# **AIMBAT Documentation**

Release 0.1.2

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#### INTRODUCTION

#### 1.1 About AIMBAT

AIMBAT (Automated and Interactive Measurement of Body wave Arrival Times) is an open-source software package for efficiently measuring teleseismic body wave arrival times for large seismic arrays [LouVanDerLee2013]. It is based on a widely used method called MCCC (Multi-Channel Cross-Correlation) [VanDecarCrosson1990]. The package is automated in the sense of initially aligning seismograms for MCCC which is achieved by an ICCS (Iterative Cross Correlation and Stack) algorithm. Meanwhile, a GUI (graphical user interface) is built to perform seismogram quality control interactively. Therefore, user processing time is reduced while valuable input from a user's expertise is retained. As a byproduct, SAC [GoldsteinDodge2003] plotting and phase picking functionalities are replicated and enhanced.

Modules and scripts included in the AIMBAT package were developed using Python programming language and its open-source modules on the Mac OS X platform since 2009. The original MCCC [VanDecarCrosson1990] code was transcribed into Python. The GUI of AIMBAT was inspired and initiated at the 2009 EarthScope USArray Data Processing and Analysis Short Course. AIMBAT runs on Mac OS X, Linux/Unix and Windows thanks to the platform-independent feature of Python. It has been tested on Mac OS 10.6.8 and 10.7 and Fedora 16.

The AIMBAT software package is distributed under the GNU General Public License Version 3 (GPLv3) as published by the Free Software Foundation.

#### 1.2 Associated Documents

- Seismological Research Letters Paper
- PDF Version of Manual

**TWO** 

### **INSTALLING DEPENDENCIES**

### 2.1 Getting your operating system

You may need to know .. image:: installing-images/system\_preferences.png

# 2.2 Installing Python

Shaowei Lin suggested Enthought Canopy to install all the Python packages easily. If you download the free version of Enthought Canopy, it gives you everything you need for installing AIMBAT properly. If you do not want to use Enthought Canopy, read the rest of this section to use Macports or Pip.

# 2.3 Python Dependencies

- Numpy
- Scipy
- Matplotlib
- iPython (optional)

THREE

#### **INSTALLING AIMBAT**

### 3.1 Getting the Packages

AIMBAT is released as a sub-package of pysmo in the name of pysmo.aimbat along with another sub-package pysmo.sac. The latest releases of pysmo.sac and pysmo.aimbat are available for download at the official project webpage and Github.

The packages should be installed into the Python site-packages directory. To find out where that is, in the python console, do:

```
import site;
site.getsitepackages()
```

Whatever is output there, lets call it <pkg-install-dir>. You can choose to install AIMBAT either locally or globally, depending on whether you want all users of the computer to have access to it.

Make a directory called pysmo, and place the sac and aimbat directories there.

Now that we know the location of the site-packages direction, cd into it. Call the path to it <pkg-install-dir>. Notice that in this case, the site-packages has been installed for all users on the computer, not just the current user's home directory.

Put the two Python packages inside the directory.

# 3.2 Installing pysmo.sac

Python module Distutils is used to write a setup.py script to build, distribute, and install pysmo.sac. In the directory <pkg-install-dir>/pysmo-sac-0.5>, type:

```
sudo python setup.py build
sudo python setup.py install
```

to install it and its package information file pysmo.sac-0.5-py2.7.egg-info to the global site-packages directory ctory fix>/lib/python2.7/site-packages, which is the same as Numpy, Scipy, and Matplotlib.

If you don't have write permission to the global site-packages directory, use the *-user* option to install to *<userbase>/lib/python2.7/site-packages*:

```
python setup.py install --user
```

This will install it to your home directory only, not for all users on the computer. Try not to use this option though, as installing without the sudo command has caused problems in the past.

If you successfully installed the sac module, in the python console, this should happen after you type from pysmo import sac

# 3.3 Installing pysmo.aimbat

Three sub-directories are included in the <pkg-install-dir>/pysmo/pysmo-aimbat-0.1.2> directory: example, scripts, and src, which contain example SAC files, Python scripts to run at the command line, and Python modules to install, respectively.

The core cross-correlation functions in pysmo.aimbat are written in both Python/Numpy (xcorr.py) and Fortran (xcorr.f90). Therefore, we need to use Numpy's Distutils module for enhanced support of Fortran extension. The usage is similar to the standard Disutils.

Note that some sort of Fortran compiler must already be installed first. Specify them in place of gfortran in the following commands.

In the directory <pkg-install-dir>/pysmo/pysmo-aimbat-0.1.2, type:

```
sudo python setup.py build --fcompiler=gfortran
sudo python setup.py install
```

to install the src directory.

Add <pkg-install-dir>/pysmo/pysmo-aimbat.0.1.2/scripts to environment variable PATH in a shells start-up file for command line execution of the scripts.

**Bash Shell Users:** export PATH=\$PATH:<pkg-install-dir>/pysmo/pysmo-aimbat-0.1.2/scripts in .bashrc files.

C Shell Users: setenv PATH=\$PATH:<pkg-install-dir>/pysmo/pysmo-aimbat-0.1.2/scripts in .bashrc files.

If AIMBAT has been installed, type from pysmo import aimbat in a Python shell, and no errors should appear.

### **FOUR**

### STANDING ORDER FOR DATA (SOD)

### 4.1 Installing SOD

First, download SOD.

Once you have gotten the folder for SOD, put it somewhere where you won't touch it too much. What I did was put the SOD folder in my home directory, though other places are acceptable as well, as long as its not too easy to delete it by accident.



Once you have it there, get the path to the sod folder's bin and put it in your path folder.



Inside my home directory's bash profile (you get the by typing cd), you put the path to sod-3.2.3/bin by adding in either the bash or bash\_profile or profile files:

### 4.2 Downloading Data with SOD

**Authors** Trevor Bollmann

- 1. Create a sod recipe and place it in the folder that you would like the data to download to.
  - sod -f <recipename>.xml
- 2. Run verb"sodcut.sh" to cut the seismogram around phase wanted
  - check model within cutevseis.sh
  - run using sodcut.sh <name>
  - watch sdir = processed seismograms
  - Run over the entire downloaded directory (the files sod downloaded)
- 3. Run sodpkl.sh (converts.sac files to python pickles)
  - run using sodpkl.sh [options] <directory>
  - output will automatically be zipped
  - run in DATA directory
- 4. Run ttpick.py (does travel time picking with plotting)
  - can use iccs.py but it does not have plotting capabilities
  - run using ttpick.py [options] <pkl.gz file>
  - do this one event at a time
  - use sacp2 to look at the stacking of the seismograms
  - you can sort the seismograms using the -s flag

#### 5. run getsta.py (creates a loc.sta file)

• getsta.py [options] <pkl.gz files>

#### 6. Run EITHER of these:

- FIRST CHOICE
- run mccc2delay.py (converts mccc delays to actual delays) by doing mccc2delay.py [option] <.mcp files>
- run getdelay.py (creates a delay file) by doing getdelay.py [options] <\*.px>
  - Can possibly use doplotsta.sh, plots all of the events and their station delays
- Run evmcdelay.sh
- SECOND CHOICE
  - :code:verb"ttcheck.py" to compare the delay times of the p and s waves. Should form a nice cloud with the mean value in line with the cloud.

#### 7. If you need to remove a station from an event you can use :code:verb"pklsel.py"

- Run using :code:verb"pklsel.py [pkl file] –d [stnm]" to remove one station
- Only works for one event at a time

#### 8. If you need to filter the data to be able to pick use :code:verb"evsacbp.sh"

- run using :code:verb"evsacbp.sh [pkl file] bp1 bp2"
- Automatically uses two corners
- run in the whole downloaded directory (the one with the sac directory)

**FIVE** 

### **ANALYZING DATA**

# 5.1 Seismic Analysis Code (SAC)

AIMBAT uses Seismic Analysis Code (SAC) formatting for some of the files it runs and outputs. To get SAC, you will need to fill out a software request form available on the IRIS website.

#### MEASURING TELESEISMIC BODY WAVE ARRIVAL TIMES

The core idea in using AIMBAT to measure teleseismic body wave arrival times has two parts:

- · automated phase alignment, to reduce user processing time, and
- interactive quality control, to retain valuable user inputs.

### **6.1 Automated Phase Alignment**

The ICCS algorithm calculates an array stack from predicted time picks, cross-correlates each seismogram with the array stack to Find the time lags at maximum cross-correlation, then use the new time picks to update the array stack in an iterative process. The MCCC algorithm cross-correlates each possible pair of seismograms and uses a least-squares method to calculate an optimized set of relative arrival times. Our method is to combine ICCS and MCCC in a four-step procedure using four anchoring time picks  $_0T_i$ ,  $_1T_i$ ,  $_2T_i$ , and  $_3T_i$ .

- 1. Coarse alignment by ICCS
- 2. Pick phase arrival at the array stack
- 3. Refined alignment by ICCS
- 4. Final alignment by MCCC

The one-time manual phase picking at the array stack in step (b) allows the measurement of absolute arrival times. The detailed methodology and procedure can be found in [LouVanDerLee2013].

Table 6.1: Time picks and their SAC headers used in the procedure for measuring teleseismic body wave arrival times.

Step	Algorithm Time Window	Input			Output	
Бер			Time Pick	Time Header	Time Pick	Time Header
1.	ICCS	$W_a$	$_{0}T_{i}$	T0	$1T_i$	T1
2	ICCS	$W_b$	$_{2}T_{i}^{\prime}$	T2	$_2T_i$	T2
2.						
	MCCS	$W_b$	$\Big _{2}T_{i}$	T2	$\Big _{\ _3T_i}$	Т3
4.	Mees	,,,,	2-1	12	3-1	10

# **SEVEN**

# **INDICES AND TABLES**

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- modindex
- search