WDmodel Documentation

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Contents

| 1 | 1 WDmodel | | | |
|----|-----------|---------------------------------|----|--|
| | 1.1 | About | 3 | |
| | 1.2 | Compatibility | 3 | |
| | 1.3 | Analysis | 3 | |
| 2 | Help | | 5 | |
| | 2.1 | Installing WDmodel | 5 | |
| | | 2.1.1 Installation Instructions | 4 | |
| | | 2.1.2 Extra | 7 | |
| | 2.2 | Using WDmodel | 8 | |
| | | 2.2.1 Usage | 8 | |
| | | 2.2.2 Useful runtime options | Ç | |
| | 2.3 | Analyzing WDmodel | 10 | |
| | | | 10 | |
| | 2.4 | WDmodel | 14 | |
| | | 2.4.1 WDmodel package | | |
| 3 | Indic | ees and tables | 59 | |
| Рy | thon I | Module Index | 61 | |

WDmodel: Bayesian inference of white dwarf properties from spectra and photometry to establish spectrophotometric standards

Contents 1

2 Contents

CHAPTER 1

WDmodel

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1.1 About

WDmodel is a DA White Dwarf model atmosphere fitting code. It fits observed spectrophotometry of DA White Dwarfs to infer intrinsic model atmosphere parameters in the presence of dust and correlated spectroscopic flux calibration errors, thereby determining full SEDs for the white dwarf. Its primary scientific purpose is to establish a network of faint (V = 16.5-19 mag) standard stars, suitable for LSST and other wide-field photometric surveys, and tied to HST and the CALSPEC system, defined by the three primary standards, GD71, GD153 and G191B2B.

Click on the badges above for code, licensing and documentation.

1.2 Compatibility

The code has been tested on Python 2.7 and 3.6 on both OS X (El Capitan and Sierra) and Linux (Debian-derivatives). Send us email or open an issue if you need help!

1.3 Analysis

We're working on a publication with the results from our combined Cycle 22 and Cycle 20 data, while ramping up for Cycle 25! A full data release of Cycle 20 and 22 HST data, and ground-based imaging and spectroscopy will accompany the publication. Look for an updated link here!

You can read the first version of our analysis of four of the Cycle 20 objects here

That analysis was intended as a proof-of-concept and used custom IDL routines from Jay Holberg (U. Arizona) to infer DA intrinsic parameters and custom python code to fit the reddening parameters. This code is intended to (significantly) improve on that analysis.

CHAPTER 2

Help

This document will help get you up and running with the WDmodel package.

For the most part, you can simply execute code in grey boxes to get things up and running, and ignore the text initially. Come back to it when you need help, or to configure the fitter.

2.1 Installing WDmodel

This document will step you through getting the WDmodel package installed on your system.

- Installation instructions
 - Get python
 - Get the code
 - Install everything
 - Get auxillary pysynphot files
 - Install the code
- Some extra notes

2.1.1 Installation Instructions

Here's a minimal set of instructions to get up and running. We will eventually get this package up on PyPI and conda-forge, and that should make this even easier.

0. Install python:

We recommend using the anaconda python distribution to run this package. If you don't have it already, follow the instructions here

Make sure you added the conda/bin dir to your path!

If you elect to use your system python, or some other distribution, we will assume you know what you are doing, and you can, skip ahead.

1. Get the code:

Clone this repository

```
git clone https://github.com/gnarayan/WDmodel.git
cd WDmodel
```

2. Install everything:

a. Create a new environment from specification (Preferred! All dependencies resolved!)

```
conda env create -f docs/env/conda_environment_py[27|36]_[osx64|i686].yml
```

or

b. Create a new environment from scratch (Let conda figure out dependencies and you sort out potential issues)

```
cp docs/env/condarc.example ~/.condarc
conda create -n WDmodel
source activate WDmodel
conda install --yes --file dependencies_py[27|36].txt
```

Setting up an environment vs setting up a known good environment

The env folder contains files to help get you setup using a consistent environment with all packages specified.

The requirements_py[27|36].txt files contains a list of required python packages and known working versions for each. They differ from the dependencies_py[27|36].txt files in the root directory in that those files specify packages and version ranges, rather than exact versions, to allow conda to resolve dependecies and pull updated versions.

Of course, the environment really needs more than just python packages, while pip only manages python packages. The conda environment files, conda_environment_py[27|37]_[osx64|i686].yml files can be used to create conda environments with exact versions of all the packages for python 2.7 or 3.6 on OS X or linux. This is the most reliable way to recreate the entire environment.

3. Get the latest HST CDBS files:

These are available over FTP from [ftp://archive.stsci.edu/pub/hst/pysynphot/]

Untar them wherever you like, and set the PYSYN_CDBS environment variable. You need at least symphot1.tar.gz and symphot6.tar.gz.

```
export PYSYN_CDBS=place_you_untarred_the_files
```

4. Install the package [optional]:

python setup.py install

2.1.2 Extra

The instructions should be enough to get up and running, even without sudo privileges. There's a few edge cases on cluster environments though. These notes may help:

Some extra notes on installation

If you followed the installation process detailed above, you shouldn't need these notes, but they are provided for users who may be running on environments they do not manage themselves.

- Installing eigen3 without conda
- Installing OpenMPI and mpi4py without conda
- Installing on a cluster

Installing eigen3 without conda

If eigen3 isn't on your system, and installing it with conda didn't work

For OS X do:

brew install eigen

or on a linux system with apt:

apt-get install libeigen3-dev

or compile it from source

Note that if you do install it in a custom location, you may have to compile celerite yourself.

pip install celerite --global-option=build_ext --global-option=-I/path/to/eigen3

Installing OpenMPI and mpi4py without conda

if no mpi is on your system, and installing it with conda didn't work

For OS X do:

brew install [mpich|mpich2|open-mpi]

on a linux system with apt:

apt-get install openmpi-bin

and if you had to resort to brew or apt, then finish with:

pip install mpi4py

Notes from installing on the Odyssey cluster at Harvard

These may be of use to get the code up and running with MPI on some other cluster. Good luck.

Odyssey uses the lmod system for module management, like many other clusters You can module spider openmpi to find what the openmpi modules.

The advantage to using this is distributing your computation over multiple nodes. The disadvantage is that you have to compile mpi4py yourself against the cluster mpi.

```
module load gcc/6.3.0-fasrc01 openmpi/2.0.2.40dc0399-fasrc01
wget https://bitbucket.org/mpi4py/mpi4py/downloads/mpi4py-2.0.0.tar.gz
tar xvzf mpi4py-2.0.0.tar.gz
cd mpi4py-2.0.0
python setup.py build --mpicc=$(which mpicc)
python setup.py build_exe --mpicc="$(which mpicc) --dynamic"
python setup.py install
```

2.2 Using WDmodel

This document will help you get comfortable using the WDmodel package.

- Usage
- Get data
- Running single threaded
- Running with MPI
- Useful options
 - Quick analysis
 - Initializing the fitter
 - Configuring the sampler
 - Resuming the fit

2.2.1 **Usage**

This is the TL;DR version to get up and running.

1. Get the data:

Instructions will be available here when the paper is accepted. In the meantime there's a single test object in the spectroscopy directory. If you want more, Write your own HST proposal! :-P

2. Run a fit single threaded:

```
fit_WDmodel --specfile data/spectroscopy/yourfavorite.flm
```

This option is single threaded and slow, but useful to testing or quick exploratory analysis.

A more reasonable way to run things fast is to use mpi.

3. Run a fit as an MPI process:

```
mpirun -np 8 fit_WDmodel --mpi --specfile=file.flm [--ignorephot]
```

Note that --mpi MUST be specified in the options to WDmodel and you must start the process with mpirun

2.2.2 Useful runtime options

There's a large number of command line options to the fitter, and most of it's aspects can be configured. Some options make sense in concert with others, and here's a short summary of use cases.

Quick looks

The spectrum can be trimmed prior to fitting with the --trimspec option. You can also blotch over gaps and cosmic rays if your reduction was sloppy, and you just need a quick fit, but it's better to do this manually.

If there is no photometry data for the object, the fitter will barf unless ——ignorephot is specified explicitly, so you know that the parameters are only constrained by the spectroscopy.

The fitter runs a MCMC to explore the posterior distribution of the model parameters given the data. If you are running with the above two options, chances are you are at the telescope, getting spectra, and doing quick look reductions, and you just want a rough idea of temperature and surface gravity to decide if you should get more signal, and eventually get HST photometry. The MCMC is overkill for this purpose so you can --skipmcmc, in which case, you'll get results using minuit. They'll be biased, and the errors will probably be too small, but they give you a ballpark estimate.

If you do want to use the MCMC anyway, you might like it to be faster. You can choose to use only every nth point in computing the log likelihood with --everyn - this is only intended for testing purposes, and should probably not be used for any final analysis. Note that the uncertainties increase as you'd expect with fewer points.

Setting the initial state

The fitter really runs minuit to refine initial supplied guesses for parameters. Every now at then, the guess prior to running minuit is so far off that you get rubbish out of minuit. This can be fixed by explicitly supplying a better initial guess. Of course, if you do that, you might wonder why even bother with minuit, and may wish to skip it entirely. This can be disabled with the --skipminuit option. If --skipminuit is used, a dl guess **MUST** be specified.

All of the parameter files can be supplied via a JSON parameter file supplied via the <code>--param_file</code> option, or using individual parameter options. An example parameter file is available in the module directory.

Configuring the sampler

You can change the sampler type (-samptype), number of chain temperatures (--ntemps), number of walkers (--nwalkers), burn in steps (--nburnin), production steps (--nprod), and proposal scale for the MCMC (--ascale). You can also thin the chain (--thin) and discard some fraction of samples from the start (--discard). The default sampler is the ensemble sampler from the emcee package. For a more conservative approach, we recommend the ptsampler with ntemps=5, nwalkers=100, nprod=5000 (or more).

Resuming the fit

If the sampling needs to be interrupted, or crashes for whatever reason, the state is saved every 100 steps, and the sampling can be restarted with --resume. Note that you must have run at least the burn in and 100 steps for it to be possible to resume, and the state of the data, parameters, or chain configuration should not be changed externally (if

they need to be use --redo and rerun the fit). You can increase the length of the chain, and chain the visualization options when you --resume but the state of everything else is restored.

You can get a summary of all available options with --help

Useful routines

There are a few useful routines included in the WDmodel package. Using WDmodel itself will do the same thing as fit_WDmodel. If you need to look at results from a large number of fits, print_WDmodel_result_table and print_WDmodel_residual_table will print out tables of results and residuals. make_WDmodel_slurm_batch_scripts provides an example script to generate batch scripts for the SLURM system used on Harvard's Odyssey cluster. Adapt this for use with other job queue systems or clusters.

2.3 Analyzing WDmodel

This document describes the output produced by the WDmodel package.

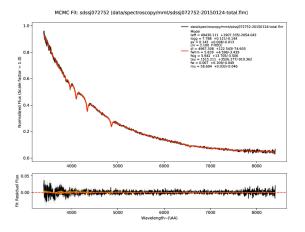
- Analysis
 - The fit
 - Spectral flux calibration errors
 - Hydrogen Balmer line diagnostics
 - Posterior distributions
 - Output files

2.3.1 Analysis

There's many different outputs (ascii files, bintables, plots) that are produced by the WDmodel package. We'll describe the plots first - it is a good idea to look at your data before using numbers from the analysis.

1. The fit:

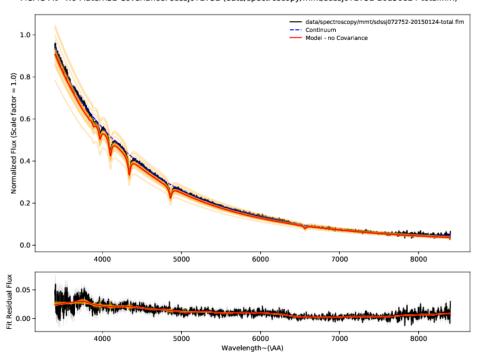
All the plots are stored in <spec basename>_mcmc.pdf in the output directory that is printed as you run the WDmodel fitter (default: out/<object name>/<spec basename>/.



The first plots show you the bottom line - the fit of the model (red) to the data - the observed photometry and spectroscopy (black). Residuals for both are shown in the lower panel. The model parameters inferred from the data are shown in the legend of the spectrum plot. Draws from the posterior are shown in orange. The number of these is customizable with --ndraws. Observational uncertainties are shown as shaded grey regions.

If both of these don't look reasonable, then the inferred parameters are probably meaningless. You should look at why the model is not in good agreement with the data. We've found this tends to happen if there's a significant flux excess at some wavelengths, indicating a companion or perhaps variability.

2. Spectral flux calibration errors:

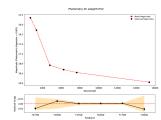


MCMC Fit - No Matern32 Covariance: sdssj072752 (data/spectroscopy/mmt/sdssj072752-20150124-total.flm)

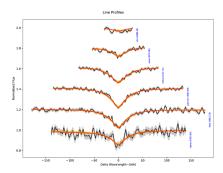
The WDmodel package uses a Gaussian process to model correlated flux calibration errors in the spectrum. These arise from

a variety of sources (flat-fielding, background subtraction, extraction of the 2D spectrum with a polynomial, telluric feature removal, and flux calibration relative to some other spectrophotometric standard, which in turn is probably only good to a few percent). However, most of the processes that generate these errors would cause smooth and continuous deformations to the observed spectrum, and a single stationary covariance kernel is a useful and reasonable way to model the effects. The choice of kernel is customizable (--covtype, with default Matern32 which has proven more than sufficient for data from four different spectrographs with very different optical designs).

The residual plot shows the difference between the spectrum and best fit model without the Gaussian process applied. The residuals therefore show our estimate of the flux calibration errors and the Gaussian process model for them.

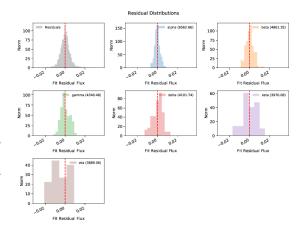


3. Hydrogen Balmer line diagnostics:



These plots illustrate the spectroscopic data and model specifically in the region of the Hydrogen Balmer lines. While the entire spectrum and all the photometry is fit simultaneously, we extract the Balmer lines, normalize their local continua to unity, and illustrate them separately here, offsetting each vertically a small amount for clarity.

With the exception of the SDSS autofit package used for their white dwarf analysis (which isn't public in any case), every white dwarf atmosphere fitting code takes the approach of only fitting the Hydrogen Balmer lines to determine model parameters. This includes our own proof-of-concept analysis of Cycle 20 data. The argument goes that the inferred model parameters aren't sensitive to reddening if the local continuum is divided out, and the line profiles determine the temperature and surface gravity.



In reality, reddening also changes the shape of the line profiles,

and to divide out the local continuum, a model for it had to be fit (typically a straight line across the line from regions defined "outside" the line profile wings). The properties of this local continuum *are* strongly correlated with reddening, and errors in the local continuum affect the inference of the model parameters, particularly at low S/N. This is the regime our program to observe faint white dwarfs operates in - low S/N with higher reddening. Any reasonable analysis of systematic errors should illustrate significant bias resulting from the naive analysis in the presence of correlated errors.

In other words, the approach doesn't avoid the problem, so much as shove it under a rug with the words "nuisance parameters" on top. This is why we adopted the more complex forward modeling approach in the WDmodel package. Unfortunately, Balmer profile fits are customary in the field, so after we forward model the data, we make a simple polynomial fit to the continuum (using our best understanding of what SDSS' autofit does), and extract out the Balmer lines purely for this visualization. This way the polynomial continuum model does have no affect on the inference, and if it goes wrong and the Balmer line profile plots look wonky, it doesn't actually matter.

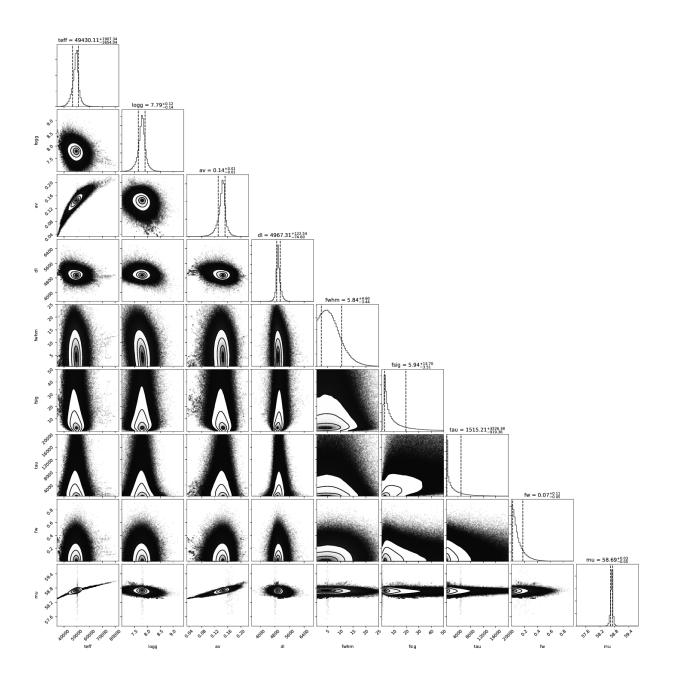
If you mail the author about these plots, he will get annoyed and grumble about you, and probably reply with snark. He had no particular desire to even include these plots.

4. Posterior Distributions:

A corner plot of the posterior distribution. If the model and data are not in good agreement, then this is a good place to look. If you are running with --samptype=ensemble (the default), you might consider --samptype=pt --ntemps=5 --nwalkers=100 --nprod=5000 --thin 10 to better sample the posterior, and map out any multi-modality.

5. Output Files:

This table describes the output files produced by the fitter.



| File | Description |
|--|---|
| <pre><spec basename="">_inputs.hdf5</spec></pre> | All inputs to fitter and visualization module. Restored on |
| | resume |
| <pre><spec basename="">_params.json</spec></pre> | Initial guess parameters. Refined by minuit if notskipminuit |
| <pre><spec basename="">_minuit.pdf</spec></pre> | Plot of initial guess model, if refined by minuit |
| <pre><spec basename="">_mcmc.hdf5</spec></pre> | Full Markov Chain - positions, log posterior, chain attributes |
| <pre><spec basename="">_mcmc.pdf</spec></pre> | Plot of model and data after MCMC |
| <pre><spec basename="">_result.json</spec></pre> | Summary of inferred model parameters, errors, uncertainties after |
| | MCMC |
| <pre><spec basename="">_spec_model.</spec></pre> | Table of the observed spectrum and inferred model spectrum |
| dat | |
| <pre><spec basename="">_phot_model.</spec></pre> | Table of the observed photometry and inferred model photometry |
| dat | |
| <pre><spec basename="">_full_model.</spec></pre> | Derived normalized SED of the object |
| hdf5 | |

See *Useful routines* for some useful routines to summarize fit results.

2.4 WDmodel

2.4.1 WDmodel package

WDmodel: Bayesian inference of white dwarf properties from spectra and photometry to establish spectrophotometric standards

Submodules

WDmodel.WDmodel module

DA White Dwarf Atmosphere Models and SED generator.

Model grid originally from J. Holberg using I. Hubeny's Tlusty code (v200) and custom Synspec routines, repackaged into HDF5 by G. Narayan.

class WDmodel.WDmodel (grid_file=None, grid_name=None, rvmodel=u'f99')
 Bases: object

DA White Dwarf Atmosphere Model and SED generator

Base class defines the routines to generate and work with DA White Dwarf model spectra. Requires a grid file of DA White Dwarf atmospheres. This grid file is included along with the package - TlustyGrids.hdf5 - and is the default.

Parameters

- **grid_file** (str, optional) Filename of the HDF5 grid file to read. See \(\bar{WDmodel.io.read_model_grid()} \) for format of the grid file. Default is \(\text{TlustyGrids.hdf5}, \text{included with the } \bar{WDmodel} \text{package}. \)
- **grid_name** (*str*, *optional*) Name of the HDF5 group containing the white dwarf model atmosphere grids in grid_file. Default is default

• rvmodel ({'ccm89','od94','f99','custom'}, optional) - Specify parametrization of the reddening law. Default is 'f99'.

| rvmodel | parametrization | |
|----------|--|--|
| 'ccm89' | Cardelli, Clayton and Mathis (1989, ApJ, 345, 245) | |
| 'od94' | O'Donnell (1994, ApJ, 422, 158) | |
| 'f99' | Fitzpatrick (1999, PASP, 111, 63) | |
| 'custom' | Custom law from Jay Holberg (email, 20180424) | |

lines

dictionary mapping Hydrogen Balmer series line names to line number, central wavelength in Angstrom, approximate line width and continuum region width around line. Used to extract Balmer lines from spectra for visualization.

Type dict

grid file

Filename of the HDF5 grid file that was read.

Type str

_grid_name

Name of the HDF5 group containing the white dwarf model atmosphere

Type str

_wave

Array of model grid wavelengths in Angstroms, sorted in ascending order

Type array-like

_ggrid

Array of model grid surface gravity values in dex, sorted in ascending order

Type array-like

_tgrid

Array of model grid temperature values in Kelvin, sorted in ascending order

Type array-like

_nwave

Size of the model grid wavelength array, _wave

Type int

_ngrav

Size of the model grid surface gravity array, _ggrid

Type int

$_\mathtt{ntemp}$

Size of the model grid temperature array, _tgrid

Type int

_flux

Array of model grid fluxes, shape (_nwave, _ntemp, _ngrav)

Type array-like

lwave

Array of model grid log10 wavelengths for interpolation

```
Type array-like

_lflux
Array of model grid log10 fluxes for interpolation, shape (_ntemp, _ngrav, _nwave)

Type array-like
_law

Type extinction function corresponding to rvmodel

Returns out
Return type WDmodel.WDmodel instance
Raises ValueError - If the supplied rvmodel is unknown
```

Notes

Virtually none of the attributes should be used directly since it is trivially possible to break the model by redefining them. Access to them is best through the functions connected to the models.

A custom user-specified grid file can be specified. See <code>WDmodel.io.read_model_grid()</code> for the format of the grid file.

Uses scipy.interpolate.RegularGridInterpolator to interpolate the models.

The class contains various convenience methods that begin with an underscore (_) that will not be imported by default. These are intended for internal use, and do not have the sanity checking and associated overhead of the public methods.

```
__WDmodel___init___rvmodel (rvmodel=u'f99')
__WDmodel___init__tlusty (grid_file=None, grid_name=None)
__call___ (teff, logg, wave=None, log=False, strict=True)
Returns the model flux given teff and logg at wavelengths wave

Wraps WDmodel.WDmodel.WDmodel.__get__model() adding checking of inputs.
```

Parameters

- **teff** (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- wave (array-like, optional) Desired wavelengths at which to compute the model atmosphere flux. If not supplied, the full model wavelength grid is returned.
- log (bool, optional) Return the log10 flux, rather than the flux
- **strict** (bool, optional)—If strict, teff and logg out of model grid range raise a ValueError, otherwise raise a RuntimeWarning and set teff, logg to the nearest grid value.

Returns

- wave (array-like) Valid output wavelengths
- flux (array-like) Interpolated model flux at teff, logg and wavelengths wave

Raises ValueError - If teff or logg are out of range of model grid and strict is True or if there are any invalid wavelengths, or the requested wavelengths to do not overlap with the model grid

Notes

Unlike the corresponding private methods, the public methods implement checking of the inputs and returns the wavelengths in addition to the flux. Internally, we only use the private methods as the inputs only need to be checked once, and their state is not altered anywhere after.

```
__init__ (grid_file=None, grid_name=None, rvmodel=u'f99')
x.__init__(...) initializes x; see help(type(x)) for signature
```

```
_custom_extinction(wave, av, rv=3.1, unit=u'aa')
```

Return the extinction for av, rv at wavelengths wave for the custom reddening law defined by J. Holberg

Mimics the interface provided by $WDmodel.WDmodel.WDmodel._law$ to calculate the extinction as a function of wavelength (in Angstroms), A_{λ} .

Parameters

- wave (array-like) Array of wavelengths in Angstrom at which to compute extinction, sorted in ascending order
- **av** (float) Extinction in the V band, A_V
- rv (float, optional) Fixed to 3.1 for J. Holberg's custom reddening law

Returns out - Extinction at wavelengths wave for av and rv

Return type array-like

Notes

av should be >= 0.

_extract_from_indices (w, f, ZE, df=None)

Extracts slices of multiple arrays for the same set of indices.

Convenience function to extract elements of wavelength w, flux f and optionally flux uncertainty df using indices ZE

Parameters

- w (array-like) Wavelength array from which to extract indices ZE
- f (array-like) Flux array from which to extract indices ZE
- ZE (array-like) indices to extract
- **df** (None or array-like, optional) If array-like, extracted elements of this array are also returned

Returns

- w (array-like) elements of input wavelength array at indices ZE
- f (array-like) elements of input flux array at indices ZE
- [df] (array-like) elements of input flux uncertainty array at indices ZE if optional input df is supplied

```
_extract_spectral_line(w, f, line, df=None)
```

Extracts slices of multiple arrays corresponding to a hydrogen Balmer line

Convenience function to extract elements of wavelength w, flux f and optionally flux uncertainty df for a hydrogen Balmer line. Wraps WDmodel.WDmodel.WDmodel.get_line_indices() and

WDmodel.WDmodel.wDmodel._extract_from_indices(), both of which have their own reasons for existence as well.

Parameters

- w (array-like) Wavelength array from which to extract elements corresponding to hydrogen Balmer line
- **f** (array-like) Flux array from which to extract elements corresponding to hydrogen Balmer line
- line ({'alpha', 'beta', 'gamma', 'delta', 'zeta', 'eta'}) Name of hydrogen Balmer line to extract. Properties are pre-defined in WDmodel. WDmodel._lines
- **df** (None or array-like, optional) If array-like, extracted elements of this array are also returned

Returns

- w (array-like) elements of input wavelength array for hydrogen Balmer feature line
- f (array-like) elements of input flux array for hydrogen Balmer feature line
- [df] (array-like) elements of input flux uncertainty array for hydrogen Balmer feature line if optional input df is supplied

Notes

Same as WDmodel.WDmodel.WDmodel.extract_spectral_line() without checking of inputs and therefore corresponding overhead. Used internally.

_get_full_obs_model(teff, logg, av, fwhm, wave, rv=3.1, log=False, pixel_scale=1.0)

Returns the observed model flux given teff, logg, av, rv, fwhm (for Gaussian instrumental broadening) at wavelengths, wave as well as the full SED.

Convenience function that does the same thing as <code>WDmodel.WDmo</code>

Uses $\mbox{WDmodel.WDmodel.get_model}$ () to get the unreddened model, and reddens it with $\mbox{WDmodel.WDmodel.WDmodel.reddening}$ () and convolves it with a Gaussian kernel using scipy.ndimage.filters.gaussian_filter1d()

Parameters

- **teff** (*float*) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- **av** (float) Extinction in the V band, A_V
- **fwhm** (float) Instrumental FWHM in Angstrom
- wave (array-like) Desired wavelengths at which to compute the model atmosphere flux.
- \mathbf{rv} (float, optional) The reddening law parameter, R_V , the ration of the V band extinction A_V to the reddening between the B and V bands, E(B-V). Default is 3.1, appropriate for stellar SEDs in the Milky Way.
- log (bool, optional) Return the log10 flux, rather than the flux (what's actually interpolated)

• pixel_scale (float, optional) – Jacobian of the transformation between wavelength in Angstrom and pixels. In principle, this should be a vector, but virtually all spectral reduction packages resample the spectrum onto a uniform wavelength scale that is close to the native pixel scale of the spectrograph. Default is 1.

Returns

- flux (array-like) Interpolated model flux at teff, logg with reddening parametrized by av, rv and broadened by a Gaussian kernel defined by fwhm at wavelengths wave
- mod (numpy.recarray with dtype=[('wave', '<f8'), ('flux', '<f8')]) Full model SED at teff, logg with reddening parametrized by av, rv

Notes

fwhm and pixel_scale must be > 0

classmethod _get_indices_in_range(wave, WA, WB, W0=None)

Returns indices of wavelength between blue and red wavelength limits and the central wavelength

Parameters

- wave (array-like) Wavelengths array from which to extract indices
- **WA** (float) blue limit of wavelengths to extract
- **WB** (float) red limit of wavelenghts to extract
- WO (float or None, optional) None or a central wavelength of range [WA, WB] to return. If None, the central wavelength is computed, else the input is simply returned.

Returns

- **W0** (*float*) central wavelength of range [WA, WB]
- **ZE** (array-like) indices of wave in range [WA, WB]

_get_line_indices(wave, line)

Returns the central wavelength and indices of wavelength corresponding to a hydrogen Balmer line

Parameters

- wave (array-like) Wavelengths array from which to extract indices
- line ({'alpha', 'beta', 'gamma', 'delta', 'zeta', 'eta'}) Name of hydrogen Balmer line to extract. Properties are pre-defined in WDmodel. WDmodel.wDmodel._lines

Returns

- W0 (float) central wavelength of line
- **ZE** (array-like) indices of wave of line

Notes

No checking of input - will throw KeyError if line is not accepted value

```
_get_model (teff, logg, wave=None, log=False)
```

Returns the model flux given teff and logg at wavelengths wave

Simple 3-D interpolation of model grid. Computes unreddened, unnormalized, unconvolved, interpolated model flux. Uses scipy.interpolate.RegularGridInterpolator to generate the interpolated model. This output has been tested against WDmodel.WDmodel.WDmodel._get_model_nosp().

Parameters

- **teff** (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- wave (array-like, optional) Desired wavelengths at which to compute the model atmosphere flux. If not supplied, the full model wavelength grid is returned.
- log (bool, optional) Return the log10 flux rather than the flux.

Returns flux – Interpolated model flux at teff, logg and wavelengths wave.

Return type array-like

Notes

Inputs teff, logg and wave must be within the bounds of the grid. See WDmodel.

```
_get_model_nosp(teff, logg, wave=None, log=False)
```

Returns the model flux given teff and logg at wavelengths wave

Simple 3-D interpolation of model grid. Computes unreddened, unnormalized, unconvolved, interpolated model flux. Not used, but serves as check of output of interpolation of scipy.interpolate. RegularGridInterpolator output.

Parameters

- **teff** (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- wave (array-like, optional) Desired wavelengths at which to compute the model atmosphere flux. If not supplied, the full model wavelength grid is returned.
- log (bool, optional) Return the log10 flux, rather than the flux.

Returns flux – Interpolated model flux at teff, logg and wavelengths wave

Return type array-like

Notes

Inputs teff, logg and wave must be within the bounds of the grid. See WDmodel.

This restriction is not imposed here for performance reasons, but is implicitly set by routines that call this method. The user is expected to verify this condition if this method is used outside the context of the WDmodel package. Caveat emptor.

_get_obs_model (teff, logg, av, fwhm, wave, rv=3.1, log=False, pixel_scale=1.0)

Returns the observed model flux given teff, logg, av, rv, fwhm (for Gaussian instrumental broadening) and wavelengths wave

Uses WDmodel.WDmodel.WDmodel._get_model() to get the unreddened model, and reddens it with WDmodel.WDmodel.WDmodel.reddening() and convolves it with a Gaussian kernel using scipy.ndimage.filters.gaussian_filter1d()

Parameters

- **teff** (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- **av** (float) Extinction in the V band, A_V
- fwhm (float) Instrumental FWHM in Angstrom
- wave (array-like) Desired wavelengths at which to compute the model atmosphere flux.
- **rv** (float, optional) The reddening law parameter, R_V , the ration of the V band extinction A_V to the reddening between the B and V bands, E(B-V). Default is 3.1, appropriate for stellar SEDs in the Milky Way.
- log (bool, optional) Return the log10 flux, rather than the flux (what's actually interpolated)
- pixel_scale (float, optional) Jacobian of the transformation between wavelength in Angstrom and pixels. In principle, this should be a vector, but virtually all spectral reduction packages resample the spectrum onto a uniform wavelength scale that is close to the native pixel scale of the spectrograph. Default is 1.

Returns flux – Interpolated model flux at teff, logg with reddening parametrized by av, rv and broadened by a Gaussian kernel defined by fwhm at wavelengths wave

Return type array-like

Notes

fwhm and pixel_scale must be > 0

```
_get_red_model (teff, logg, av, wave, rv=3.1, log=False)
```

Returns the reddened model flux given teff, logg, av, rv at wavelengths wave

Uses WDmodel.WDmodel.WDmodel.get_model() to get the unreddened model, and reddens it with WDmodel.WDmodel.WDmodel.reddening()

Parameters

- **teff** (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- av (float) Extinction in the V band, A_V
- wave (array-like) Desired wavelengths at which to compute the model atmosphere flux.
- \mathbf{rv} (float, optional) The reddening law parameter, R_V , the ration of the V band extinction A_V to the reddening between the B and V bands, E(B-V). Default is 3.1, appropriate for stellar SEDs in the Milky Way.

• log (bool, optional) - Return the log10 flux, rather than the flux (what's actually interpolated)

Returns flux – Interpolated model flux at teff, logg with reddening parametrized by av, rv at wavelengths wave

Return type array-like

classmethod _wave_test(wave)

Raises an error if wavelengths are not valid

Parameters wave (array-like) - Array of wavelengths to test for validity

Raises ValueError - If wavelength array is empty, has negative values, or is not monotonic

```
extinction (wave, av, rv=3.1)
```

Return the extinction for av, rv at wavelengths wave

Uses the extinction function corresponding to the rvmodel parametrization set as WDmodel. WDmodel. UDmodel. UDmod

Parameters

- wave (array-like) Array of wavelengths in Angstrom at which to compute extinction, sorted in ascending order
- av (float) Extinction in the V band, A_V
- rv(float, optional) The reddening law parameter, R_V , the ration of the V band extinction A_V to the reddening between the B and V bands, E(B-V). Default is 3.1, appropriate for stellar SEDs in the Milky Way.

Returns out – Extinction at wavelengths wave for av and rv

Return type array-like

Notes

av should be >= 0.

extract_spectral_line (w, f, line, df=None)

Extracts slices of multiple arrays corresponding to a hydrogen Balmer line

Convenience function to extract elements of wavelength w, flux f and optionally flux uncertainty df for a hydrogen Balmer line. Wraps WDmodel.WDmodel.WDmodel.extract_spectral_line() adding checking of inputs.

Parameters

- **w** (array-like) Wavelength array from which to extract elements corresponding to hydrogen Balmer line
- **f** (array-like) Flux array from which to extract elements corresponding to hydrogen Balmer line
- line ({'alpha', 'beta', 'gamma', 'delta', 'zeta', 'eta'}) Name of hydrogen Balmer line to extract. Properties are pre-defined in WDmodel. WDmodel.wDmodel.lines
- **df** (None or array-like, optional) If array-like, extracted elements of this array are also returned

Returns

- w (array-like) elements of input wavelength array for hydrogen Balmer feature line
- f (array-like) elements of input flux array for hydrogen Balmer feature line
- [df] (array-like) elements of input flux uncertainty array for hydrogen Balmer feature line if optional input df is supplied

Raises ValueError – If line is not one of the first six of the Balmer series or If wavelengths are invalid of If there's a difference in shape of any of the arrays

get_model (teff, logg, wave=None, log=False, strict=True)

Returns the model flux given teff and logg at wavelengths wave

Wraps WDmodel.WDmodel.WDmodel._get_model() adding checking of inputs.

Parameters

- **teff** (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- wave (array-like, optional) Desired wavelengths at which to compute the model atmosphere flux. If not supplied, the full model wavelength grid is returned.
- log (bool, optional) Return the log10 flux, rather than the flux
- **strict** (bool, optional) If strict, teff and logg out of model grid range raise a ValueError, otherwise raise a RuntimeWarning and set teff, logg to the nearest grid value.

Returns

- wave (array-like) Valid output wavelengths
- flux (array-like) Interpolated model flux at teff, logg and wavelengths wave

Raises ValueError - If teff or logg are out of range of model grid and strict is True or if there are any invalid wavelengths, or the requested wavelengths to do not overlap with the model grid

Notes

Unlike the corresponding private methods, the public methods implement checking of the inputs and returns the wavelengths in addition to the flux. Internally, we only use the private methods as the inputs only need to be checked once, and their state is not altered anywhere after.

get_obs_model (teff, logg, av, fwhm, rv=3.1, wave=None, log=False, strict=True, pixel_scale=1.0)
Returns the observed model flux given teff, logg, av, rv, fwhm (for Gaussian instrumental broadening) and wavelengths wave

Uses WDmodel.WDmodel.WDmodel.get_red_model() to get the reddened model and convolves it with a Gaussian kernel using scipy.ndimage.filters.gaussian_filter1d()

Parameters

- teff (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- **av** (float) Extinction in the V band, A_V
- **fwhm** (float) Instrumental FWHM in Angstrom

- \mathbf{rv} (float, optional) The reddening law parameter, R_V , the ration of the V band extinction A_V to the reddening between the B and V bands, E(B-V). Default is 3.1, appropriate for stellar SEDs in the Milky Way.
- wave (array-like, optional) Desired wavelengths at which to compute the model atmosphere flux. If not supplied, the full model wavelength grid is returned.
- log (bool, optional) Return the log10 flux, rather than the flux (what's actually interpolated)
- **strict** (bool, optional) If strict, teff and logg out of model grid range raise a ValueError, otherwise raise a RuntimeWarning and set teff, logg to the nearest grid value.
- pixel_scale (float, optional) Jacobian of the transformation between wavelength in Angstrom and pixels. In principle, this should be a vector, but virtually all spectral reduction packages resample the spectrum onto a uniform wavelength scale that is close to the native pixel scale of the spectrograph. Default is 1.

Returns

- wave (array-like) Valid output wavelengths
- flux (array-like) Interpolated model flux at teff, logg with reddening parametrized by av, rv broadened by a Gaussian kernel defined by fwhm at wavelengths wave

Notes

fwhm and pixel_scale must be > 0

get_red_model (teff, logg, av, rv=3.1, wave=None, log=False, strict=True)
Returns the reddened model flux given teff, logg, av, rv at wavelengths wave

Uses WDmodel.WDmodel.WDmodel.get_model() to get the unreddened model, and reddens it with WDmodel.WDmodel.WDmodel.reddening()

Parameters

- **teff** (float) Desired model white dwarf atmosphere temperature (in Kelvin)
- logg (float) Desired model white dwarf atmosphere surface gravity (in dex)
- av (float) Extinction in the V band, A_V
- rv (float, optional) The reddening law parameter, R_V , the ration of the V band extinction A_V to the reddening between the B and V bands, E(B-V). Default is 3.1, appropriate for stellar SEDs in the Milky Way.
- wave (array-like, optional) Desired wavelengths at which to compute the model atmosphere flux. If not supplied, the full model wavelength grid is returned.
- log (bool, optional) Return the log10 flux, rather than the flux (what's actually interpolated)
- **strict** (bool, optional) If strict, teff and logg out of model grid range raise a ValueError, otherwise raise a RuntimeWarning and set teff, logg to the nearest grid value.

Returns

• wave (array-like) – Valid output wavelengths

• flux (array-like) – Interpolated model flux at teff, logg with reddening parametrized by av, rv at wavelengths wave

```
Raises ValueError – If av < 0 or rv not in [1.7, 5.1]
```

```
reddening (wave, flux, av, rv=3.1)
```

Redden a 1-D spectrum with extinction

Uses the extinction function corresponding to the rvmodel parametrization set in $WDmodel.WDmodel.WDmodel.WDmodel.WDmodel.wDmodel.init_rvmodel() to calculate the extinction as a function of wavelength (in Angstroms), <math>A_{\lambda}$.

Parameters

- wave (array-like) Array of wavelengths in Angstrom at which to compute extinction, sorted in ascending order
- flux (array-like) Array of fluxes at wave at which to apply extinction
- av (float) Extinction in the V band, A_V
- \mathbf{rv} (float, optional) The reddening law parameter, R_V , the ration of the V band extinction A_V to the reddening between the B and V bands, E(B-V). Default is 3.1, appropriate for stellar SEDs in the Milky Way.

Returns out – The reddened spectrum

Return type array-like

Notes

av and flux should be $\geq = 0$.

WDmodel.covariance module

Parametrizes the noise of the spectrum fit using a Gaussian process.

```
class WDmodel.covariance.WDmodel_CovModel(errscale, covtype=u'Matern32', coveps=1e-12)

Bases: object
```

Parametrizes the noise of the spectrum fit using a Gaussian process.

This class models the covariance of the spectrum fit using a stationary Gaussian process conditioned on the spectrum flux residuals and spectrum flux uncertainties. The class allows the kernel of the Gaussian process to be set in a single location. A few different stationary kernels are supported. These choices are defined in celerite.terms.

Parameters

- **errscale** (*float*) Chracteristic scale of the spectrum flux uncertainties. The kernel amplitude hyperparameters are reported as fractions of this number. If the spectrum flux is rescaled, this must be set appropriately to get the correct uncertainties. The *WDmodel* package uses the median of the spectrum flux uncertainty internally.
- covtype ({'Matern32', 'SHO', 'Exp', 'White}) The model to use to parametrize the covariance. Choices are defined in celerite.terms All choices except 'White' parametrize the covariance using a stationary kernel with a characteristic amplitude fsig and scale tau + a white noise component with amplitude fw. Only the white noise component is used to condition the Gaussian process if covtype is 'White'. If not specified or unknown, 'Matern32' is used and a RuntimeWarning is raised.

```
Matern32Term is used to approximate a Matern32 kernel with precision coveps.
            The default is 1e-12. Ignored if any other covtype is specified.
errscale
    The input errscale
        Type float
covtype
    The input covtype
        Type str
coveps
    The input coveps
        Type float
ndim
    The dimensionality of kernel used to parametrize the covariance
         Type int
_k1
    The non-trivial stationary component of the kernel
         Type None or a term instance from celerite.terms
_k2
    The white noise component of the kernel
        Type celerite.terms.JitterTerm
_logQ
    1/sqrt(2) - only set if covtype is 'SHO'
        Type float, conditional
    Returns
```

Return type A WDmodel.covariance.WDmodel_CovModelinstance

Notes

Virtually none of the attributes should be used directly since it is trivially possible to break the model by redefining them. Access to them is best through the functions connected to the models.

• coveps (float) - If covtype is 'Matern32' a celerite.terms.

```
__init__ (errscale, covtype=u'Matern32', coveps=1e-12)
x.__init__(...) initializes x; see help(type(x)) for signature

getgp (wave, flux_err, fsig, tau, fw)
Return the celerite.GP instance
```

Precomputes the covariance matrix of the Gaussian process specified by the functional form of the stationary kernel and the current values of the hyperparameters. Wraps celerite.GP.

Parameters

- wave (array-like, optional) Wavelengths at which to condition the Gaussian process
- **flux_err** (array-like) Flux uncertainty array on which to condition the Gaussian process

- **fsig** (float) The fractional amplitude of the non-trivial stationary kernel. The true amplitude is scaled by WDmodel.covariance.WDmodel_CovModel._errscale
- tau (float) The characteristic length scale of the non-trivial stationary kernel.
- **fw** (float) The fractional amplitude of the white noise component of the kernel. The true amplitude is scaled by WDmodel.covariance.WDmodel_CovModel.errscale

Returns gp – The Gaussian process with covariance matrix precomputed at the location of the

Return type celerite. GP instance

Notes

fsig, tau and fw all must be > 0. This constraint is not checked here, but is instead imposed by the samplers/optimizers used in the <code>WDmodel.fit</code> methods, and by bounds used to construct the <code>WDmodel.likelihood.WDmodel_Likelihood</code> instance using the <code>WDmodel.likelihood.setup_likelihood()</code> method.

lnlikelihood(wave, res, flux_err, fsig, tau, fw)

Return the log likelihood of the Gaussian process

Conditions the Gaussian process specified by the functional form of the stationary kernel and the current values of the hyperparameters on the data, and computes the log likelihood. Wraps $celerite.GP.log_likelihood()$.

Parameters

- wave (array-like, optional) Wavelengths at which to condition the Gaussian process
- **res** (array-like) Flux residual array on which to condition the Gaussian process. The kernel parametrization assumes that the mean model has been subtracted off.
- **flux_err** (array-like) Flux uncertaintyarray on which to condition the Gaussian process
- fsig (float) The fractional amplitude of the non-trivial stationary kernel.

 The true amplitude is scaled by WDmodel.covariance.WDmodel_CovModel.

 _errscale
- tau (float) The characteristic length scale of the non-trivial stationary kernel.
- fw (float) The fractional amplitude of the white noise component of the kernel.

 The true amplitude is scaled by WDmodel.covariance.WDmodel_CovModel.

 _errscale

Returns Inlike – The log likelihood of the Gaussian process conditioned on the data.

Return type float

See also:

```
getgp()
```

```
predict (wave, res, flux_err, fsig, tau, fw, mean_only=False)
    Return the prediction for the Gaussian process
```

Conditions the Gaussian process specified by the parametrized with the functional form of the stationary kernel and the current values of the hyperparameters on the data, and computes returns the prediction at the same location as the data. Wraps celerite.GP.predict().

Parameters

- wave (array-like, optional) Wavelengths at which to condition the Gaussian process
- **res** (array-like) Flux residual array on which to condition the Gaussian process. The kernel parametrization assumes that the mean model has been subtracted off.
- **flux_err** (array-like) Flux uncertaintyarray on which to condition the Gaussian process
- fsig (float) The fractional amplitude of the non-trivial stationary kernel.

 The true amplitude is scaled by WDmodel.covariance.WDmodel_CovModel.

 _errscale
- tau (float) The characteristic length scale of the non-trivial stationary kernel.
- fw (float) The fractional amplitude of the white noise component of the kernel.

 The true amplitude is scaled by WDmodel.covariance.WDmodel_CovModel.

 _errscale
- mean_only (bool, optional) Return only the predicted mean, not the covariance matrix

Returns

- wres (*array-like*) The prediction of the Gaussian process conditioned on the data at the same location i.e. the model.
- **cov** (*array-like*, *optional*) The computed covariance matrix of the Gaussian process using the parametrized stationary kernel evaluated at the locations of the data.

See also:

getgp()

WDmodel.fit module

Core data processing and fitting/sampling routines

```
WDmodel.fit.blotch_spectrum(spec, linedata)

Automagically remove cosmic rays and gaps from spectrum
```

Parameters

- **spec** (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- linedata (numpy.recarray) The observations of the spectrum corresponding to the hydrogen Balmer lines. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8'), ('line_mask', 'i4'), ('line_ind', 'i4')] Produced by orig_cut_lines()

Return type numpy.recarray

Notes

Some spectra have nasty cosmic rays or gaps in the data. This routine does a reasonable job blotching these by Wiener filtering the spectrum, marking features that differ significantly from the local variance in the region, and replace them with the filtered values. The hydrogen Balmer lines are preserved, so if your gap/cosmic ray lands on a line it will not be filtered. Additionally, filtering has edge effects, and these data are preserved as well. If you do blotch the spectrum, it is highly recommended that you use the bluelimit and redlimit options to trim the ends of the spectrum. Note that the spectrum will be rejected if it has flux or flux errors that are not finite or below zero. This is often the case with cosmic rays and gaps, so you will likely have to do some manual removal of these points.

YOU SHOULD PROBABLY PRE-PROCESS YOUR DATA YOURSELF BEFORE FITTING IT AND NOT BE LAZY! THIS ROUTINE ONLY EXISTS TO FIT QUICK LOOK SPECTRUM AT THE TELESCOPE, BEFORE FINAL REDUCTIONS!

```
WDmodel.fit.fit_model (spec, phot, model, covmodel, pbs, params, objname, outdir, specfile, phot_dispersion=0.0, samptype=u'ensemble', ascale=2.0, ntemps=1, nwalk-ers=300, nburnin=50, nprod=1000, everyn=1, thin=1, pool=None, resume=False, redo=False)
```

Core routine that models the spectrum using the white dwarf model and a Gaussian process with a stationary kernel to account for any flux miscalibration, sampling the posterior using a MCMC.

Parameters

- **spec** (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- phot (numpy.recarray) The photometry of objname with dtype=[('pb',
 'str'), ('mag', '<f8'), ('mag_err', '<f8')]</pre>
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- **pbs** (dict) Passband dictionary containing the passbands corresponding to phot.pb' and generated by WDmodel.passband.get_pbmodel().
- params (dict) A parameter dict such as that produced by WDmodel.io. read_params()
- objname (str) object name used to save output with correct name
- outdir (str) controls where the chain file s written
- **specfile** (str) Used in the title, and to set the name of the outfile
- phot_dispersion (float, optional) Excess photometric dispersion to add in quadrature with the photometric uncertainties phot.mag_err. Use if the errors are grossly underestimated. Default is 0.
- samptype ({ 'ensemble', 'pt', 'gibbs'}) Which sampler to use. The default is ensemble.
- ascale (float) The proposal scale for the sampler. Default is 2.
- ntemps (int) The number of temperatures to run walkers at. Only used if samptype is in {'pt', 'gibbs'} and set to 1. for ensemble. See a short summary review for details. Default is 1.
- nwalkers (int) The number of Goodman and Weare walkers. Default is 300.
- **nburnin** (*int*) The number of steps to discard as burn-in for the Markov-Chain. Default is 500.

- nprod (int) The number of production steps in the Markov-Chain. Default is 1000.
- **everyn** (*int*, *optional*) If the posterior function is evaluated using only every nth observation from the data, this should be specified. Default is 1.
- **thin** (*int*) Only save every thin steps to the output Markov Chain. Useful, if brute force way of reducing correlation between samples.
- **pool** (None or :py:class`emcee.utils.MPIPool`) If running with MPI, the pool object is used to distribute the computations among the child process
- resume (bool) If True, restores state and resumes the chain for another nprod iterations.
- **redo** (bool) If True, and a chain file and state file exist, simply clobbers them.

Returns

- **free_param_names** (*list*) names of parameters that were fit for. Names correspond to keys in params and the order of parameters in samples.
- **samples** (*array-like*) The flattened Markov Chain with the parameter positions. Shape is (ntemps*nwalkers*nprod, nparam)
- **samples_Inprob** (*array-like*) The flattened log of the posterior corresponding to the positions in samples. Shape is (ntemps*nwalkers*nprod, 1)
- everyn (int) Specifies sampling of the data used to compute the posterior. Provided in case we are using resume to continue the chain, and this value must be restored from the state file, rather than being supplied as a user input.
- **shape** (*tuple*) Specifies the shape of the un-flattened chain. (ntemps, nwalkers, nprod, nparam) Provided in case we are using resume to continue the chain, and this value must be restored from the state file, rather than being supplied as a user input.

Raises RuntimeError - If resume is set without the chain having been run in the first place.

Notes

Uses an Ensemble MCMC (implemented by emcee) to generate samples from the posterior. Does a short burn-in around the initial guess model parameters - either minuit or user supplied values/defaults. Model parameters may be frozen/fixed. Parameters can have bounds limiting their range. Then runs a full production change. Chain state is saved after every 100 production steps, and may be continued after the first 100 steps if interrupted or found to be too short. Progress is indicated visually with a progress bar that is written to STDOUT.

See also:

```
WDmodel.likelihood WDmodel.covariance
```

WDmodel.fit.fix_pos (pos, free_param_names, params)

Ensures that the initial positions of the emcee walkers are out of bounds

Parameters

- **pos** (array-like) starting positions of all the walkers, such as that produced by utils.sample_ball
- **free_param_names** (*iterable*) names of parameters that are free to float. Names must correspond to keys in params.
- params (dict) A parameter dict such as that produced by WDmodel.io. read_params()

Returns pos – starting positions of all the walkers, fixed to guarantee that they are within bounds defined in params

Return type array-like

Notes

emcee.utils.sample_ball() creates random walkers that may be initialized out of bounds. These walkers get stuck as there is no step they can take that will make the change in loglikelihood finite. This makes the chain appear strongly correlated since all the samples of one walker are at a fixed location. This resolves the issue by assuming that the parameter value was within bounds to begin with. This routine does not do any checking of types, values or bounds. This check is done by \(\textit{WDmodel.io.get_params_from_argparse()} \) before the fit. If you setup the fit using an external code, you should check these values.

See also:

```
emcee.utils.sample_ball() WDmodel.io.get_params_from_argparse()
```

WDmodel.fit.get_fit_params_from_samples(param_names, samples, samples_lnprob, params, ntemps=1, nwalkers=300, nprod=1000, discard=5)

Get the marginalized parameters from the sample chain

Parameters

- param_names (list) names of parameters that were fit for. Names correspond to keys in params and the order of parameters in samples.
- **samples** (array-like) The flattened Markov Chain with the parameter positions. Shape is (ntemps*nwalkers*nprod, nparam)
- **samples_Inprob** (array-like) The flattened log of the posterior corresponding to the positions in samples. Shape is (ntemps*nwalkers*nprod, 1)
- params (dict) A parameter dict such as that produced by WDmodel.io. read params()
- ntemps (int) The number of temperatures chains were run at. Default is 1.
- **nwalkers** (*int*) The number of Goodman and Weare walkers used in the fit. Default is 300.
- **nprod** (*int*) The number of production steps in the Markov-Chain. Default is 1000.
- **discard** (*int*) percentage of nprod steps from the start of the chain to discard in analyzing samples

Returns

- **mcmc_params** (*dict*) The output parameter dictionary with updated parameter estimates, errors and a scale. params.
- **out_samples** (*array-like*) The flattened Markov Chain with the parameter positions with the first %discard tossed.
- out_samples_Inprob (array-like) The flattened log of the posterior corresponding to the positions in samples with the first %discard samples tossed.

See also:

```
fit_model()
```

```
WDmodel.fit.hyper_param_guess (spec, phot, model, pbs, params)

Makes a guess for the parameter mu after the initial fit by quick_fit_spec_model()
```

Parameters

- **spec** (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- phot (numpy.recarray) The photometry of objname with dtype=[('pb',
 'str'), ('mag', '<f8'), ('mag_err', '<f8')]</pre>
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- **pbs** (dict) Passband dictionary containing the passbands corresponding to phot.pb' and generated by WDmodel.passband.get_pbmodel().
- params (dict) A parameter dict such as that produced by WDmodel.io. read_params()

Returns out_params – The output parameter dictionary with an initial guess for mu

Return type dict

Notes

Uses the initial guess of parameters from the spectrum fit by <code>quick_fit_spec_model()</code> to construct an initial guess of the SED, and computes mu (which looks like a distance modulus, but also includes a normalization for the radius of the DA white dwarf, and it's radius) as the median difference between the observed and synthetic photometry.

```
WDmodel.fit.orig_cut_lines(spec, model)
```

Cut out the hydrogen Balmer spectral lines defined in $\begin{subarray}{l} WDmodel. \begin{subarray}{l} WDmodel. \begin{subarray}{l$

The masking of Balmer lines is basic, and not very effective at high surface gravity or low temperature, or in the presence of non hydrogen lines. It's used to get a roughly masked set of data suitable for continuum detection, and is effective in the context of our ground-based spectroscopic followup campaign for HST GO 12967 and 13711 programs.

Parameters

- **spec** (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator

Returns

- linedata (numpy.recarray) The observations of the spectrum corresponding to the hydrogen Balmer lines. Has dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8'), ('line_mask', 'i4'), (line_ind', 'i4')]
- continuumdata (numpy.recarray) The continuum data. Has dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]

Notes

Does a coarse cut to remove hydrogen absorption lines from DA white dwarf spectra The line centroids, and widths are fixed and defined with the model grid This is insufficient, and particularly at high surface gravity and

low temperatures the lines are blended. This routine is intended to provide a rough starting point for the process of continuum determination.

```
WDmodel.fit.polyfit_continuum(continuumdata, wave)
```

Fit a polynomial to the DA white dwarf continuum to normalize it - purely for visualization purposes

Parameters

- continuumdata (numpy.recarray) The continuum data. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')] Produced by running the spectrum through WDmodel.fit. orig_cut_lines() and extracting the pre-defined lines in the WDmodel.WDmodel.WDmodel.WDmodel.w
- wave (array-like) The full spectrum wavelength array on which to interpolate the continuum model

Return type numpy.recarray

Notes

Roughly follows the algorithm described by the SDSS SSPP for a global continuum fit. Fits a red side and blue side at 5500 A separately to get a smooth polynomial representation. The red side uses a degree 5 polynomial and the blue side uses a degree 9 polynomial. Then "splices" them together - I don't actually know how SDSS does this, but we simply assert the two bits are the same function - and fits the full continuum to a degree 9 polynomial.

WDmodel.fit.pre_process_spectrum(spec, bluelimit, redlimit, model, params, lamshift=0.0, vel=0.0, rebin=1, blotch=False, rescale=False)

Pre-process the spectrum before fitting

Parameters

- **spec** (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- bluelimit (None or float) Trim wavelengths bluer than this limit. Uses the bluest wavelength of spectrum if None
- redlimit (None or float) Trim wavelengths redder than this limit. Uses the reddest wavelength of spectrum if None
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- params (dict) A parameter dict such as that produced by WDmodel.io. read_params() Will be modified to adjust the spectrum normalization parameters dl limits if rescale is set
- lamshift (float, optional) Apply a flat wavelength shift to the spectrum. Useful if the target was not properly centered in the slit, and the shift is not correlated with wavelength. Default is 0.
- vel (float, optional) Apply a velocity shift to the spectrum. Default is 0.
- **rebin** (*int*, *optional*) Integer factor by which to rebin the spectrum. Default is 1 (no rebinning).

- **blotch** (bool, optional) Attempt to remove cosmic rays and gaps from spectrum. Only to be used for quick look analysis at the telescope.
- rescale (bool, optional) Rescale the spectrum to make the median noise ~1. Has no effect on fitted parameters except spectrum flux normalization parameter dl but makes residual plots, histograms more easily interpretable as they can be compared to an N(0, 1) distribution.

Return type numpy.recarray

See also:

```
orig_cut_lines() blotch_spectrum() rebin_spec_by_int_factor()
polyfit_continuum()
```

```
WDmodel.fit.quick_fit_spec_model(spec, model, params)
```

Does a quick fit of the spectrum to get an initial guess of the fit parameters

Uses iminuit to do a rough diagonal fit - i.e. ignores covariance. For simplicity, also fixed FWHM and Rv (even when set to be fit). Therefore, only teff, logg, av, dl are fit for (at most). This isn't robust, but it's good enough for an initial guess.

Parameters

- spec (numpy.recarray) The spectrum with dtype=[('wave', '<f8'),
 ('flux', '<f8'), ('flux_err', '<f8')]</pre>
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- params (dict) A parameter dict such as that produced by WDmodel.io. read_params()

Returns migrad_params – The output parameter dictionary with updated initial guesses stored in the value key. Same format as params.

Return type dict

Raises

- RuntimeError If all of teff, logg, av, dl are set as fixed there's nothing to fit
- RuntimeWarning If minuit.Minuit.migrad() or minuit.Minuit. hesse() indicate that the fit is unreliable

Notes

None of the starting values for the parameters maybe None EXCEPT c. This refines the starting guesses, and determines a reasonable value for c

```
WDmodel.fit.rebin_spec_by_int_factor (spec, f=1)
Rebins a spectrum by an integer factor f
```

Parameters

• **spec** (numpy.recarray) - The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]

• **f** (*int*, *optional*) – an integer factor to rebin the spectrum by. Default is 1 (no rebinning)

Notes

If the spectrum is not divisible by f, the edges are trimmed by discarding the remainder measurements from both ends. If the remainder itself is odd, the extra measurement is discarded from the blue side.

WDmodel.io module

I/O methods. All the submodules of the WDmodel package use this module for almost all I/O operations.

```
WDmodel.io._read_ascii (filename, **kwargs)
Read ASCII files
```

Read space separated ASCII file, with column names provided on first line (leading # optional). kwargs are passed along to numpy.genfromtxt(). Forces any string column data to be encoded in ASCII, rather than Unicode.

Parameters

- **filename** (str) Filename of the ASCII file. Column names must be provided on the first line.
- **kwargs** (*dict*) Extra options, passed directly to numpy.genfromtxt()

Returns out – Record array with the data. Field names correspond to column names in the file.

```
Return type numpy.recarray
```

See also:

```
numpy.genfromtxt()
WDmodel.io.copy params(params)
```

Returns a deep copy of a dictionary. Necessary to ensure that dictionaries that nest dictionaries are properly updated.

```
Parameters params (dict or Object) – Any python object for which a deepcopy needs to be created. Typically a parameter dictionary such as that from WDmodel.io.read_params()
```

Returns params – A deepcopy of the object

Return type Object

Notes

Simple wrapper around copy.deepcopy()

```
WDmodel.io.get filepath(infile)
```

Returns the full path to a file. If the path is relative, it is converted to absolute. If this file does not exist, it is treated as a file within the WDmodel package. If that file does not exist, an error is raised.

Parameters infile (str) – The name of the file to set the full path for

Returns pkgfile - The path to the file

Return type str

Raises IOError – If the infile could not be found at location or inside the WDmodel package.

WDmodel.io.get_options (args, comm)

Get command line options for the WDmodel fitter package

Parameters

- args (array-like) list of the input command line arguments, typically from sys. argv
- comm (None or mpi4py.mpi.MPI instance) Used to communicate options to all child processes if running with mpi

Returns

- args (Namespace) Parsed command line options
- **pool** (None or :py:class'emcee.utils.MPIPool') If running with MPI, the pool object is used to distribute the computations among the child process

Raises ValueError - If any input value is invalid

WDmodel.io.get_outfile (outdir, specfile, ext, check=False, redo=False, resume=False)

Formats the output directory, spectrum filename, and an extension into an output filename.

Parameters

- outdir (str) The output directory name for the output file
- **specfile** (str) The spectrum filename
- ext (str) The output file's extension
- check (bool, optional) If True, check if the output file exists
- redo (bool, optional) If False and the output file already exists, an error is raised
- resume (bool, optional) If False and the output file already exists, an error is raised

Returns outfile - The output filename

Return type str

Raises IOError - If check is True, redo and resume are False, and outfile exists.

Notes

We set the output file based on the spectrum name, since we can have multiple spectra per object.

If outdir is configured by set_objname_outdir_for_specfile() for specfile, it'll include the object name.

See also:

```
set_objname_outdir_for_specfile()
```

WDmodel.io.get_params_from_argparse(args)

Converts an argparse. Namespace into an ordered parameter dictionary.

Returns params – The parameter dictionary

```
Return type collections.OrderedDict
```

Raises RuntimeError — If format of argparse. Namespace is invalid. or If parameter is fixed but value is None. or If parameter value is out of bounds.

Notes

Assumes that the argument parser options were names

- <param> value : Value of the parameter (float or None)
- <param>_fix: Bool specifying if the parameter
- <param>_bounds : tuple with lower limit and upper limit

where ram> is one of WDmodel.io._PARAMETER_NAMES

See also:

```
WDmodel.io.read_params() WDmodel.io.get_options()
```

```
WDmodel.io.get_phot_for_obj(objname, filename)
```

Gets the measured photometry for an object from a photometry lookup table.

Parameters

- **objname** (str) Object name to look for photometry for
- **filename** (str) The spectrum FWHM lookup table filename

```
Returns phot - The photometry of objname with dtype=[('pb', 'str'), ('mag',
    '<f8'), ('mag_err', '<f8')]</pre>
```

Return type numpy.recarray

Raises

- RuntimeError If there are no matches in the photometry lookup file or if there are multiple matches for an object in the photometry lookup file
- ValueError If the photometry or the photometry uncertainty values are not finite or if the photometry uncertainties are less <= 0

Notes

The lookup file must be readable by read_phot()

The column name with the object name objname expected to be obj

If column names for magnitudes are named <passband>, the column names for errors in magnitudes in passband must be 'd'+<passband name>.

```
WDmodel.io.get_pkgfile(infile)
```

Returns the full path to a file inside the WDmodel package

Parameters infile (str) – The name of the file to set the full package filename for

Returns pkgfile – The path to the file within the package.

Return type str

Raises IOError – If the pkgfile could not be found inside the WDmodel package.

Notes

This allows the package to be installed anywhere, and the code to still determine the location to a file included with the package, such as the model grid file.

WDmodel.io.get_spectrum_resolution (specfile, spectable, fwhm=None, lamshift=None) Gets the measured FWHM from a spectrum lookup table.

Parameters

- **specfile** (*str*) The spectrum filename
- **spectable** (*str*) The spectrum FWHM lookup table filename
- **fwhm** (None or float, optional) If specified, this overrides the resolution provided in the lookup table. If None lookups the resultion from spectable.
- lamshift (None or float, optional) If specified, this overrides the wavelength shift provided in the lookup table. If None lookups the wavelength shift from spectable.

Returns

- **fwhm** (*float*) The FWHM of the spectrum file. This is typically used as an initial guess to the WDmodel. fit fitter routines.
- **lamshift** (*float*) The wavelength shift to apply additively to the spectrum. This is not a fit parameter, and is treated as an input

Raises RuntimeWarning – If the spectable cannot be read, or the specfile name indicates that this is a test, or if there are no or multiple matches for specfile in the spectable

Notes

If the specfile is not found, it returns a default resolution of 5 Angstroms, appropriate for the instruments used in our program.

Note that there there's some hackish internal name fixing since T. Matheson's table spectrum names didn't match the spectrum filenames.

WDmodel.io.make_outdirs (dirname, redo=False, resume=False)
Makes output directories

Parameters

- **dirname** (str) The output directory name to create
- redo (bool, optional) If False the directory will not be created if it already exists, and an error is raised
- resume (bool, optional) If False the directory will not be created if it already exists, and an error is raised

Returns None - If the output directory dirname is successfully created

Return type None

Raises

- IOError If the output directory exists
- OSError If the output directory could not be created

Notes

If the options are parsed by $get_options()$ then only one of redo or resume can be set, as the options are mutually exclusive. If redo is set, the fit is redone from scratch, while resume restarts the MCMC sampling from the last saved chain position.

WDmodel.io.read fit inputs(input file)

Read the fit input HDF5 file produced by write_fit_inputs() and return numpy.recarray instances with the data.

Parameters input_file (str) - The HDF5 fit inputs filename

Returns

- spec (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- **cont_model** (numpy.recarray) The continuuum model. Has the same structure as spec.
- linedata (numpy.recarray) The observations of the spectrum corresponding to the hydrogen Balmer lines. Has dtype=[('wave', '<f8'), ('flux', '<f8'), ('line_mask', 'i4')]
- continuumdata (numpy.recarray) Data used to generate the continuum model. Has the same structure as spec.
- phot (None or numpy.recarray) None or the photometry with dtype=[('pb', 'str'), ('mag', '<f8'), ('mag_err', '<f8')]
- fit_config (dict) -

Dictionary with various keys needed to configure the fitter

- rvmodel: {'ccm89','od94','f99', 'custom'} Parametrization of the reddening law.
- covtype: {'Matern32', 'SHO', 'Exp', 'White'}- kernel type used to
 parametrize the covariance
- coveps: float Matern32 kernel precision
- phot_dispersion: float Excess dispersion to add in quadrature with photometric uncertainties
- scale_factor: float Flux scale factor

Raises

- IOError If all the fit inputs could not be restored from the HDF5 input_file
- RuntimeWarning If the input_file includes a phot group, but the data cannot be loaded.

See also:

```
write fit inputs()
```

WDmodel.io.read_full_model(input_file)

Read the full SED model from an output file.

Parameters input_file (str) - Input HDF5 SED model filename

Return type numpy.recarray

Raises

- KeyError If any of wave, flux or flux_err is not found in the file
- ValueError If any value is not finite or if flux or flux_err have any values <= 0

WDmodel.io.read_mcmc(input_file)

Read the saved HDF5 Markov chain file and return samples, sample log probabilities and chain parameters

Parameters input_file (str) - The HDF5 Markov chain filename

Returns

- samples (array-like) The model parameter sample chain
- samples_Inprob (array-like) The log posterior corresponding to each of the samples
- chain_params (dict) -

The chain parameter dictionary

- param_names : list list of model parameter names
- samptype: { 'ensemble', 'pt', 'gibbs' } the sampler to use
- ntemps: int the number of chain temperatures
- nwalkers: int the number of Goodman & Ware walkers
- nprod: int the number of production steps of the chain
- ndim: int the number of model parameters in the chain
- thin: int the chain thinning if any
- everyn: int the sparse of spectrum sampling step size
- ascale: float the proposal scale for the sampler

Raises IOError - If a key in the fit_config output is missing

 $\verb|WDmodel.io.read_model_grid| (\textit{grid_file} = None, \textit{grid_name} = None)|$

Read the Tlusty/Hubeny grid file

Parameters

- grid_file (None or str) Filename of the Tlusty model grid HDF5 file. If None reads the TlustyGrids.hdf5 file included with the WDmodel package.
- **grid_name** (*None or str*) Name of the group name in the HDF5 file to read the grid from. If None uses default

Returns

- **grid_file** (*str*) Filename of the HDF5 grid file
- grid_name (str) Name of the group within the HDF5 grid file with the grid arrays
- wave (array-like) The wavelength array of the grid with shape (nwave,)
- ggrid (array-like) The surface gravity array of the grid with shape (ngrav,)
- tgrid (array-like) The temperature array of the grid with shape (ntemp,)
- flux (array-like) The DA white dwarf model atmosphere flux array of the grid. Has shape (nwave, ngrav, ntemp)

Notes

There are no easy command line options to change this deliberately because changing the grid file essentially changes the entire model, and should not be done lightly, without careful comparison of the grids to quantify differences.

See also:

```
WDmodel.WDmodel
```

```
WDmodel.io.read_params (param_file=None)
```

Read a JSON file that configures the default guesses and bounds for the parameters, as well as if they should be fixed.

Parameters param_file (str, optional) - The name of the input parameter file. If not the default file provided with the package, WDmodel_param_defaults.json, is read.

Returns params - The dictionary with the parameter values, bounds, scale and if fixed. See notes for more detailed information on dictionary format and WDmodel_param_defaults.json for an example file for param_file.

Return type dict

Notes

params is a dict the parameter names, as defined with WDmodel.io._PARAMETER_NAMES as keys

Each key must have a dictionary with keys:

- value: value
- fixed: a bool specifying if the parameter is fixed (True) or allowed to vary (False)
- scale: a scale parameter used to set the step size in this dimension
- bounds: An upper and lower limit on parameter values

The default bounds are set by the grids available for the DA White Dwarf atmospheres, and by reasonable plausible ranges for the other parameters. Don't muck with them unless you really have good reason to.

This routine does not do any checking of types, values or bounds. This is done by <code>WDmodel.io.get_params_from_argparse()</code> before the fit. If you setup the fit using an external code, you should check these values.

```
WDmodel.io.read_pbmap (filename, **kwargs)
    Read passband obsmode mapping table - wraps _read_ascii()
WDmodel.io.read_phot (filename, **kwargs)
    Read photometry - wraps _read_ascii()
WDmodel.io.read_reddening (filename, **kwargs)
    Read J. Holberg's custom reddening function - wraps _read_ascii()
WDmodel.io.read_spec (filename, **kwargs)
    Read a spectrum
```

Wraps _read_ascii(), adding testing of the input arrays to check if the elements are finite, and if the errors and flux are strictly positive.

Parameters

• **filename** (*str*) - Filename of the ASCII file. Must have columns wave, flux, flux_err

```
• kwargs (dict) - Extra options, passed directly to numpy.genfromtxt()
```

Return type numpy.recarray

Raises ValueError – If any value is not finite or if flux or flux_err have any values <= 0

See also:

```
numpy.genfromtxt() _read_ascii()
```

WDmodel.io.read_spectable (filename, **kwargs)

Read spectrum FWHM table - wraps _read_ascii()

WDmodel.io.set_objname_outdir_for_specfile (specfile, outdir=None, outroot=None, redo=False, resume=False, nocreate=False)

Sets the short human readable object name and output directory

Parameters

- **specfile** (str) The spectrum filename
- outdir (None or str, optional) The output directory name to create. If None this is set based on specfile
- outroot (None or str, optional) The output root directory under which to store the fits. If None the default is 'out'
- redo (bool, optional) If False the directory will not be created if it already exists, and an error is raised
- **resume** (bool, optional) If False the directory will not be created if it already exists, and an error is raised
- nocreate (bool, optional) If True then creation of output directories is not even attempted

Returns

- **objname** (*str*) The human readable object name based on the spectrum
- **dirname** (*str*) The output directory name created if successful

See also:

```
make_outdirs()
```

WDmodel.io.write_fit_inputs (spec, phot, cont_model, linedata, continuumdata, rvmodel, covtype, coveps, phot_dispersion, scale_factor, outfile)

Save all the inputs to the fitter to a file

This file is enough to resume the fit with the same input, redoing the output, or restoring from a failure.

Parameters

- **spec** (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- phot (None or numpy.recarray) None or the photometry with dtype=[('pb', 'str'), ('mag', '<f8'), ('mag_err', '<f8')]
- cont_model (numpy.recarray) The continuuum model. Must have the same structure as spec. Produced by WDmodel.fit.pre_process_spectrum(). Used by WDmodel.viz

- linedata (numpy.recarray) The observations of the spectrum corresponding to the hydrogen Balmer lines. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8'), ('line_mask', 'i4')] Produced by WDmodel.fit.pre_process_spectrum() Used by WDmodel.viz
- continuumdata (numpy.recarray) Data used to generate the continuum model. Must have the same structure as spec. Produced by WDmodel.fit. pre_process_spectrum()
- rvmodel ({'ccm89', 'od94', 'f99', 'custom'}) Parametrization of the reddening law. Used to initialize WDmodel.WDmodel.WDmodel() instance.
- covtype ({'Matern32', 'SHO', 'Exp', 'White'}) stationary kernel type used to parametrize the covariance in WDmodel.covariance.WDmodel_CovModel
- **coveps** (*float*) If covtype is 'Matern32' a celerite.terms. Matern32Term is used to approximate a Matern32 kernel with precision *coveps*.
- phot_dispersion (float, optional) Excess photometric dispersion to add in quadrature with the photometric uncertainties phot.mag_err in WDmodel. likelihood.WDmodel_Likelihood.
- **scale_factor** (*float*) Factor by which the flux must be scaled. Critical to getting the right uncertainties.
- outfile (str) Output HDF5 filename

Notes

The outputs are stored in a HDF5 file with groups

- spec storing the spectrum and scale_factor
- \bullet cont_model stores the continuum model
- linedata stores the hydrogen Balmer line data
- continuumdata stores the data used to generate cont_model
- fit_config stores covtype, coveps and rvmodel as attributes
- phot only created if phot is not None, stores phot, phot_dispersion

WDmodel.io.write_full_model(full_model, outfile)

Write the full SED model to an output file.

Parameters

- full_model (numpy.recarray) The SED model with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- outfile (str) Output HDF5 SED model filename

Notes

The output is written into a group model with datasets

- ullet wave: array-like the SED model wavelength
- flux: array-like the SED model flux
- flux_err: array-like the SED model flux uncertainty

```
WDmodel.io.write_params (params, outfile)
```

Dumps the parameter dictionary params to a JSON file

Parameters

- params (dict) A parameter dict such as that produced by WDmodel.io. read_params()
- **outfile** (str) Output filename to save the parameter dict as a JSON file.

Notes

params is a dict the parameter names, as defined with WDmodel.io._PARAMETER_NAMES as keys

Each key must have a dictionary with keys:

- value: value
- fixed: a bool specifying if the parameter is fixed (True) or allowed to vary (False)
- scale: a scale parameter used to set the step size in this dimension
- bounds: An upper and lower limit on parameter values

Any extra keys are simply written as-is JSON doesn't preserve ordering necessarily. This is imposed by WDmodel.io.read_params()

See also:

```
WDmodel.io.read_params()
```

```
WDmodel.io.write_phot_model (phot, model_mags, outfile)
```

Write the photometry, model photometry and residuals to an output file.

Parameters

- phot (None or numpy.recarray) None or the photometry with dtype=[('pb', 'str'), ('mag', '<f8'), ('mag_err', '<f8')]
- model_mags (None or numpy.recarray) The model magnitudes. Has dtype=[('pb', 'str'), ('mag', '<f8')]
- **outfile** (str) Output space-separated text filename

Notes

The data is saved to a space-separated ASCII text file with 6 decimal places of precision.

The order of the columns is

- pb: array-like the observation's passband
- mag: array-like the observed magnitude
- mag_err: array-like the observed magnitude uncertainty
- model_mag: array-like the model magnitude
- res mag: array-like the magnitude residual

${\tt WDmodel.io.write_spectrum_model} \ (spec, model_spec, outfile)$

Write the spectrum and the model spectrum and residuals to an output file.

Parameters

```
spec (numpy.recarray) - The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]</li>
model_spec (numpy.recarray) - The model spectrum. Has dtype=[('wave', '<f8'), ('flux', '<f8'), ('norm_flux', '<f8'), ('flux_err', '<f8')]</li>
```

• outfile (str) - Output space-separated text filename

Notes

The data is saved to a space-separated ASCII text file with 8 decimal places of precision.

The order of the columns is

- wave : array-like the spectrum wavelength
- flux: array-like the observed flux
- flux_err: array-like the observed flux uncertainty
- norm_flux: array-like the model flux without the Gaussian process covariance model
- model_flux: array-like the model flux
- model_flux_err: array-like the model flux uncertainty
- res_flux: array-like the flux residual

WDmodel.likelihood module

Classes defining the likelihood and the posterior probability of the model given the data

Bases: object

Classes defining the posterior probability of the model given the data

An instance of this class is used to store the data and model, and evaluate the likelihood and prior to compute the posterior.

Parameters

- **spec** (numpy.recarray) The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- phot (None or numpy.recarray) The photometry with dtype=[('pb',
 'str'), ('mag', '<f8'), ('mag_err', '<f8')]</pre>
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- **covmodel** (WDmodel.covariance.WDmodel_CovModel instance) The parametrized model for the covariance of the spectrum spec
- **pbs** (dict) Passband dictionary containing the passbands corresponding to phot.pb and generated by WDmodel.passband.get_pbmodel().
- Inlike (WDmodel_Likelihood instance) Instance of the likelihood function class, such as that produced by WDmodel.likelihood.setup likelihood()

- pixel_scale (float, optional) Jacobian of the transformation between wavelength in Angstrom and pixels. In principle, this should be a vector, but virtually all spectral reduction packages resample the spectrum onto a uniform wavelength scale that is close to the native pixel scale of the spectrograph. Default is 1.
- phot_dispersion (float, optional) Excess photometric dispersion to add in quadrature with the photometric uncertainties phot.mag_err. Use if the errors are grossly underestimated. Default is 0.

spec

```
The spectrum with dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
```

```
Type numpy.recarray
```

wave_scale

length of the wavelength array wave in Angstroms

```
Type float
```

phot

```
The photometry with dtype=[('pb', 'str'), ('mag', '<f8'), ('mag_err',
'<f8')]</pre>
```

```
Type None or numpy . recarray
```

model

The DA White Dwarf SED model generator

```
Type WDmodel.WDmodel.WDmodelinstance
```

covmodel

The parametrized model for the covariance of the spectrum spec

```
Type WDmodel.covariance.WDmodel CovModelinstance
```

pbs

Passband dictionary containing the passbands corresponding to phot.pb and generated by WDmodel. passband.get_pbmodel().

```
Type dict
```

_lnlike

Instance of the likelihood function class, such as that produced by WDmodel.likelihood. setup_likelihood()

```
Type WDmodel_Likelihoodinstance
```

pixel scale

Jacobian of the transformation between wavelength in Angstrom and pixels. In principle, this should be a vector, but virtually all spectral reduction packages resample the spectrum onto a uniform wavelength scale that is close to the native pixel scale of the spectrograph. Default is 1.

```
Type float
```

phot_dispersion

Excess photometric dispersion to add in quadrature with the photometric uncertainties phot.mag_err. Use if the errors are grossly underestimated. Default is 0.

Type float, optional

p0

initial values of all the model parameters, including fixed parameters

Type dict

Returns Inpost – It is this instance that is passed to the samplers/optimizers in the WDmodel.fit module. Those methods evaluate the posterior probability of the model parameters given the data.

Return type WDmodel_Posterior instance

Notes

Wraps celerite.modeling.Model.log_prior() which imposes a boundscheck and returns -inf. This is not an issue as the samplers used in the methods in WDmodel.fit.

```
__call__(theta, prior=False, likelihood=False)
```

Evalulates the log posterior of the model parameters given the data

Parameters

- theta (array-like) Vector of the non-frozen model parameters. The order of the parameters is defined by WDmodel_Likelihood.parameter_names.
- **prior** (bool, optional) Only return the value of the log prior given the model parameters
- **likelihood** (bool, optional) Only return the value of the log likelihood given the model parameters if the prior is finite

Returns Inpost – the log posterior of the model parameters given the data

```
Return type float
```

```
__init__ (spec, phot, model, covmodel, pbs, lnlike, pixel_scale=1.0, phot_dispersion=0.0)
x.__init__(...) initializes x; see help(type(x)) for signature
```

_lnprior()

Evalulates the log likelihood of the model parameters given the data.

Implements an Inprior function which imposes weakly informative priors on the model parameters.

Parameters theta (array-like) - Vector of the non-frozen model parameters. The order of the parameters is defined by WDmodel_Likelihood.parameter_names.

Returns Inprior – the log likelihood of the model parameters given the data

Return type float

Notes

The prior on av is the 'glos' prior

The prior on rv is a Gaussian with mean 3.1 and standard deviation 0.18. This is adopted from Schlafly et al., 2014 PS1 analysis. Note that they report 3.31, but they aren't really measuring E(B-V) with PS1. Their sigma should be consistent despite the different filter set.

The prior on fsig and fw - the fractional amplitudes of the non-trivial stationary and white components of the kernel used to parametrize the covariance is half-Cauchy since we don't want it to be less than zero

There is no explicit prior on tau i.e. a tophat prior, defined by the bounds

The fwhm has a lower bound set at the value below which the spectrum isn't being convolved anymore. We never run into this bound since real spectra have physical instrumental broadening. This prevents fwhm from going to zero for fitting poorly simulated spectra generated from simply resampling the model grid.

The prior on all other parameters are broad Gaussians

Wraps celerite.modeling.Model.log_prior() which imposes a boundscheck and returns -inf. This is not an issue as the samplers used in the methods in WDmodel.fit.

lnlike(theta)

Evalulates the log likelihood of the model parameters given the data.

Convenience function that can return the value of the likelihood even if the prior is not finite unlike <code>WDmodel_Posterior.__call__()</code> for debugging.

Parameters theta (array-like) - Vector of the non-frozen model parameters. The order of the parameters is defined by WDmodel_Likelihood.parameter_names.

Returns Inlike – the log likelihood of the model parameters given the data

Return type float

lnprior(theta)

Evalulates the log prior of the model parameters.

Convenience function that can return the value of the prior defined to make the interface consistent with the <code>WDmodel_Posterior.lnlike()</code> method. Just a thin wrapper around <code>WDmodel_Posterior._lnlike()</code> method. Just a thin wrapper around <code>WDmodel_Posterior._call__()</code>.

Parameters theta (array-like) - Vector of the non-frozen model parameters. The order of the parameters is defined by WDmodel_Likelihood.parameter_names.

Returns Inprior – the log prior of the model parameters

Return type float

WDmodel.likelihood.setup_likelihood(params)

Setup the form of the likelihood of the data given the model.

Parameters params (dict) - A parameter dictionary used to configure the WDmodel_Likelihood instance. The format of the dict is defined by WDmodel. io.read_params().

Returns Inlike – An instance of the likelihood function class.

Return type WDmodel.likelihood.WDmodel_Likelihood

WDmodel.main module

The WDmodel package is designed to infer the SED of DA white dwarfs given spectra and photometry. This main module wraps all the other modules, and their classes and methods to implement the alogrithm.

```
WDmodel.main.main(inargs=None)
```

Entry point for the WDmodel fitter package.

Parameters inargs (dict, optional) – Input arguments to configure the fit. If not specified sys.argv is used. inargs must be parseable by WDmodel.io.get_options().

Raises RuntimeError - If user attempts to resume the fit without having run it first

Notes

The package is structured into several modules and classes

| Module | Model Component |
|---------------------------------------|---------------------|
| WDmodel.io | I/O methods |
| WDmodel.WDmodel | SED generator |
| WDmodel.passband | Throughput model |
| WDmodel.covariance.WDmodel_CovModel | Noise model |
| WDmodel.likelihood.WDmodel_Likelihood | Likelihood function |
| WDmodel.likelihood.WDmodel_Posterior | Posterior function |
| WDmodel.fit | "Fitting" methods |
| WDmodel.viz | Viz methods |

This method implements our algorithm to infer the DA White Dwarf properties and construct the SED model given the data using the methods and classes listed above. Once the data is read, the model is configured, and the liklihood and posterior functions constructed, the fitter methods evaluate the model parameters given the data, using the sampler in <code>emcee.WDmodel.mossampler</code> provides an overloaded <code>emcee.PTSampler</code> with a more reliable auto-correlation estimate. Finally, the result is output along with various plots.

WDmodel.main.mpi_excepthook (excepttype, exceptvalue, traceback)

Overload sys.excepthook() when using mpi4py.MPI to terminate all MPI processes when an Exception is raised.

WDmodel.mossampler module

Overridden PTSampler with random Gibbs selection, more-reliable acor.

Original Author: James Guillochon for the mosfit package

Modified to update kwargs, docstrings for full compatibility with PTSampler by G. Narayan

WDmodel.passband module

Instrumental throughput models and calibration and synthetic photometry routines

WDmodel.passband.chop_syn_spec_pb(spec, model_mag, pb, model)

Trims the pysynphot bandpass pb to non-zero throughput, computes the zeropoint of the passband given the SED spec, and model magnitude of spec in the passband

Parameters

- **spec** (numpy.recarray) The spectrum. Typically a standard which has a known model_mag. This can be a real source such as Vega, BD+174708, or one of the three CALSPEC standards, or an idealized synthetic source such as AB. Must have dtype=[('wave', '<f8'), ('flux', '<f8')]
- model_mag (float) The apparent magnitude of the spectrum through the passband. The difference between the apparent magnitude and the synthetic magnitude is the synthetic zeropoint.
- **pb** (numpy.recarray) The passband transmission. Must have dtype=[('wave', '<f8'), ('throughput', '<f8')]
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator

Returns

- outpb (numpy.recarray) The passband transmission with zero throughput entries trimmed. Has dtype=[('wave', '<f8'), ('throughput', '<f8')]
- outzp (float) The synthetic zeropoint of the passband pb such that the source with spectrum spec will have apparent magnitude model_mag through pb. With the synthetic zeropoint computed, the synthetic magnitude of any source can be converted into an apparent magnitude and can be passed to WDmodel.passband.synphot().

See also:

```
WDmodel.passband.interp_passband() WDmodel.passband.synphot()
```

WDmodel.passband.get_model_synmags(model_spec, pbs, mu=0.0)

Computes the synthetic magnitudes of spectrum model_spec through the passbands pbs, and optionally applies a common offset, mu

Wrapper around WDmodel.passband.synphot().

Parameters

- **spec** (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8')]
- **pbs** (dict) Passband dictionary containing the passbands corresponding to phot.pb' and generated by WDmodel.passband.get_pbmodel().
- mu (float, optional) Common achromatic photometric offset to apply to the synthetic magnitudes in al the passbands. Would be equal to the distance modulus if model_spec were normalized to return the true absolute magnitude of the source.

Returns model_mags - The model magnitudes. Has dtype=[('pb', 'str'), ('mag',
 '<f8')]</pre>

Return type None or numpy.recarray

WDmodel.passband.get_pbmodel (pbnames, model, pbfile=None, mag_type=None, mag_zero=0.0)

Converts passband names pbnames into passband models based on the mapping of name to pysynphot obsmode strings in pbfile.

Parameters

- **pbnames** (array-like) List of passband names to get throughput models for Each name is resolved by first looking in pbfile (if provided) If an entry is found, that entry is treated as an obsmode for pysynphot. If the entry cannot be treated as an obsmode, we attempt to treat as an ASCII file. If neither is possible, an error is raised.
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator All the passbands are interpolated onto the wavelengths of the SED model.
- **pbfile** (*str*, *optional*) Filename containing mapping between pbnames and pysynphot obsmode string, as well as the standard that has 0 magnitude in the system (either "Vega" or "AB"). The obsmode may also be the fullpath to a file that is readable by pysynphot
- mag_type (str, optional) One of "vegamag" or "abmag" Used to specify the standard that has mag_zero magnitude in the passband. If magsys is specified in pbfile, that overrides this option. Must be the same for all passbands listed in pbname that do not have magsys specified in pbfile If pbnames require multiple mag_type, concatentate the output.
- mag_zero (float, optional) Magnitude of the standard in the passband If magzero is specified in pbfile, that overrides this option. Must be the same for all

passbands listed in pbname that do not have magzero specified in pbfile If pbnames require multiple mag_zero, concatentate the output.

Returns out – Output passband model dictionary. Has passband name pb from pbnames as key.

Return type dict

Raises RuntimeError - If a bandpass cannot be loaded

Notes

Each item of out is a tuple with

- pb: (numpy.recarray) The passband transmission with zero throughput entries trimmed. Has dtype=[('wave', '<f8'), ('throughput', '<f8')]
- transmission: (array-like) The non-zero passband transmission interpolated onto overlapping model wavelengths
- ind: (array-like) Indices of model wavelength that overlap with this passband
- zp : (float) mag_type zeropoint of this passband
- avgwave : (float) Passband average/reference wavelength

pbfile must be readable by WDmodel.io.read_pbmap() and must return a numpy.recarray
with 'dtype=[('pb', 'str'),('obsmode', 'str')]''

If there is no entry in pbfile for a passband, then we attempt to use the passband name pb as obsmode string as is.

Trims the bandpass to entries with non-zero transmission and determines the VEGAMAG/ABMAG zeropoint for the passband - i.e. zp that gives mag_Vega/AB=mag_zero in all passbands.

See also:

```
WDmodel.io.read_pbmap() WDmodel.passband.chop_syn_spec_pb()
```

WDmodel.passband.interp_passband(wave, pb, model)

Find the indices of the wavelength array wave, that overlap with the passband pb and interpolates the passband onto the wavelengths.

Parameters

- wave (array-like) The wavelength array. Must satisfy WDmodel.WDmod
- **pb** (numpy.recarray) The passband transmission. Must have dtype=[('wave', '<f8'), ('throughput', '<f8')]
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator

Returns

- **transmission** (*array-like*) The transmission of the passband interpolated on to overlapping elements of wave
- ind (array-like) Indices of wavelength wave that overlap with the passband pb. Produced by WDmodel.WDmodel.WDmodel.get_indices_in_range() Satisfies transmission.shape == wave[ind].shape

Notes

The passband pb is interpolated on to the wavelength arrray wave. wave is typically the wavelengths of a spectrum, and have much better sampling than passband transmission curves. Only the wavelengths wave that overlap the passband are taken, and the passband transmission is then linearly interpolated on to these wavelengths. This prescription has been checked against pysynphot to return synthetic magnitudes that agree to be < 1E-6, while WDmodel.passband.synphot() is very significantly faster than pysynphot. observation.Observation.effstim().

WDmodel.passband.synflux(spec, ind, pb)

Compute the synthetic flux of spectrum spec through passband pb

Parameters

- spec (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8')]
- ind (array-like) Indices of spectrum spec that overlap with the passband pb. Can be produced by WDmodel.passband.interp_passband()
- **pb** (array-like) The passband transmission. Must satisfy pb.shape == spec[ind].flux.shape

Returns flux – The normalized flux of the spectrum through the passband

Return type float

Notes

The passband is assumed to be dimensionless photon transmission efficiency.

Routine is intended to be a mch faster implementation of pysynphot.observation.Observation.effstim(), since it is called over and over by the samplers as a function of model parameters.

Uses numpy.trapz() for interpolation.

See also:

```
WDmodel.passband.interp_passband()
```

WDmodel.passband.synphot (spec, ind, pb, zp=0.0)

Compute the synthetic magnitude of spectrum spec through passband pb

Parameters

- spec (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8')]
- ind (array-like) Indices of spectrum spec that overlap with the passband pb. Can be produced by WDmodel.passband.interp_passband()
- **pb** (array-like) The passband transmission. Must satisfy pb.shape == spec[ind].flux.shape
- **zp** (float, optional) The zeropoint to apply to the synthetic flux

Returns mag – The synthetic magnitude of the spectrum through the passband

Return type float

See also:

```
\label{lower_model} \textit{WDmodel.passband.interp\_passband()} \\
```

WDmodel.viz module

Routines to visualize the DA White Dwarf model atmosphere fit

WDmodel.viz.plot_mcmc_line_fit (spec, linedata, model, cont_model, draws, balmer=None)
Plot a comparison of the normalized hydrogen Balmer lines of the spectrum and model

Note that we fit the full spectrum, not just the lines. The lines are extracted using a coarse continuum fit in <code>WDmodel.fit.pre_process_spectrum()</code>. This fit is purely cosmetic and in no way contributes to the likelihood. It's particularly useful to detect small velocity offsets or wavelength calibration errors.

Parameters

- spec (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux err', '<f8')]
- linedata (numpy.recarray) The observations of the spectrum corresponding to the hydrogen Balmer lines. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8'), ('line_mask', 'i4'), ('line_ind', 'i4')]
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- cont_model (numpy.recarray) The continuuum model. Must have the same structure as spec Produced by WDmodel.fit.pre_process_spectrum()
- **draws** (array-like) produced by plot_mcmc_spectrum_fit() see notes for content.
- balmer(array-like, optional)—list of Balmer lines to plot elements must be in range [1, 6] These correspond to the lines defined in WDmodel.WDmodel.WDmodel.lines. Default is range (1, 7)

Returns

- fig (matplotlib.figure.Figure instance) The output figure containing the line profile plot
- fig2 (matplotlib.figure.Figure instance) The output figure containing histograms of the line residuals

See also:

```
WDmodel.viz.plot_mcmc_spectrum_fit()
```

WDmodel.viz.plot_mcmc_model (spec, phot, linedata, scale_factor, phot_dispersion, objname, outdir, specfile, model, covmodel, cont_model, pbs, params, param_names, samples, samples_lnprob, covtype=u'Matern32', balmer=None, ndraws=21, everyn=1, savefig=False)

Make all the plots to visualize the full fit of the DA White Dwarf data

Wraps $plot_mcmc_spectrum_fit()$, $plot_mcmc_photometry_res()$, $plot_mcmc_spectrum_nogp_fit()$, $plot_mcmc_line_fit()$ and corner.corner() and saves all the plots to a combined PDF, and optionally individual PDFs.

Parameters

- spec (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- phot (None or numpy.recarray) The photometry. Must have dtype=[('pb',
 'str'), ('mag', '<f8'), ('mag_err', '<f8')]</pre>

- linedata (numpy.recarray) The observations of the spectrum corresponding to the hydrogen Balmer lines. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8'), ('line_mask', 'i4'), ('line_ind', 'i4')]
- scale_factor (float) factor by which the flux was scaled for y-axis label
- phot_dispersion (float, optional) Excess photometric dispersion to add in quadrature with the photometric uncertainties phot.mag_err. Use if the errors are grossly underestimated. Default is 0.
- **objname** (str) object name used to title plots
- outdir (str) controls where the plot is written out if savefig=True
- **specfile** (*str*) Used in the title, and to set the name of the outfile if savefig=True
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- **covmodel** (WDmodel.covariance.WDmodel_CovModel instance) The parametrized model for the covariance of the spectrum spec
- cont_model (numpy.recarray) The continuuum model. Must have the same structure as spec Produced by WDmodel.fit.pre_process_spectrum()
- **pbs** (dict) Passband dictionary containing the passbands corresponding to phot.pb and generated by WDmodel.passband.get_pbmodel().
- params (dict) dictionary of parameters with keywords value, fixed, scale, bounds for each. Same format as returned from WDmodel.io.read_params()
- param_names (array-like) Ordered list of free parameter names
- **samples** (array-like) Samples from the flattened Markov Chain with shape (N, len(param_names))
- **samples_Inprob** (*array-like*) Log Posterior corresponding to samples from the flattened Markov Chain with shape (N,)
- covtype ({'Matern32', 'SHO', 'Exp', 'White'}) stationary kernel type used to parametrize the covariance in WDmodel.covariance.WDmodel_CovModel
- balmer (array-like, optional) list of Balmer lines to plot elements must be in range [1, 6] These correspond to the lines defined in WDmodel.WDmodel.WDmodel._lines. Default is range (1, 7)
- ndraws (int, optional) Number of draws to make from the Markov Chain to overplot. Higher numbers provide a better sense of the uncertainty in the model at the cost of speed and a larger, slower to render output plot.
- **everyn** (*int*, *optional*) If the posterior function was evaluated using only every nth observation from the data, this should be specified to visually indicate the observations used
- **savefig** (bool) if True, save the individual figures

Returns

• model_spec (numpy.recarray) - The model spectrum. Has dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8'), ('norm_flux', '<f8')] and same shape as input spec. The norm_flux attribute has the model flux without the Gaussian process prediction applied.

- SED_model (numpy.recarray) The SED model spectrum. Has dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- model_mags (None or numpy.recarray) If there is observed photometry, this contains the model magnitudes. Has dtype=[('pb', 'str'), ('mag', '<f8')]

WDmodel.viz.plot_mcmc_photometry_res (objname, phot, phot_dispersion, model, pbs, draws)
Plot the observed DA white dwarf photometry as well as the "best-fit" model magnitudes

Parameters

- objname (str) object name used to title plots
- phot (None or numpy.recarray) The photometry. Must have dtype=[('pb',
 'str'), ('mag', '<f8'), ('mag_err', '<f8')]</pre>
- phot_dispersion (float, optional) Excess photometric dispersion to add in quadrature with the photometric uncertainties phot.mag_err. Use if the errors are grossly underestimated. Default is 0.
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- **pbs** (dict) Passband dictionary containing the passbands corresponding to phot.pb and generated by WDmodel.passband.get_pbmodel().
- draws (array-like) produced by plot_mcmc_spectrum_fit() see notes for content.

Returns

- fig (matplotlib.figure.Figure instance) The output figure
- mag_draws (*array-like*) The magnitudes corresponding to the parameters draws from the Markov Chain used in fig

Notes

Each element of mag_draws contains

- wres the difference between the observed and synthetic magnitudes
- model_maqs the model magnitudes corresponding to the current model parameters
- mu the flux normalization parameter that must be added to the model mags

See also:

```
WDmodel.viz.plot_mcmc_spectrum_fit()
```

WDmodel.viz.plot_mcmc_spectrum_fit (spec, objname, specfile, scale_factor, model, covmodel, result, param_names, samples, ndraws=21, everyn=1)

Plot the spectrum of the DA White Dwarf and the "best fit" model

The full fit parametrizes the covariance model using a stationary Gaussian process as defined by <code>WDmodel.covariance.WDmodel_CovModel</code>. The posterior function constructed in <code>WDmodel.likelihood.WDmodel_Posterior</code> is evaluated by the sampler in the <code>WDmodel.fit.fit_model()</code> method. The median value is reported as the best-fit value for each of the fit parameters in <code>WDmodel.likelihood.WDmodel_Likelihood.parameter_names</code>.

Parameters

- spec (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- objname (str) object name used to title plots
- outdir (str) controls where the plot is written out if save=True
- specfile (str) Used in the title, and to set the name of the outfile if save=True
- scale_factor (float) factor by which the flux was scaled for y-axis label
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- **covmodel** (WDmodel.covariance.WDmodel_CovModel instance) The parametrized model for the covariance of the spectrum spec
- result (dict) dictionary of parameters with keywords value, fixed, scale, bounds for each. Same format as returned from WDmodel.io.read_params()
- param_names (array-like) Ordered list of free parameter names
- **samples** (array-like) Samples from the flattened Markov Chain with shape (N, len(param_names))
- ndraws (int, optional) Number of draws to make from the Markov Chain to overplot. Higher numbers provide a better sense of the uncertainty in the model at the cost of speed and a larger, slower to render output plot.
- **everyn** (*int*, *optional*) If the posterior function was evaluated using only every nth observation from the data, this should be specified to visually indicate the observations used.

Returns

- fig (matplotlib.figure.Figure instance) The output figure
- draws (array-like) The actual draws from the Markov Chain used in fig

Notes

It's faster to draw samples from the posterior in one location, and pass along the same samples to all the methods in WDmodel.viz.

Consequently, most require draws as an input. This makes all the plots connected, and none will return if an error is thrown here, but this is the correct behavior as all of them are visualizing one aspect of the same fit.

Each element of draws contains

- smoothedmod the model spectrum
- wres the prediction from the Gaussian process
- \bullet wres_err the diagonal of the covariance matrix for the prediction from the Gaussian process
- full mod the full model SED, in order to compute the synthetic photometry
- out_draw the dictionary of model parameters from this draw. Same format as result.

WDmodel.viz.plot_mcmc_spectrum_nogp_fit (spec, objname, specfile, scale_factor, cont_model, draws, covtype=u'Matern32', everyn=1)

Plot the spectrum of the DA White Dwarf and the "best fit" model without the Gaussian process

Unlike plot_mcmc_spectrum_fit () this version does not apply the prediction from the Gaussian process to the spectrum model to match the observed spectrum. This visualization is useful to indicate if the Gaussian process - i.e. the kernel choice covtype used to parametrize the covariance is - is appropriate.

Parameters

- spec (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- objname (str) object name used to title plots
- outdir (str) controls where the plot is written out if save=True
- **specfile** (str) Used in the title, and to set the name of the outfile if save=True
- scale_factor (float) factor by which the flux was scaled for y-axis label
- cont_model (numpy.recarray) The continuuum model. Must have the same structure as spec Produced by WDmodel.fit.pre_process_spectrum()
- draws (array-like) produced by plot_mcmc_spectrum_fit() see notes for content.
- covtype ({'Matern32', 'SHO', 'Exp', 'White'}) stationary kernel type used to parametrize the covariance in WDmodel.covariance.WDmodel_CovModel
- **everyn** (*int*, *optional*) If the posterior function was evaluated using only every nth observation from the data, this should be specified to visually indicate the observations used.

Returns fig – The output figure

Return type matplotlib.figure.Figure instance

See also:

```
WDmodel.viz.plot_mcmc_spectrum_fit()
```

WDmodel.viz.plot_minuit_spectrum_fit (spec, objname, outdir, specfile, scale_factor, model, result, save=True)

Plot the MLE fit of the spectrum with the model, assuming uncorrelated noise.

Parameters

- spec (numpy.recarray) The spectrum. Must have dtype=[('wave', '<f8'), ('flux', '<f8'), ('flux_err', '<f8')]
- objname (str) object name used to title plots
- outdir (str) controls where the plot is written out if save=True
- specfile (str) Used in the title, and to set the name of the outfile if save=True
- scale_factor (float) factor by which the flux was scaled for y-axis label
- model (WDmodel.WDmodel.WDmodel instance) The DA White Dwarf SED model generator
- result (dict) dictionary of parameters with keywords value, fixed, scale, bounds for each. Same format as returned from WDmodel.io.read params()
- **save** (bool) if True, save the file

Returns fig

Return type matplotlib.figure.Figure instance

Notes

The MLE fit uses iminuit.Minuit.migrad() to fit the spectrum with the model. This fit doesn't try to account for the covariance in the data, and is not expected to be great - just fast, and capable of setting a reasonable initial guess. If it is apparent from the plot that this fit is very far off, refine the initial guess to the fitter.

$\mathsf{CHAPTER}\,3$

Indices and tables

- genindex
- modindex
- search

Python Module Index

W

```
WDmodel, 14
WDmodel.covariance, 25
WDmodel.fit, 28
WDmodel.io, 35
WDmodel.likelihood, 45
WDmodel.main, 48
WDmodel.mossampler, 49
WDmodel.passband, 49
WDmodel.viz, 53
WDmodel.WDmodel, 14
```

62 Python Module Index

Index

| Symbols | _get_obs_model() (WDmodel.WDmodel.WDmodel |
|--|--|
| _WDmodelinitrvmodel() (WD- | method), 20 |
| model.WDmodel.WDmodel method), 16 | _get_red_model() (WDmodel.WDmodel.WDmodel |
| _WDmodelinittlusty() (WD- | method), 21 |
| model.WDmodel.WDmodel method), 16 | _ggrid (WDmodel.WDmodel.WDmodel attribute), 15 |
| call() (WDmodel.WDmodel.WDmodel method), | _grid_file (WDmodel.WDmodel at- |
| 16 | tribute), 15 |
| call() (WDmodel.likelihood.WDmodel_Posterior | _grid_name (WDmodel.WDmodel at- |
| method), 47 | tribute), 15 |
| init() (WDmodel.WDmodel.WDmodel method), | _k1 (WDmodel.covariance.WDmodel_CovModel at- |
| 17 | tribute), 26 |
| init() (WDmodel.covariance.WDmodel_CovMode | _{el} _k2 (WDmodel.covariance.WDmodel_CovModel at- |
| method), 26 | tribute), 26 |
| init() (WDmodel.likelihood.WDmodel_Posterior | _law (WDmodel.WDmodel attribute), 16 |
| method), 47 | _lflux (WDmodel.WDmodel.WDmodel attribute), 16 |
| _coveps (WDmodel.covariance.WDmodel_CovModel | _lines (WDmodel.WDmodel.WDmodel attribute), 15 |
| attribute), 26 | _lnlike (WDmodel.likelihood.WDmodel_Posterior at- |
| _covtype (WDmodel.covariance.WDmodel_CovModel | tribute), 46 |
| attribute), 26 | _lnprior()(WDmodel.likelihood.WDmodel_Posterior |
| _custom_extinction() (WD- | method), 47 |
| model.WDmodel.WDmodel method), 17 | _logQ (WDmodel.covariance.WDmodel_CovModel at- |
| $_{\tt errscale} ({\it WD model.covariance.WD model_CovModel} $ | tribute), 26 |
| attribute), 26 | _lwave (WDmodel.WDmodel.WDmodel attribute), 15 |
| _extract_from_indices() (WD- | ndim (WDmodel.covariance.WDmodel_CovModel at- |
| model.WDmodel.WDmodel method), 17 | tribute), 26 |
| _extract_spectral_line() (WD- | _ngrav (WDmodel.WDmodel.WDmodel attribute), 15 |
| model.WDmodel.WDmodel method), 17 | _ntemp(WDmodel.WDmodel.WDmodel attribute), 15 |
| _flux (WDmodel.WDmodel.WDmodel attribute), 15 | _nwave (WDmodel.WDmodel.WDmodel attribute), 15 |
| _get_full_obs_model() (WD- | _read_ascii() (in module WDmodel.io), 35 |
| model.WDmodel.WDmodel method), 18 | _tgrid(WDmodel.WDmodel.WDmodel attribute), 15 |
| _get_indices_in_range() (WD- | _wave (WDmodel.WDmodel attribute), 15 |
| model.WDmodel.WDmodel class method), 19 | _wave_test() (WDmodel.WDmodel.WDmodel class method), 22 |
| _get_line_indices() (WD- | В |
| model.WDmodel.WDmodel method), 19 | Ь |
| _get_model() (WDmodel.WDmodel.WDmodel | blotch_spectrum() (in module WDmodel.fit), 28 |
| method), 19 | |
| _get_model_nosp() (WD- | C |
| model.WDmodel.WDmodel method), 20 | chop_syn_spec_pb() (in module WD-model.passband), 49 |

| copy_params() (in module WDmodel.io), 35 | M | |
|--|---|--|
| covmodel (WDmodel.likelihood.WDmodel_Posterior attribute), 46 | <pre>main() (in module WDmodel.main), 48 make_outdirs() (in module WDmodel.io), 38 model (WDmodel.likelihood.WDmodel_Posterior)</pre> | |
| E | attribute), 46 | |
| extinction() (WDmodel.WDmodel.WDmodel method), 22 | <pre>mpi_excepthook() (in module WDmodel.main), 49</pre> | |
| extract_spectral_line() (WD- | 0 | |
| model.WDmodel.WDmodel method), 22 | orig_cut_lines() (in module WDmodel.fit), 32 | |
| F | P | |
| fit_model() (in module WDmodel.fit), 29 fix_pos() (in module WDmodel.fit), 30 | p0 (WDmodel.likelihood.WDmodel_Posterior attribute), 46 | |
| G | pbs (WDmodel.likelihood.WDmodel_Posterior at- tribute), 46 | |
| <pre>get_filepath() (in module WDmodel.io), 35 get_fit_params_from_samples() (in module</pre> | phot (WDmodel.likelihood.WDmodel_Posterior at- tribute), 46 | |
| <pre>WDmodel.fit), 31 get_model() (WDmodel.WDmodel.WDmodel method), 23</pre> | phot_dispersion (WD-model.likelihood.WDmodel_Posterior at-tribute), 46 | |
| <pre>get_model_synmags() (in module WD- model.passband), 50</pre> | pixel_scale (WDmodel.likelihood.WDmodel_Posterio.attribute), 46 | |
| <pre>get_obs_model() (WDmodel.WDmodel.WDmodel</pre> | <pre>plot_mcmc_line_fit() (in module WDmodel.viz),</pre> | |
| <pre>get_options() (in module WDmodel.io), 36 get_outfile() (in module WDmodel.io), 36 get_params_from_argparse() (in module WD-</pre> | <pre>plot_mcmc_model() (in module WDmodel.viz), 53 plot_mcmc_photometry_res() (in module WD-</pre> | |
| model.io), 36 get_pbmodel() (in module WDmodel.passband), 50 | plot_mcmc_spectrum_fit() (in module WD-model.viz),55 | |
| <pre>get_phot_for_obj() (in module WDmodel.io), 37 get_pkgfile() (in module WDmodel.io), 37</pre> | plot_mcmc_spectrum_nogp_fit() (in module WDmodel.viz), 56 | |
| <pre>get_red_model() (WDmodel.WDmodel.WDmodel</pre> | plot_minuit_spectrum_fit() (in module WD-model.viz), 57 | |
| get_spectrum_resolution() (in module WD-model.io), 38 | polyfit_continuum() (in module WDmodel.fit), 33 pre_process_spectrum() (in module WD- | |
| getgp() (WDmodel.covariance.WDmodel_CovModel method), 26 | model.fit), 33 predict() (WDmodel.covariance.WDmodel_CovModel | |
| Н | method), 27 | |
| hyper_param_guess() (in module WDmodel.fit), 31 | Q | |
| I | <pre>quick_fit_spec_model() (in module WD- model.fit), 34</pre> | |
| <pre>interp_passband() (in module WD- model.passband), 51</pre> | R | |
| L | <pre>read_fit_inputs() (in module WDmodel.io), 39 read_full_model() (in module WDmodel.io), 39</pre> | |
| <pre>lnlike() (WDmodel.likelihood.WDmodel_Posterior method), 48</pre> | read_mcmc() (in module WDmodel.io), 40 read_model_grid() (in module WDmodel.io), 40 | |
| lnlikelihood() (WD- | read_params() (in module WDmodel.io), 41 | |
| model.covariance.WDmodel_CovModel | read_pbmap() (in module WDmodel.io), 41 | |
| <pre>method), 27 lnprior() (WDmodel.likelihood.WDmodel_Posterior</pre> | <pre>read_phot() (in module WDmodel.io), 41 read_reddening() (in module WDmodel.io), 41</pre> | |
| method), 48 | read_spec() (in module WDmodel.io), 41 | |

64 Index

```
read_spectable() (in module WDmodel.io), 42
rebin_spec_by_int_factor() (in module WD-
        model.fit), 34
                    (WD model. WD model. WD model
reddening()
        method), 25
S
set_objname_outdir_for_specfile()
                                            (in
        module WDmodel.io), 42
setup_likelihood()
                                           WD-
                          (in
                                module
        model.likelihood), 48
       (WDmodel.likelihood.WDmodel_Posterior
spec
                                            at-
        tribute), 46
synflux() (in module WDmodel.passband), 52
synphot () (in module WDmodel.passband), 52
wave_scale(WDmodel.likelihood.WDmodel_Posterior
        attribute), 46
WDmodel (class in WDmodel.WDmodel), 14
WDmodel (module), 1, 14
WDmodel.covariance (module), 25
WDmodel.fit (module), 28
WDmodel.io (module), 35
WDmodel.likelihood (module), 45
WDmodel.main (module), 48
WDmodel.mossampler (module), 49
WDmodel.passband (module), 49
WDmodel.viz (module), 53
WDmodel.WDmodel (module), 14
WDmodel_CovModel (class in WDmodel.covariance),
WDmodel_Posterior (class in WDmodel.likelihood),
write_fit_inputs() (in module WDmodel.io), 42
write_full_model() (in module WDmodel.io), 43
write_params() (in module WDmodel.io), 44
write_phot_model() (in module WDmodel.io), 44
write_spectrum_model() (in module
        model.io), 44
```

Index 65