via `load_*` functions, organized in submodules by the software package that generated them. The current set of submodules are:

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<code>gmx</code>	Gromacs molecular dynamics simulation datasets.
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As an example, we can access the *Gromacs: Benzene in water* dataset with:

```
>>> from alchemtest.gmx import load_benzene  
>>> bz = load_benzene()
```

and use the resulting `Bunch` object to introspect what this dataset includes. In particular, it features a `DESCR` attribute with a human-readable description of the dataset:

```
>>> print(bz.DESCR)  
Gromacs: Benzene in water  
=====
```

Notes

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Data Set Characteristics:

- :Number of Legs: 2 (Coulomb, VDW)
- :Number of Windows: 5 for Coulomb, 16 for VDW
- :Length of Windows: 40ns

- :Missing Values: None
- :Creator: \I. Kenney
- :Donor: Ian Kenney (ian.kenney@asu.edu)
- :Date: March 2017

Benzene in water, alchemically turned into benzene in vacuum separated from water

This dataset was generated using `MDPOW <<https://github.com/Becksteinlab/MDPOW>>`\_, ↴with the `Gromacs <<http://www.gromacs.org/>>`\_ molecular dynamics engine.

as well as the dataset itself:

```
>>> bz.data.keys()  
['VDW', 'Coulomb']
```

which consists in this case of two alchemical legs, each having several files. For this dataset each file happens to correspond to a simulation sampling a particular  $\lambda$ :

```
>>> bz.data['Coulomb']  
['/usr/local/python3.6/site-packages/alchemtest/gmx/benzene/Coulomb/0000/dhdl.xvg.bz2  
↳',  
 '/usr/local/python3.6/site-packages/alchemtest/gmx/benzene/Coulomb/0250/dhdl.xvg.bz2  
↳',  
 '/usr/local/python3.6/site-packages/alchemtest/gmx/benzene/Coulomb/0500/dhdl.xvg.bz2  
↳',  
 '/usr/local/python3.6/site-packages/alchemtest/gmx/benzene/Coulomb/0750/dhdl.xvg.bz2  
↳',  
 '/usr/local/python3.6/site-packages/alchemtest/gmx/benzene/Coulomb/1000/dhdl.xvg.bz2  
↳']
```

These paths can be read by any appropriate parser for further analysis. For this particular dataset, see [alchemlyb.parsing.gmx](#) for a good set of parsers.

# CHAPTER 3

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## Gromacs datasets

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Gromacs molecular dynamics simulation datasets.

The `alchemlyb.gmx` module features datasets generated using the `Gromacs` molecular dynamics engine. They can be accessed using the following accessor functions:

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`load_benzene()`

Load the Gromacs benzene dataset.

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### 3.1 Gromacs: Benzene in water

#### 3.1.1 Notes

**Data Set Characteristics:**

**Number of Legs** 2 (Coulomb, VDW)

**Number of Windows** 5 for Coulomb, 16 for VDW

**Length of Windows** 40ns

**System Size** 1668 atoms

**Temperature** 300 K

**Pressure** 1 bar

**Alchemical Pathway** vdw + coul → vdw → vacuum

**Experimental Hydration Free Energy** -0.90 +- 0.2 kcal/mol

**Missing Values** None

**Creator** I. Kenney

**Donor** Ian Kenney ([ian.kenney@asu.edu](mailto:ian.kenney@asu.edu))

**Date** March 2017

Benzene in water, alchemically turned into benzene in vacuum separated from water

This dataset was generated using [MDPOW](#), with the [Gromacs](#) molecular dynamics engine.

Experimental value sourced from [Mobley2013].

`alchemtest.gmx.load_benzene()`

Load the Gromacs benzene dataset.

**Returns data** – Dictionary-like object, the interesting attributes are: - ‘data’ : the data files by alchemical leg - ‘DESCR’: the full description of the dataset

**Return type** Bunch

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## Bibliography

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[Mobley2013] Mobley, David L. (2013). Experimental and Calculated Small Molecule Hydration Free Energies. UC Irvine: Department of Pharmaceutical Sciences, UCI. Retrieved from: <http://escholarship.org/uc/item/6sd403pz>



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