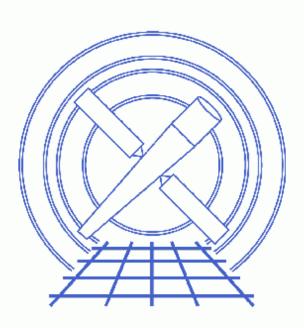
Fitting Grating Data



Sherpa Threads (CIAO 3.4)

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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 <u>Fitting Multiple Orders of HRC-S/LETG Data</u> thread helpful for their analysis.

Proceed to the <u>HTML</u> or hardcopy (PDF: <u>A4 / letter</u>) 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 <u>Create Grating RMFs for ACIS Observations thread</u>. Creating observation–specific gRMFs is optional, and is discussed further in the <u>Building the Instrument Responses</u> section.

The data files are available in sherpa.tar.gz, as explained in the Sherpa Getting Started thread.

Reading the Spectrum Files

The data are input to *Sherpa* with the <u>data</u> command (a shorthand version of "<u>read_data</u>"):

```
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_heq_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 <u>rsp</u> 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[megm1]
sherpa> rsp[megm1]
sherpa> rsp[megp1]
sherpa> hegm1.arf = 459_heg_m1.arf
sherpa> hegm1.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 <u>instrument</u> command:

sherpa>	instrument	1	=	hegm1
sherpa>	instrument	2	=	hegp1
sherpa>	instrument	3	=	megml
sherpa>	instrument	4	=	megp1

The current definition of the instrument model may be examined using <u>show</u> instrument:

```
sherpa> show instrument 1
Instrument 1: rspld[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. <u>ahelp instrument</u> contains more information on "dummy" instruments.

The datasets may now be plotted:

```
sherpa> <u>lplot</u> 4 data 1 data 2 data 3 data 4
Figure 1 or shows the resulting plot.
```

Filtering the Data

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

sherpa>	<u>ignore</u>	allsets	all	
sherpa>	<u>notice</u>	allsets	wave	1:15

The <u>ignore</u> command is used to ignore all the data in every dataset, then <u>notice</u> is used to select the 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> <u>lplot</u> 4 data 3 data 4 data 3 data 4	sh	erpa>	<u>lplot</u>	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 to 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 (<u>bplld</u>) absorbed by the interstellar medium (<u>atten</u>). The background will be modeled by a one-dimensional power law (<u>powlawld</u>), 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> bpl1d[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]
powld.gamma parameter value [1]
powld.ref parameter value [7]
powld.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 <u>show</u> command:

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sherpa> show models										
Defined sour	rce/background m	odel compone	nts:							
	(integrate: off									
	Type Value			Units						
	 nawed 1e+20									
	thawed 1e-0									
	thawed 1e-									
	(integrate: on									
Param	Type Value	Min	Max	Units						
	nawed 0	-10	10							
	nawed 0									
3 eb th	nawed 7.9963	1.0075	14.985							
	rozen 7									
5 ampl th	nawed 2.383e-02	2.383e-04	2.383							
powlaw[powld] (integrate: on)										
	Type Value		Max	Units						
1 gamma th	nawed 1	-10	10							
	rozen 7									
3 ampl th	hawed 2.383e-02	2.383e-04	2.383							

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

```
sherpa> abs.hcol=1.81e20
sherpa> <u>freeze</u> abs
```

The hydrogen column density (hcol) is set to the Galactic value. All the abs parameters are then <u>frozen</u>, 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:

chernas	gource	1:4	_	abs*bpow
sherpa/	Source	T • 4	-	absopow

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

sherpa> <u>background</u> 1:4 = abs*powld

Both model definitions can be listed with the show command:

sherpa> <u>show</u> source Source 1: (abs * bp				
Atten[abs] (integr	ate: off)			
Param Type	Value	Min	Max	Units
1 hcol frozen	1.81e+20	1e+17	1e+24	
2heiRatio frozen	1e-01	0	1	
3heiiRatio frozen	1e-02	0	1	
<pre>bpl1d[bpow] (integ</pre>	grate: on)			
Param Type	Value	Min	Max	Units
1 gammal thawed	0	-10	10	
2 gamma2 thawed	0	-10	10	
3 eb thawed	7.9963	1.0075	14.985	

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4 ref frozen	7	1	14.985	
5 ampl thawed	2.383e-02 2	2.383e-04	2.383	
sherpa> <u>show</u> back	kground 1			
Background 1: (at	os * powld)			
Atten[abs] (inte	egrate: off)			
Param Type	Value	Min	Max	Units
1 hcol frozen	1.81e+20	1e+17	1e+24	
2heiRatio frozer	n 1e-01	0	1	
3heiiRatio froze	en 1e-02	0	1	
powlaw[powld] (i	integrate: on)	1		
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	2.383e-04	2.383	

Examining Method & Statistic Settings

Next we check the current method and statistics settings:

	sherpa> show <u>method</u> 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					
she	sherpa> show <u>statistic</u>									
Sta	atistic:	Chi-S	quared Gehre	els						

For this fit, the default fitting and statistic settings will be used. More information is available from <u>ahelp</u> <u>lev-mar</u> and <u>ahelp</u> chigehrels. For a list of all the available methods and statistic settings, see <u>ahelp</u> <u>method</u> and <u>ahelp</u> statistic, respectively.

Fitting

The datasets are now fit:

```
sherpa> <u>fit</u>
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 powbkgmh.ampl 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
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 l all green
sherpa> redraw
```

The ChIPS commands are used to add labels to the drawing areas. The plot is shown in Figure 3 0.

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

sherpa> <u>lplot</u> 2 fit 1 delchi

This plot is shown in Figure 4 10. After creating a plot, it may be saved as a PostScript file:

```
sherpa> <u>print</u> postfile 459_1_fit_delchi.ps
```

Examining Fit Results

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

```
sherpa> <u>qoodness</u>
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.
                 = 260.716
Statistic value
Probability [Q-value] = 0.722871
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 <u>uncertainty</u>, <u>covariance</u>, and <u>projection</u> commands can be used to estimate confidence intervals for the thawed parameters:

sherpa> covariance									
Computed for sherpa.cov.sigma = 1									
Parameter Na	ame Best-Fit	Lower Bound	Upper Bound						
bpow.gamma bpow.gamma bpow.eb bpow.ampl powld.gamma powld.ampl	a2 -0.0607528 11.7837 0.00237397		+0.0654054						

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> <u>save</u> 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 <u>use</u> command.

Finally, quit the session:

sherpa> <u>quit</u>

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

URL: http://cxc.harvard.edu/sherpa/threads/grating/

Last modified: 1 Dec 2006

Image 1: Plotting the four orders

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

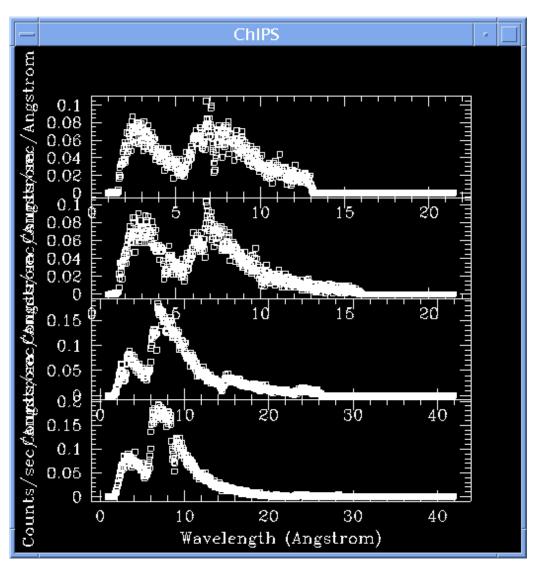
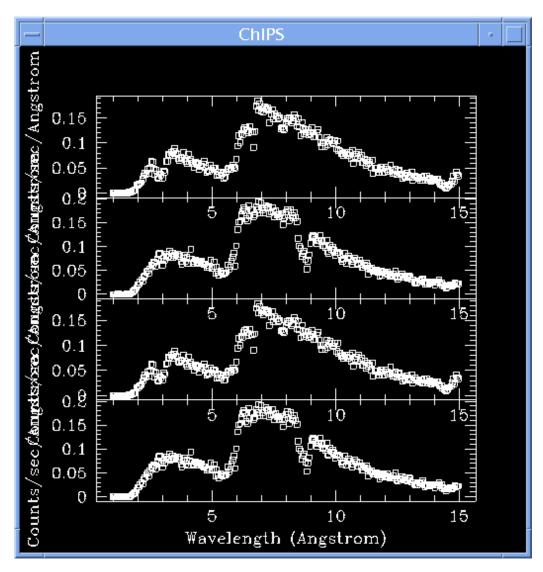


Image 2: Filtering the datasets

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



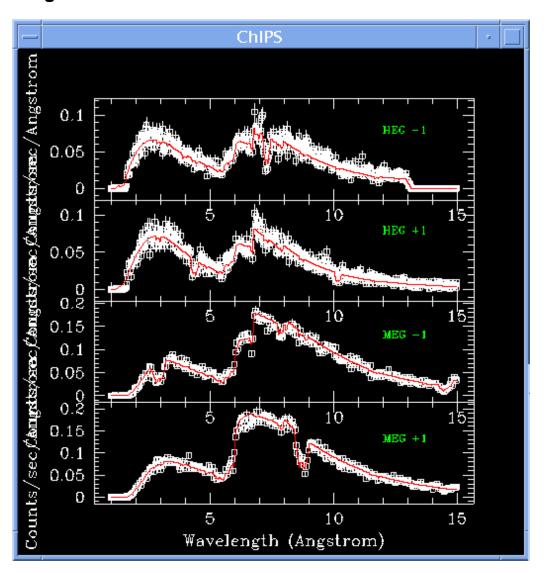


Image 3: Results of simultaneous fit

