Tutorial on BayesVP Documentation

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CHAPTER

ONE

BAYESVP: TUTORIAL

bayesvp is a Bayesian MCMC parallel Voigt profile fitting routine written by Cameron Liang & Andrey Kravtsov.

bayesvp provides a number of helpful executable scripts that work with command line arguments (saved in your environment PATH). The main functionality is the MCMC Voigt profile fitting (bvpfit) where the user supplies a config file that specifies parameters for the fitting. These include parameter priors, number of walkers, parallel threads, line spread function, continuum model, Bayesian model comparisons, and etc. There are utility functions that allow users to quickly create an example config file, process and plot the chains, process and plot the best fit models and more. For details on the code, refer to the papers Liang & Kravtsov 2017 and Liang et al. 2017.

The following is a short tutorial on how to use bayesup fitting routine.

1.1 Installation

Install bayesvp using pip with the --user flag:

```
pip install bayesvp --user
```

For users with Python 3 and above, you need to convert all the "bayesvp" scripts (written originally in Python 2.7) to Python 3 using the Python package '2to3 https://pypi.org/project/2to3/.

After installing bayesvp we should run its unit tests to ensure the package works as expected. The simplest way to do this is inside a python shell:

```
from bayesvp.tests import run_tests
```

The output should look something like this:

```
The output should look something like this:

test_config_file_exists (bayesvp.tests.test_config.TCConfigFile) ... ok

test_default_no_continuum_params (bayesvp.tests.test_config.TCConfigFile) .

... ok

test_example_mcmc_params (bayesvp.tests.test_config.TCConfigFile) ... ok

...

test_prior (bayesvp.tests.test_likelihood.TCPosterior) ... ok

test_general_intensity (bayesvp.tests.test_model.TCSingleVP) ... ok

test_simple_spec (bayesvp.tests.test_model.TCSingleVP) ... ok
```

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```
Ran 13 tests in 3.654s
OK
```

This would ensure bayesup package is installed.

1.2 Fit a simple absorption line with bayesvp

We describe a simple tutorial on how to use bayesvp package. It can be run in an interactive python session or using a python script.

To start bayesvp, first we need to setup a configuration file. bayesvp is meant to run with this file in background as it can take few minutes for MCMC sampling depending on the chosen number of walkers, steps, and parallel processes.

Below we illustrate a simple interactive use of the code and setting up of a configuration file to fit an O VI transition with rest wavelength of 1031.926 Å.

1.2.1 Setup a 'config file'

First step is to import bayesvp package. Next we import an object, bvp_write_config, that interactively asks the user a few questions to create a config file.

```
In [1]: import bayesvp
In [2]: from bayesvp.scripts import bvp_write_config as wc
```

Let us assume that the spectrum in question is located in the following directory:

```
In [3]: spectrum_path = '/home/<username>/bayesvp_tutorial/codes/examples/'
```

Suppose the file name of the example spectrum is OVI.spec with three of columns of data: wave, flux, error. We can use the bvp_write_config routine to set up the config file as:

```
In [4]: config_writer = wc.WriteBayesVPConfig().print_to_file("-i")
```

On executing the above command, the routine will ask the user few questions:

The questions are on the left, while on the right, as an example, the answers are written. These answers can change based on different user preferences.

```
Path to spectrum: /home/<username>/bayesvp_tutorial/codes/examples
Spectrum filename: OVI.spec
filename for output chain: o6
Enter the name of the ion to be fitted: first atom name, next its ionization state
atom: O
```

```
state: VI
```

Enter the number of components to used in the fitting routine. If you wish to use more components then type in the number

```
Maximum number of components to try: 1
```

Enter the observed wavelength region to be used in the fitting routine

```
Starting wavelength(A): 1030
Ending wavelength(A): 1033
Enter the priors:
```

Enter the assumed minimum and maximum values of column density for the absorption feature to be fitted

```
min logN [cm^{-2}] = 10
max logN [cm^{-2}] = 18
```

Similarly, enter the assumed minimum and maximum values of Doppler b parameter for the absorption feature to be fitted

```
min b [km/s] = 0
max b [km/s] = 100
```

Enter the central redshift of the absorption feature corresponding to the ion to be fitted

```
central redshift = 0
velocity range [km/s] = 300
```

Here we enter parameters to be used for MCMC sampling

```
Enter the MCMC parameters:

Number of walkers: 400

Number of steps: 2000

Number of processes: 8

Model selection method bic(default), aic, bf: bic

MCMC sampler kombine(default), emcee: kombine
```

On completion of the above step, the configuration file is automatically written within a sub-directory where the spectrum is located.

Written config file: /home/<username>/bayesvp_tutorial/codes/examples/bvp_configs/config_OVI.dat

The saved configuration file 'config_OVI.dat' will look like the following:

```
spec_path /home/<username>/bayesvp_tutorial/codes/examples
output o6
mcmc 400 2000 8 bic kombine
%% OVI.spec 1030.000000 1033.000000
% O VI 15 30 0.0000000
logN 10.00 18.00
b     0.00 100.00
z     0.000000 300.00
```

To deal with HST/COS data, which requires incorporation of line-spread-functions (LSFs), the above saved **config_OVI.dat** needs to be modified as such:

```
spec_path /home/<username>/bayesvp_tutorial/codes/examples
output o6
mcmc 400 2000 8 bic kombine
%% OVI.spec 1030.000000 1033.000000
% O VI 15 30 0.0000000
lsf COS_res<central_wavel>.dat
logN 10.00 18.00
b    0.00 100.00
z    0.000000 300.00
```

Note the changes made above. An additional statement: lsf <LSF kernel at the central wavelength> is added.

The HST/COS LSFs should be saved in a sub-directory (named **database**) where the spectrum is located.

1.2.2 Run a MCMC fit

To start MCMC fit, firstly we need to import three more objects:

```
In [5]: from bayesvp.config import DefineParams
In [6]: from bayesvp.mcmc_setup import bvp_mcmc_single as mc_single
In [7]: from bayesvp.mcmc_setup import bvp_mcmc as mcmc
```

bayesvp can be run by supplying the full path to the config file as a command line argument. bayesvp will print to screen the relevant information from the config file