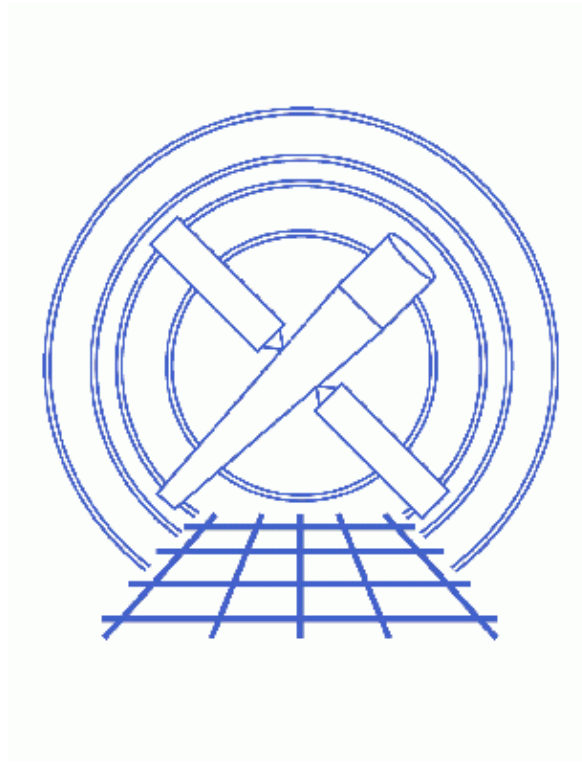


# Fitting Grating Data



## Sherpa Threads (CIAO 3.4)

# Table of Contents

- *Getting Started*
- *Reading the Spectrum Files*
- *Building the Instrument Responses*
- *Filtering the Data*
- *Defining the Source and Background Models*
- *Examining Method & Statistic Settings*
- *Fitting*
- *Examining Fit Results*
- *Saving and Quitting the Session*
- *History*
- *Images*
  - ◆ *Plotting the four orders*
  - ◆ *Filtering the datasets*
  - ◆ *Results of simultaneous fit*
  - ◆ *Fit and residuals for the HEG –1 order spectrum*

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# Fitting Grating Data

## *Sherpa Threads*

### Overview

*Last Update:* 1 Dec 2006 – reviewed for CIAO 3.4: no changes

#### *Synopsis:*

This thread provides a general introduction to fitting grating data in *Sherpa*. Loading and filtering data are covered, as well as defining instrument and source models.

Users working with HRC–S/LETG grating data will also find the [Fitting Multiple Orders of HRC–S/LETG Data thread](#) helpful for their analysis.

*Proceed to the [HTML](#) or [hardcopy \(PDF: \[A4\]\(#\) / \[letter\]\(#\)\)](#) version of the thread.*

---

## Getting Started

*Sample ObsID used:* 459 (HETG/ACIS–S, 3C 273)

The files used in this example were created by following several of the [CIAO Grating threads](#):

- [Obtain Grating Spectra from HETG/ACIS–S Data](#)
- [Compute HETG/ACIS–S Grating ARFs](#)
- [Grouping PHA Data before Fitting](#)

Here is a list of all the necessary files:

```
spectra:
459_heg_m1_bin10.pha
459_heg_p1_bin10.pha
459_meg_m1_bin10.pha
459_meg_p1_bin10.pha

gARFs:
459_heg_m1.arf
459_heg_p1.arf
459_meg_m1.arf
459_meg_p1.arf
```

The spectrum that will be used in this session has been binned by a factor of 10.

Users may also choose to run the [Create Grating RMFs for ACIS Observations thread](#). Creating observation–specific gRMFs is optional, and is discussed further in the [Building the Instrument Responses section](#).

The data files are available in [sherpa.tar.gz](http://sherpa.tar.gz), as explained in the [Sherpa Getting Started thread](#).

---

## Reading the Spectrum Files

The data are input to *Sherpa* with the `data` command (a shorthand version of "`read_data`"):

```
sherpa> data 1 459_heg_m1_bin10.pha
The inferred file type is PHA.  If this is not what you want, please
specify the type explicitly in the data command.
Warning: could not find SYS_ERR column
WARNING: statistical errors specified in the PHA file.
        These are currently IGNORED.  To use them, type:
        READ ERRORS "<filename>[cols CHANNEL,STAT_ERR]" fitsbin
WARNING: backgrounds UP and DOWN are being read from this file,
        and are being combined into a single background dataset.
Warning: could not find SYS_ERR column

sherpa> data 2 459_heg_p1_bin10.pha
The inferred file type is PHA.  If this is not what you want, please
specify the type explicitly in the data command.
Warning: could not find SYS_ERR column
WARNING: statistical errors specified in the PHA file.
        These are currently IGNORED.  To use them, type:
        READ ERRORS "<filename>[cols CHANNEL,STAT_ERR]" fitsbin
WARNING: backgrounds UP and DOWN are being read from this file,
        and are being combined into a single background dataset.
Warning: could not find SYS_ERR column

sherpa> data 3 459_meg_m1_bin10.pha
The inferred file type is PHA.  If this is not what you want, please
specify the type explicitly in the data command.
Warning: could not find SYS_ERR column
WARNING: statistical errors specified in the PHA file.
        These are currently IGNORED.  To use them, type:
        READ ERRORS "<filename>[cols CHANNEL,STAT_ERR]" fitsbin
WARNING: backgrounds UP and DOWN are being read from this file,
        and are being combined into a single background dataset.
Warning: could not find SYS_ERR column

sherpa> data 4 459_meg_p1_bin10.pha
The inferred file type is PHA.  If this is not what you want, please
specify the type explicitly in the data command.
Warning: could not find SYS_ERR column
WARNING: statistical errors specified in the PHA file.
        These are currently IGNORED.  To use them, type:
        READ ERRORS "<filename>[cols CHANNEL,STAT_ERR]" fitsbin
WARNING: backgrounds UP and DOWN are being read from this file,
        and are being combined into a single background dataset.
Warning: could not find SYS_ERR column
```

*Sherpa* now refers to the spectra as follows:

- HEG, -1 order = dataset 1
  - HEG, +1 order = dataset 2
  - MEG, -1 order = dataset 3
  - MEG, +1 order = dataset 4
-

## Building the Instrument Responses

First, the instrument models are established by the `rsp` command. The `arf` parameter value is then set to the corresponding file for each order and arm:

```
sherpa> paramprompt off

sherpa> rsp[hegm1]
sherpa> rsp[hegp1]
sherpa> rsp[megm1]
sherpa> rsp[megp1]

sherpa> hegm1.arf = 459_heg_m1.arf
sherpa> hegp1.arf = 459_heg_p1.arf
sherpa> megm1.arf = 459_meg_m1.arf
sherpa> megp1.arf = 459_meg_p1.arf
```

This message will be printed after each gARF is entered:

```
The inferred file type is ARF.  If this is not what you want, please
specify the type explicitly in the data command.
```

In order to convolve the input datasets with the response model components that have been established, they must be defined as the instrument models. This involves pairing up the gARF and spectrum for each order, via the `instrument` command:

```
sherpa> instrument 1 = hegm1
sherpa> instrument 2 = hegp1
sherpa> instrument 3 = megm1
sherpa> instrument 4 = megp1
```


The current definition of the instrument model may be examined using `show instrument`:

```
sherpa> show instrument 1
Instrument 1: rsp1d[hegm1]
  Param  Type  Value
  -----
  1      rmf string: "none" (N_E=8192,N_PHA=8192)
  2      arf string: "459_heg_m1.arf" (N_E=8192)
```

Notice that *Sherpa* has defined properties for the `rmf` parameter, even though we did not enter a file. *Sherpa* has support for "dummy" instruments: if data have been input and the instrument stack contains only an ARF, a dummy RMF will be created that maps the ARF bins to the data bins, if possible. `ahelp instrument` contains more information on "dummy" instruments.

The datasets may now be plotted:

```
sherpa> lplot 4 data 1 data 2 data 3 data 4
```

Figure 1  shows the resulting plot.

## Filtering the Data

We choose to filter the data to focus on an area of interest:

```
sherpa> ignore allsets all
sherpa> notice allsets wave 1:15
```


The `ignore` command is used to ignore all the data in every dataset, then `notice` is used to select the

## Fitting Grating Data – Sherpa

desired regions. You may wish to adjust the limits to exclude more or less of your data.

Each filtered dataset may then be plotted:

```
sherpa> lplot 4 data 3 data 4 data 3 data 4
```

Notice that the plot now includes only the data in the specified wavelength regions. [Figure 2](#)  shows the resulting plot.

---

## Defining the Source and Background Models

We plan on simultaneously fitting the background data (rather than subtracting it), so we need to create a model expression for the source and the background. We model this source with a broken power law (`bp11d`) absorbed by the interstellar medium (`atten`). The background will be modeled by a one-dimensional power law (`powlaw1d`), also absorbed by the ISM (the same `atten` model).

First, we set up each model component. The absorption model will be referred to as "abs", the broken power law will be "bpow", and the 1D power law will be "pow1d".

```
sherpa> paramprompt on
Model parameter prompting is on

sherpa> atten[abs]
abs.hcol parameter value [1e+20]
abs.heiRatio parameter value [0.1]
abs.heiiRatio parameter value [0.01]

sherpa> bp11d[bpow]
bpow.gamma1 parameter value [0]
bpow.gamma2 parameter value [0]
bpow.eb parameter value [7.99625]
bpow.ref parameter value [7]
bpow.ampl parameter value [0.0238299]

sherpa> powlaw1d[pow1d]
pow1d.gamma parameter value [1]
pow1d.ref parameter value [7]
pow1d.ampl parameter value [0.0238299]
```

Note that since a dataset has already been input, *Sherpa* estimates the initial parameter values for the models based on the data. These values can also be listed with the `show` command:

```
sherpa> show models
-----
Defined source/background model components:
-----

Atten[abs] (integrate: off)
  Param  Type      Value      Min      Max      Units
  -----
  1 hcol  thawed    1e+20    1e+17    1e+24
  2 heiRatio thawed    1e-01      0        1
  3 heiiRatio thawed    1e-02      0        1

bp11d[bpow] (integrate: on)
  Param  Type      Value      Min      Max      Units
  -----
  1 gamma1 thawed      0        -10     10
  2 gamma2 thawed      0        -10     10
```

## Fitting Grating Data – Sherpa

```

3   eb thawed      7.9963      1.0075      14.985
4   ref frozen      7              1      14.985
5   ampl thawed   2.383e-02   2.383e-04      2.383

powlaw[powld] (integrate: on)
  Param  Type      Value      Min      Max      Units
  -----
1  gamma thawed      1      -10      10
2  ref frozen      7              1      14.985
3  ampl thawed   2.383e-02   2.383e-04      2.383

```

Next we modify the initial parameter value for `abs.hcol`:

```

sherpa> abs.hcol=1.81e20
sherpa> freeze abs

```

The hydrogen column density (`hcol`) is set to the Galactic value. All the `abs` parameters are then frozen, which means they will not be allowed to vary during the fit.

Now that the model components have been established, the product of `abs` and `bpow` is assigned as the source model for all datasets:

```

sherpa> source 1:4 = abs*bpow

```

while the background model is set as the product of `abs` and `powld`:

```

sherpa> background 1:4 = abs*powld

```

Both model definitions can be listed with the `show` command:

```

sherpa> show source 1
Source 1: (abs * bpow)
Atten[abs] (integrate: off)
  Param  Type      Value      Min      Max      Units
  -----
1  hcol frozen   1.81e+20   1e+17    1e+24
2 2heiRatio frozen  1e-01      0         1
3 3heiiRatio frozen  1e-02      0         1
bppld[bpow] (integrate: on)
  Param  Type      Value      Min      Max      Units
  -----
1  gamma1 thawed      0      -10      10
2  gamma2 thawed      0      -10      10
3  eb thawed      7.9963      1.0075      14.985
4  ref frozen      7              1      14.985
5  ampl thawed   2.383e-02   2.383e-04      2.383

sherpa> show background 1
Background 1: (abs * powld)
Atten[abs] (integrate: off)
  Param  Type      Value      Min      Max      Units
  -----
1  hcol frozen   1.81e+20   1e+17    1e+24
2 2heiRatio frozen  1e-01      0         1
3 3heiiRatio frozen  1e-02      0         1
powlaw[powld] (integrate: on)
  Param  Type      Value      Min      Max      Units
  -----
1  gamma thawed      1      -10      10
2  ref frozen      7              1      14.985
3  ampl thawed   2.383e-02   2.383e-04      2.383

```

## Examining Method & Statistic Settings

Next we check the current method and statistics settings:

```
sherpa> show method
Optimization Method: Levenberg-Marquardt

      Name      Value      Min      Max      Description
      ----      -
1  iters      2000      1      10000  Maximum number of iterations
2  eps       1e-03     1e-09     1      Absolute accuracy
3  smplx      0          0          1      Refine fit with simplex (0=no)
4  smplxep   1          1e-04     1000   Switch-to-simplex eps factor
5  smplxit   3          1          20     Switch-to-simplex iters factor

sherpa> show statistic
Statistic:      Chi-Squared Gehrels
```

For this fit, the default fitting and statistic settings will be used. More information is available from [ahelp lev-mar](#) and [ahelp chigehrels](#). For a list of all the available methods and statistic settings, see [ahelp method](#) and [ahelp statistic](#), respectively.

---

## Fitting

The datasets are now fit:

```
sherpa> fit
LVMQT: V2.0
LVMQT: initial statistic value = 1.06211e+10
LVMQT: final statistic value = 789026 at iteration 114
      bpow.gamma1  0.420507
      bpow.gamma2 -0.0786515
      bpow.eb     11.7909
      bpow.ampl   0.00225955
      powld.gamma 0.276206
      powld.ampl  0.000238299

WARNING:
  The value of powld.ampl within 0.01% of the powld.ampl.min limit boundary.
  You may wish to consider changing min/max values and refitting.
```

As the warning says, we reset the minimum boundary of `powld.ampl.min` and refit the data:

```
sherpa> powld.ampl.min=2.383e-10

sherpa> fit
LVMQT: V2.0
LVMQT: initial statistic value = 789026
LVMQT: final statistic value = 1954.98 at iteration 6
      bpow.gamma1  0.414261
      bpow.gamma2 -0.0607346
      bpow.eb     11.7837
      bpow.ampl   0.00237397
      powld.gamma 0.21045
      powld.ampl  9.79897e-06
```

To plot the fits:

```
sherpa> lplot 4 fit 3 fit 4 fit 3 fit 4
```




## Fitting Grating Data – Sherpa

```
sherpa> d 1,3,4 ylabel ""
sherpa> title "3C 273 (ObsID 459)"


sherpa> d 1 label 12 0.075 "HEG -1"
sherpa> d 2 label 12 0.075 "HEG +1"
sherpa> d 3 label 12 0.125 "MEG -1"
sherpa> d 4 label 12 0.125 "MEG +1"
sherpa> d all 1 all green

sherpa> redraw
```

The *ChIPS* commands are used to add labels to the drawing areas. The plot is shown in [Figure 3](#) .

It is also useful to plot the fit with the residuals:

```
sherpa> lplot 2 fit 1 delchi
```

This plot is shown in [Figure 4](#) . After creating a plot, it may be saved as a PostScript file:

```
sherpa> print postfile 459_1_fit_delchi.ps
```

---

## Examining Fit Results

There are several methods available in *Sherpa* for examining fit results. The goodness command reports information on the chi-square goodness-of-fit:

```
sherpa> goodness
Goodness: computed with Chi-Squared Gehrels

DataSet 1: 561 data points -- 555 degrees of freedom.
Statistic value      = 473.223
Probability [Q-value] = 0.994862
Reduced statistic    = 0.852654

Background for DataSet 1: 561 data points -- 559 degrees of freedom.
Statistic value      = 114.649
Probability [Q-value] = 1
Reduced statistic    = 0.205097

DataSet 2: 561 data points -- 555 degrees of freedom.
Statistic value      = 499.407
Probability [Q-value] = 0.956151
Reduced statistic    = 0.899832

Background for DataSet 2: 561 data points -- 559 degrees of freedom.
Statistic value      = 98.6341
Probability [Q-value] = 1
Reduced statistic    = 0.176447

DataSet 3: 281 data points -- 275 degrees of freedom.
Statistic value      = 279.114
Probability [Q-value] = 0.419589
Reduced statistic    = 1.01496

Background for DataSet 3: 281 data points -- 279 degrees of freedom.
Statistic value      = 119.737
Probability [Q-value] = 1
Reduced statistic    = 0.429164

DataSet 4: 281 data points -- 275 degrees of freedom.
Statistic value      = 260.716
Probability [Q-value] = 0.722871
```

## Fitting Grating Data – Sherpa

```
Reduced statistic      = 0.948058

Background for DataSet 4: 281 data points -- 279 degrees of freedom.
Statistic value       = 109.496
Probability [Q-value] = 1
Reduced statistic     = 0.392459

Total   : 3368 data points -- 3362 degrees of freedom.
Statistic =          1954.98
Probability =          1
Reduced statistic = 0.581492
```

The uncertainty, covariance, and projection commands can be used to estimate confidence intervals for the thawed parameters:

```
sherpa> covariance

Computed for sherpa.cov.sigma = 1
-----
Parameter Name      Best-Fit Lower Bound      Upper Bound
-----
bpow.gamma1         0.414261 -0.00746802      +0.00746802
bpow.gamma2         -0.0607528 -0.119213        +0.119213
bpow.eb              11.7837 -0.243908        +0.243908
bpow.ampl            0.00237397 -8.36986e-06     +8.36986e-06
powld.gamma          0.21045 -0.0654054       +0.0654054
powld.ampl           9.79894e-06 -2.59661e-07     +2.59661e-07
```

---

## Saving and Quitting the Session

Before exiting *Sherpa*, you may wish to save the session in order to return to the analysis at a later point:

```
sherpa> save all 459_fitting_session.shp
```

All the information about the current session is written to `459_fitting_session.shp`, an ASCII file. It may be loaded into *Sherpa* again with the use command.

Finally, quit the session:

```
sherpa> quit
```

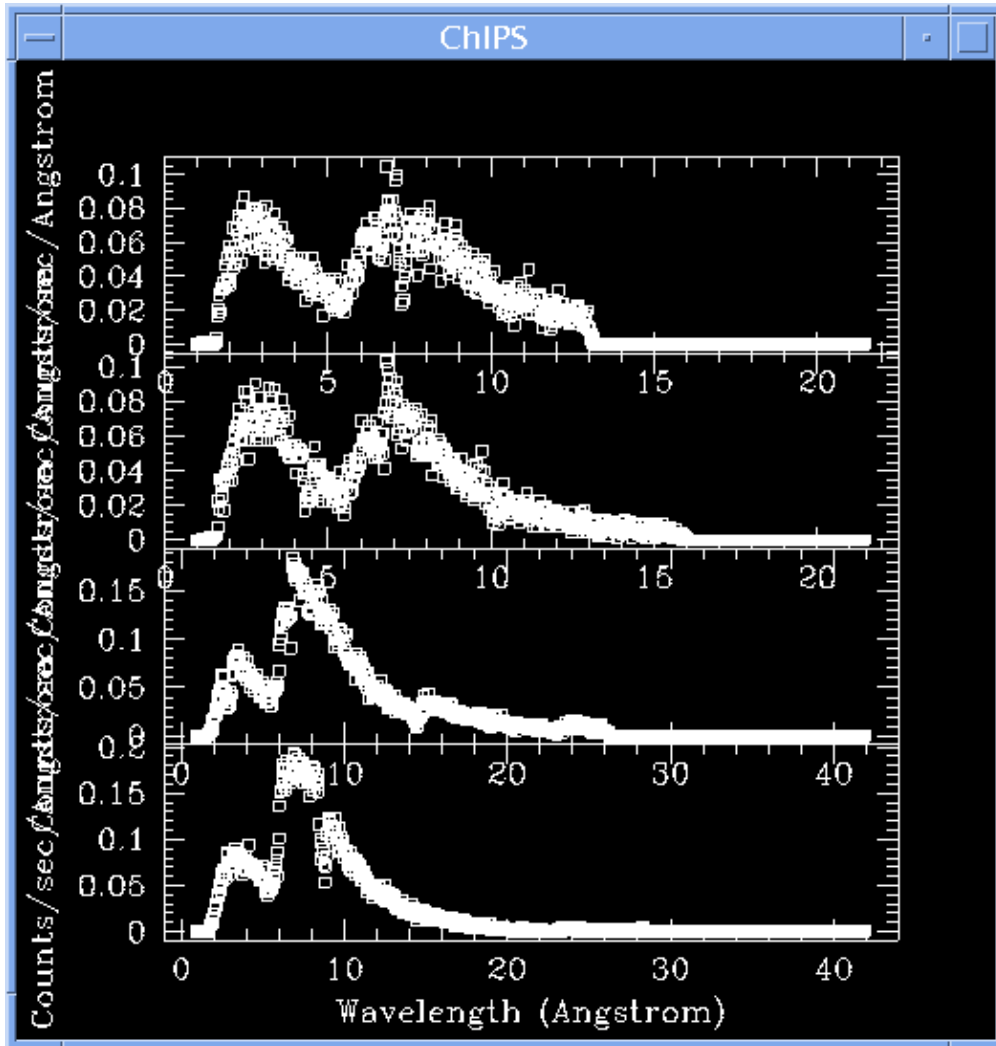
---

## History

- 14 Jan 2005 updated for CIAO 3.2: minor changes to screen output
- 11 Jul 2005 overall revision to thread, changes to screen output
- 21 Dec 2005 reviewed for CIAO 3.3: no changes
- 01 Dec 2006 reviewed for CIAO 3.4: no changes

### Image 1: Plotting the four orders

The plots show, from top to bottom, the HEG -1, HEG +1, MEG -1, and MEG +1 orders.



## Image 2: Filtering the datasets

All four plots have been restricted to the 1–15 Å range.

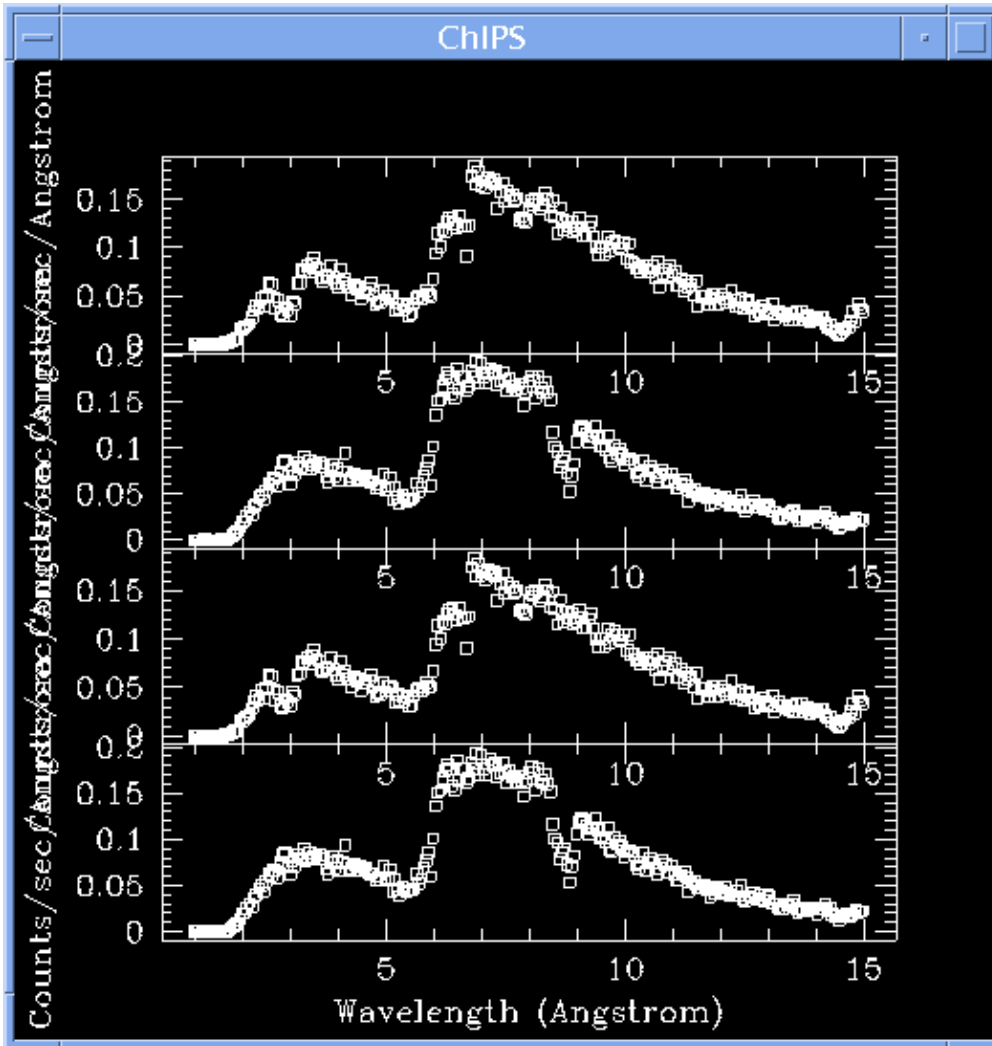


Image 3: Results of simultaneous fit

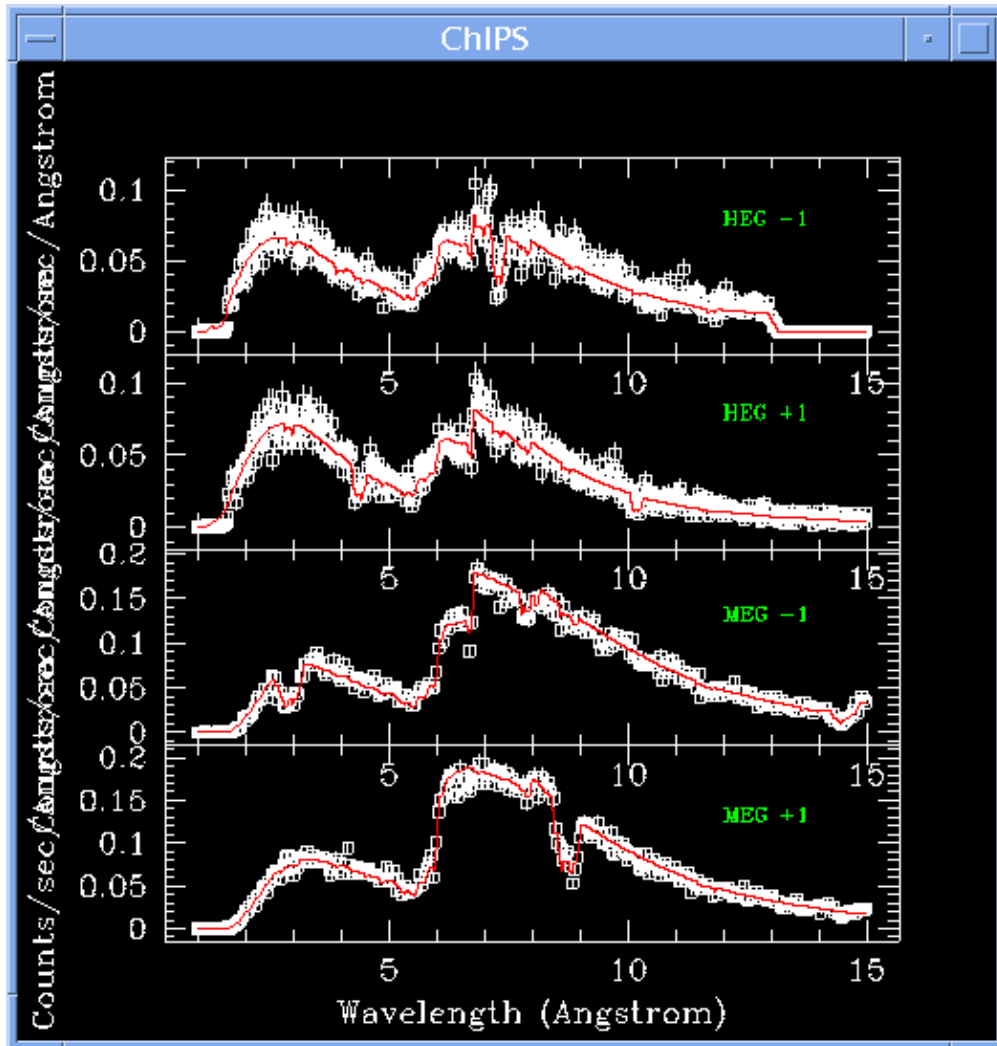


Image 4: Fit and residuals for the HEG -1 order spectrum

