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# **pytc Documentation**

*Release 1.1.5*

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Open source python software for extracting thermodynamic information from Isothermal Titration Calorimetry experiments.

- We welcome bug reports, patches, and new thermodynamic models. See the [contributing](#) page for how to get involved.
- If you find pytc useful, please star and/or fork the project on [github](#).
- *Citation:* Duvvuri H, Wheeler LC, Harms MJ. (2018) “pytc: Open-Source Python Software for Global Analyses of Isothermal Titration Calorimetry Data” *Biochemistry*. doi://10.1021/acs.biochem.7b01264



# CHAPTER 1

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[Download](#)

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[Windows GUI](#) | [Mac GUI](#) | [Linux GUI](#) | [API](#) | [Source code](#)





## CHAPTER 2

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### Philosophy

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- **Open source and cross platform.** The full source code should be available. The program should not require proprietary software to run.
- **Rigorous.** Program should use best practices and current algorithms for performing fits and assessing fit quality.
- **Ease of use.** Fitting basic models should be easy. Implementing completely new thermodynamic models should be straightforward.
- **Accessible for users and programmers.** It should have both a graphical user interface and a well-designed API.



## CHAPTER 3

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### Features

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- Simple, cross-platform graphical user interface.
- Clean, pythonic API
- New models can be defined using a few lines of python code
- Easy integration with [jupyter](#) notebooks for writing custom fitting scripts



- Installation
- Fitting models using the GUI
- Fitting models using the API
- **Reference:**
  - Thermodynamic models for individual experiments
  - Global fits
  - Fit strategies
  - Fit statistics
  - Defining new models
  - Contributing to the project

**Warning:** `pytc` will fit all sorts of complicated models to your data. It is up to you to make sure the fit is justified by the data. See the [Fitting and statistics](#) section to see what `pytc` reports to help in this decision making.



## CHAPTER 5

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### GUI Demo

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An animated gif showing the GUI in action.





Fit a single-site binding model to a  $Ca^{2+}/EDTA$  binding experiment.

```
import pytc

# Load in integrated heats from an ITC experiment
e = pytc.ITCExperiment("demos/ca-edta/tris-01.DH",
                      pytc.indiv_models.SingleSite)

# Create the global fitter, add the experiment
g = pytc.GlobalFit()
g.add_experiment(e)

# Do the fit
g.fit()

# Print the results out
g.plot()
print(g.fit_as_csv)
```



Our implementation is built on `python3` extended with `numpy`, `scipy`, `matplotlib` and `emcee`. The GUI is built on `pyqt5`.

## 7.1 `pytc` package

### 7.1.1 Subpackages

`pytc.experiments` package

Submodules

`pytc.experiments.base` module

Module contents

`pytc.global_connectors` package

Submodules

`pytc.global_connectors.base` module

`pytc.global_connectors.num_protons` module

`pytc.global_connectors.vant_hoff` module

`pytc.global_connectors.vant_hoff_extended` module

## Module contents

pytc.indiv\_models package

### Submodules

pytc.indiv\_models.base module

pytc.indiv\_models.binding\_polynomial module

pytc.indiv\_models.blank module

pytc.indiv\_models.single\_site module

pytc.indiv\_models.single\_site\_competitor module

## Module contents

pytc.util package

### Submodules

pytc.util.util module

## Module contents

### 7.1.2 Submodules

7.1.3 pytc.fit\_param module

7.1.4 pytc.global\_fit module

### 7.1.5 Module contents

## CHAPTER 8

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### Indices and tables

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- `modindex`
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