

IRAF 一维光谱处理入门

(内容来自多方, 在此不详细注明出处)

IRAF 入门

之一维光谱

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第一章：IRAF 基本操作

1.1 Fits 文件

老版本的 IRAF 需要进行文件格式转换，新版本的 IRAF 可以直接对 fits 或 fit 进行操作，无需转换成 imh 格式。

1.2 批处理

使用 list 进行批处理，可以节省大量时间。

```
ecl> ls *.fits > l1
```

将本地所有的 fits 文件名添加到 l1 文件中，对列表进行操作时只需在列表名前增加@符号，如删除 l1 中的文件

```
ecl> del @l1 **现在版本的IRAF不再区别imdel和del
```

1.3 图像拷贝

使用 imcopy 命令可以方便地进行图像拷贝以及裁剪等操作。

```
ecl> imcopy raw.fits new.fits **将raw.fits复制到new.fits
```

```
ecl> imcopy raw.fits[600:1200,700:3600] new2.fits **将raw.fits x轴(600:1200)y轴(700:3600)区域内的数据复制到new2.fits,对YFOOSC长缝光谱而言,这一区域经常被用到
```

在处理 YFOOSC 数据时，由于 fits 文件有两层，而只有第一层有数据，因此在进行操作时需要指明是对第一层进行。比如显示一幅 YFOOSC 数据：

```
ecl> display YFub090021.fits[1] 1 **[1]即指明是第一层
```

而在一些处理中，IRAF 不能很好的识别[1]符号，因此可以用 imcopy 将原始的具有两层结果的 fits 文件，转换为单层 fits 文件。

```
ecl> imcopy YFub090021.fits[1] object.fits
```

利用批处理，一次转换全部数据

```
ecl> ls *.fits > l1
```

在本地建立 l2 文件，将 l1 中的内容拷贝进 l2，利用查找替换，将 l1 中的 fits 替换成 fits[1]。如果要保留原始数据，用查找替换修改 l2 中的文件名。

```
ecl> imcopy @l1 @l2
```

如需进行裁剪，在 l1 中添加裁剪区域即可，fits[1][600:1200,700:3600].

下面也将介绍利用 `ccdproc` 命令的 `trim` 进行图像裁剪，本文没有考虑两者的差别。

1.4 程序包

IRAF 中有大量的程序包可供使用，为了使用程序包中的函数，需要先登录到该程序所在的程序包中，比如要使用 `ccdproc` 命令，需要知道这个函数属于哪个程序包。

```
ecl> help ccdproc
```

出来帮助信息，第一行就给出了该函数所在的程序包。

```
CCDPROC (Dec93)          noao.imred.ccdred          CCDPROC (Dec93)
```

```
ecl> noao  **输入最上层程序包名—noao，进入后出现一系列第二层程序包
```

```
artdata.      digiphot.      nobsolete.      onedspec.
astcat.       focas.          nproto.        rv.
astrometry.   imred.          observatory    surfphot.
astutil.      mtlocal.        obsutil.       twodspec.
```

```
noao> imred **输入第二层程序包名—imred，出现第三层程序包
```

```
argus.        crutil.         echelle.       iids.          kpnocoude.    specred.
bias.         ctioslit.       generic.       irred.         kpnoslit.     vtel.
ccdred.       dtoi.           hydra.         irs.           quadred.
```

```
im> ccdred **进入到imred程序包，出现下列函数
```

```
badpixmap     ccdmask         flatcombine    mkskyflat
ccdgroups     ccdproc        mkfringecor    setinstrument
cdhedit       ccdtest        mkillumcor     zerocombine
ccdinstrument combine         mkillumflat
cdclist       darkcombine    mskskycor
```

```
ccd> ccdproc **现在可以调用ccdproc函数了
```

对于一些常用函数、子程序包或者自行编写的脚步，可以通过修改 `login.cl` 文件，在登陆 IRAF 时即刻加载。详细内容见本系列关于脚本部分的介绍文档，此次不再赘述。

第二章：图像处理

2.1 数据特点

在进行 CCD 光谱图像的预处理时，基本原理与测光内容相同，只是由于光谱观测的曝光时间较长，因此有更多地宇宙线进入 CCD，影响观测。比起测光观测，需要更加细致的剔除宇宙线。另外，光谱观测时经常使用灯光平场，这种平场在色散方向以及空间方向上都存在一定的不均匀性，需要进行归一化。

所谓的一维光谱是针对 Echell 光谱而言，即只有色散方向是一维，而不像 Echell 光谱，多级光谱分布在不同的空间位置上。因此，使用长缝所拍摄的面源光谱，也一并称作一维光谱。

2.2 数据检查

数据检查主要包括两部分内容。

首先是数据的完备性检查。待处理的数据包含有本底、平场、待测目标等几类数据，有些时候还需要标准星的观察数据。对于光谱观测而言，还需要定标灯谱，灯光平场等数据。

其次是数据质量检查。对拍摄的目标用 `display` 依次显示所有获取图像，(**在脚本部分的介绍中，将提供一个自动显示图像的脚本)，目视检测图像质量，尤其是辅助图像，如本底、平场等。去掉那些有明显问题的图像，比如平场中有星。接下来，用 `imstat` 命令统计辅助图像的平均值和标准差。统计范围一般选取图像中心的避开坏列的一块小区域。

例如：

```
ecl>imstat bias01.fits[600:1200,700:3600] **统计[600:1200,700:3600]范围内的统计结果
```

去掉那些平均值和标准差值差别很大的图像。

2.3 文件改名

在处理数据前，根据文件内容修改文件名是一个比较好的习惯。可以通过 `log` 文件来进行分类，也可以使用 `imhead` 命令，来查看文件类型。将文件按照 `bias`、`flat`、`object`、`stdstars`、`lamb` 等类型，分别设置新的名字。可以采用批处理的方法，在此不再赘述。

2.4 裁剪图像

有时候，由于光学设计以及 CCD 本身的原因，CCD 的边缘部分不可用。这时候需要对除 bias 以外的数据进行裁剪。裁剪的方法可以参考 1.3，也可以使用 ccdproc 命令。

注意，ccdproc 命令不允许 fits 头文件中有 CCDSEC、DATASEC 两个参数，如果待处理的文件中还有上述参数，需要先用 hedit 命令删除

```
PACKAGE = imutil
TASK = hedit
images =          *.fits  images to be edited
fields =    CCDSEC DATASEC  fields to be edited ** 空格号隔开
value =          value expression
(add =          no) add rather than edit fields
(addonly=        no) add only if field does not exist
(delete =        yes) delete rather than edit fields
(verify =        no) verify each edit operation
(show =          yes) print record of each edit operation
(update =        yes) enable updating of the image header
(mode =          ql)
```

:g

登陆到 noao/imred/ccdred 中：

ccd> epa ccdp ****epa 命令是editparameter的缩写，用来编辑命令中的各项参数**

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PACKAGE = ccdred

TASK = ccdproc

```
images =    imagename  List of CCD images to correct**亦可使用list进行批处理，
@II
(output =          ) List of output CCD images**若设置为空，则覆盖原
文件
(ccdtype=          ) CCD image type to correct
(max_cac=          0) Maximum image caching memory (in Mbytes)
(noproc =        no) List processing steps only?
(fixpix =        no) Fix bad CCD lines and columns?
```

(oversca= no) Apply overscan strip correction?
(trim = *yes*) Trim the image? ***裁剪图像*
(zerocor= no) Apply zero level correction?
(darkcor= no) Apply dark count correction?
(flatcor= no) Apply flat field correction?
(illumco= no) Apply illumination correction?
(fringec= no) Apply fringe correction?
(readcor= no) Convert zero level image to readout correction?
(scancor= no) Convert flat field image to scan correction?
(readaxi= line) Read out axis (column|line)
(fixfile=) File describing the bad lines and columns
(biassec=) Overscan strip image section
(trimsec= [600:1200,700:3600]) Trim data section ***裁剪尺寸*
(zero =) Zero level calibration image
(dark =) Dark count calibration image
(flat =) Flat field images
(illum =) Illumination correction images
(fringe =) Fringe correction images
(minrepl= 1.) Minimum flat field value
(scantyp= shortscan) Scan type (shortscan|longscan)
(nscan = 1) Number of short scan lines
(interac= no) Fit overscan interactively?
(funcio= chebyshev) Fitting function
(order = 1) Number of polynomial terms or spline pieces
(sample = *) Sample points to fit
(naverag= 1) Number of sample points to combine
(niterat= 1) Number of rejection iterations
(low_rej= 3.) Low sigma rejection factor
(high_re= 3.) High sigma rejection factor
(grow = 0.) Rejection growing radius
(mode = ql)

:g ****:g 运行, :q 保存并退出,:e 编辑子函数的参数

2.5 BIAS

2.5.1 合并 BIAS

利用 noao\imred\ccdred 中 *zerocombine*

cc> epa zeroc ****IRAF 中的所有命令均可使用简写, 只要该简写在程序包中以及默认加载的包中是唯一的**

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PACKAGE = ccdred

TASK = zerocombine

input = bias*.fits List of zero level images to combine ****本人习惯将所有 bias 文件改成 biasxx.fits**

(output = zero) Output zero level name

(combine= average) Type of combine operation

(reject = minmax) Type of rejection

(ccdtype=) CCD image type to combine

(process= no) Process images before combining?

(delete = no) Delete input images after combining?

(clobber= no) Clobber existing output image?

(scale = none) Image scaling

(statsec=) Image section for computing statistics

(nlow = 0) minmax: Number of low pixels to reject

(nhigh = 1) minmax: Number of high pixels to reject

(nkeep = 1) Minimum to keep (pos) or maximum to reject (neg)

(mclip = yes) Use median in sigma clipping algorithms?

(lsigma = 3.) Lower sigma clipping factor

(hsigma = 3.) Upper sigma clipping factor

(rdnoise= **6.5**) ccdclip: CCD readout noise (electrons) ****YFOSC 一号 CCD**

(gain = **0.4**) ccdclip: CCD gain (electrons/DN) ****YFOSC 一号 CCD**

(snoise = 0.) ccdclip: Sensitivity noise (fraction)

(pclip = -0.5) pclip: Percentile clipping parameter

(blank = 0.) Value if there are no pixels

(mode = ql)

:g

生成文件 zero.fits

2.5.2 减去 bias

利用 `ccproc` 命令，减去平场、目标等图像中的 bias

```
cc> epa ccdp
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = ccdred
```

```
    TASK = ccdproc
```

```
images =          imagename  List of CCD images to correct
(output =          ) List of output CCD images
(ccdtype=         ) CCD image type to correct
(max_cac=         0) Maximum image caching memory (in Mbytes)
(noproc =         no) List processing steps only?
(fixpix =         no) Fix bad CCD lines and columns?
(oversca=        no) Apply overscan strip correction?
(trim   =         no) Trim the image?
(zerocor=        yes) Apply zero level correction? ** 减本底
(darkcor=        no) Apply dark count correction?
(flatcor=        no) Apply flat field correction?
(illumco=        no) Apply illumination correction?
(fringec=        no) Apply fringe correction?
(readcor=        no) Convert zero level image to readout correction?
(scancor=        no) Convert flat field image to scan correction?
(readaxi=        line) Read out axis (column|line)
(fixfile=        ) File describing the bad lines and columns
(biassec=        ) Overscan strip image section
(trimsec=        ) Trim data section
(zero   =        Zero.fits) Zero level calibration image
(dark   =        ) Dark count calibration image
(flat   =        ) Flat field images
(illum  =        ) Illumination correction images
(fringe =        ) Fringe correction images
```

(minrepl=	1.) Minimum flat field value
(scantyp=	shortscan) Scan type (shortscan longscan)
(nscan =	1) Number of short scan lines
(interac=	no) Fit overscan interactively?
(functio=	chebyshev) Fitting function
(order =	1) Number of polynomial terms or spline pieces
(sample =	*) Sample points to fit
(naverag=	1) Number of sample points to combine
(niterat=	1) Number of rejection iterations
(low_rej=	3.) Low sigma rejection factor
(high_re=	3.) High sigma rejection factor
(grow =	0.) Rejection growing radius
(mode =	ql)

:g

2.6 平场

2.6.1 原始平场合并

cc>epa flatcombine

I R A F

Image Reduction and Analysis Facility

PACKAGE = ccdred

TASK = flatcombine

input =	flat*.fits	List of flat field images to combine
(output =	Flat.fits)	Output flat field root name
(combine=	average)	Type of combine operation
(reject =	avsigclip)	Type of rejection
(ccdtype=)	CCD image type to combine
(process=	no)	Process images before combining?
(subsets=	yes)	Combine images by subset parameter?
(delete =	no)	Delete input images after combining?
(clobber=	no)	Clobber existing output image?
(scale =	mode)	Image scaling

```

(statsec=          ) Image section for computing statistics
(nlow  =          ) 1) minmax: Number of low pixels to reject
(nhigh =          ) 1) minmax: Number of high pixels to reject
(nkeep =          ) 1) Minimum to keep (pos) or maximum to reject (neg)
(mclip =          ) yes) Use median in sigma clipping algorithms?
(lsigma =         ) 3.) Lower sigma clipping factor
(hsigma =         ) 3.) Upper sigma clipping factor
(rdnoise=         ) 6.5) ccdclip: CCD readout noise (electrons)
(gain  =          ) 0.4) ccdclip: CCD gain (electrons/DN)
(snoise =         ) 0.) ccdclip: Sensitivity noise (fraction)
(pclip =          ) -0.5) pclip: Percentile clipping parameter
(blank =          ) 1.) Value if there are no pixels
(mode  =          ) ql)

```

:g

合并后的平场 Flat.fits

2.6.2 色散方向上的平场归一化

注意：由于平场灯是黑体辐射，经过色散系统在色散方向上能量呈黑体谱分布，用 `response` 命令是来去除这种照明上的不均匀性。

```
noao\twospec\longslit\response
```

```
lo> epa response
```

```
I R A F
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = longslit
```

```
TASK = response
```

```
calibrat=          Flat  Longslit calibration images ** 上一步合并的平场
```

```
normaliz=         Flat  Normalization spectrum images
```

```
response=         reFlat Response function images ** 归一化后输出的平场
```

```
(interac=         ) yes) Fit normalization spectrum interactively?
```

```
(thresho=         ) INDEF) Response threshold
```

```
(sample =         ) *) Sample of points to use in fit
```

```
(naverag=         ) 1) Number of points in sample averaging
```

(funcio= spline3) Fitting function ** 有多种拟合函数可供选择: *legendre, chebyshev, spline1* 以及 *spline3*, 常用三次样条函数 *spline3*

(order = 7) Order of fitting function ** 拟合的阶数, 在交互模式下根据拟合情况而定, 一般是高阶拟合

(low_rej= 0.) Low rejection in sigma of fit

(high_re= 0.) High rejection in sigma of fit

(niterat= 1) Number of rejection iterations

(grow = 0.) Rejection growing radius

(graphic= stdgraph) Graphics output device

(cursor =) Graphics cursor input

(mode = ql)

:g

在弹出的交互式窗口 *irafterm* 中, 键入:f 可再次选择拟合函数的种类, 如

:f spline3

键入:o 可选择拟合阶数, 如

:o 10

按 q 退出, 此时会发现文件夹中生成了归一化平场 *reFlat.fits*

2.6.3 空间方向上的平场归一化

由于采用灯光照明时, 出射光在空间上具有不均匀性 (如 YFOSC 的光纤, 其出射光能的分布大致服从高斯分布), 为了消除这一影响, 需要使用 *illumination* 命令。

```
noao\twospec\longslit\illumination
```

```
TASK = illumination
```

```
images = reFlat.fits Longslit calibration images
```

```
illumina= ilFlat.fits Illumination function images
```

```
(interac= yes) Interactive illumination fitting?
```

```
(bins = ) Dispersion bins
```

```
(nbins = 5) Number of dispersion bins when bins = ""
```

```
(sample = *) Sample of points to use in fit
```

```
(naverag= 1) Number of points in sample averaging
```

```
(funcio= spline3) Fitting function
```

```
(order = 1) Order of fitting function ** 使用低阶拟合
```

```
(low_rej= 0.) Low rejection in sigma of fit
```

```
(high_re=          0.) High rejection in sigma of fit
(niterat=         1) Number of rejection iterations
(grow   =         0.) Rejection growing radius
(interpo=        poly3) Interpolation type
(graphic=        stdgraph) Graphics output device
(cursor =         ) Graphics cursor input
(mode   =         ql)
```

2.6.4 两个方向的平场归一化

利用 `imarith` 命令, 用 `response` 得到的平场除以 `illumination` 命令得到的平场, 生成完美的平场。

```
ec1> imarith reFlat / ilFalt perFlat **生成完美的平场perFlat.fits
```

2.6.5 对图像除归一化平场

```
lo> epa ccdp
```

```
I R A F
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = ccdred
```

```
TASK = ccdproc
```

```
images =          Object*.fits  List of CCD images to correct
(output =         ) List of output CCD images
(ccdtype=         ) CCD image type to correct
(max_cac=         0) Maximum image caching memory (in Mbytes)
(noproc =         no) List processing steps only?
(fixpix =         no) Fix bad CCD lines and columns?
(oversca=        no) Apply overscan strip correction?
(trim   =         no) Trim the image?
(zeroacor=       no) Apply zero level correction?
(darkcor=       no) Apply dark count correction?
(flatcor=       yes) Apply flat field correction?
(illumco=       no) Apply illumination correction?
(fringec=       no) Apply fringe correction?
(readcor=       no) Convert zero level image to readout correction?
(scancor=       no) Convert flat field image to scan correction?
```

(readaxi= line) Read out axis (column|line)
 (fixfile=) File describing the bad lines and columns
 (biassec=) Overscan strip image section
 (trimsec=) Trim data section
 (zero = zero) Zero level calibration image
 (dark =) Dark count calibration image
 (flat = *perFlat*) Flat field images
 (illum =) Illumination correction images
 (fringe =) Fringe correction images
 (minrepl= 1.) Minimum flat field value
 (scantyp= shortscan) Scan type (shortscan|longscan)
 (nscan = 1) Number of short scan lines
 (interac= no) Fit overscan interactively?
 (functio= chebyshev) Fitting function
 (order = 1) Number of polynomial terms or spline pieces
 (sample = *) Sample points to fit
 (naverag= 1) Number of sample points to combine
 (niterat= 1) Number of rejection iterations
 (low_rej= 3.) Low sigma rejection factor
 (high_re= 3.) High sigma rejection factor
 (grow = 0.) Rejection growing radius
 (mode = ql)

 :g

2.7 剔除宇宙线

有多种方法剔除宇宙线，在宇宙线不多的情况下，可以在抽取完光谱之后，利用 `splot` 命令手动剔除。下面介绍叠加多幅图像剔除宇宙线的办法。

2.7.1 谱对齐

拍摄一个暗弱目标的光谱，往往会曝光较长的时间。在曝光期间，宇宙射线会不可避免的在 CCD 上产生噪声。比方两个小时的曝光时间，如果只拍摄一张两个小时的图像，那么宇宙线在 CCD 上会有两个小时的累加；但如果分别拍摄两张一个小时的图片，由于宇宙线在 CCD 上得分布是随机的，因此通过把两张图像合并就可以去掉大部分的宇宙线。而由于没有比较，一张两小时曝光的照片

则无法通过这种方法去除宇宙线。因此，合并两张图像是必要的。而合并的第一步就是确保两张图像已经严格对齐；

打开两张图像：

```
lo> displ w0126012.fits 1
```

```
lo> displ w0126014.fits 2
```

在 ds9 中可以通过 Frame 中得 prev 和 next 对比查看两张图像；

```
lo> imexam
```

打开两张图像：

```
ec> displ w0424013.fits 1
```

```
ec> displ w0424014.fits 2
```

在 ds9 中可以通过 Frame 中得 prev 和 next 对比查看两张图像；

```
ec> imexam
```

将鼠标分别停在两张图像相同的坐标处，分别按 z，在 xgterm 窗口中比较两张图像上像素点随坐标的变化趋势，如图 15-1；结合对图像的观察，确定图像是否对齐；若没有，需要移动几个像素等；

比较后发现，对这组数据，不需要移动图像；若需要移动，则可使用命令 imshift；

xshift, yshift 分别为移动的大小；

输出 x 坐标 = 输入 x 坐标 + xshift

输出 y 坐标 = 输入 y 坐标 + yshift

```
cc> immatch
```

```
im> epa imshift
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = immatch
```

```
TASK = imshift
```

```
input   =   w0126014.fits  Input images to be fit
```

```
output  =   w0126014.fits  Output images
```

```
xshift  =                               Fractional pixel shift in x
```

```
yshift  =                               Fractional pixel shift in y
```

```
(shifts_ =                               ) Text file containing shifts for each image
```

```
(interp_ = linear) Interpolant (nearest,linear,poly3,poly5,spline3,sinc,drizzle)
```

```
(boundar_ = nearest) Boundary (constant,nearest,reflect,wrap)
```


(constan= 0.) Constant for boundary extension
 (mode = ql)

 :g

2.7.2 图像合并，产生一幅假图像：

im> epa imcombine

I R A F

Image Reduction and Analysis Facility

PACKAGE = immatch

TASK = imcombine

input = w0126012,w0126014 List of images to combine**输入两幅要合并的图像

output = w3com List of output images

(headers=) List of header files (optional)

(bpmask=) List of bad pixel masks (optional)

(rejmask=) List of rejection masks (optional)

(nrejmas=) List of number rejected masks (optional)

(expmask=) List of exposure masks (optional)

(sigmas =) List of sigma images (optional)

(logfile= STDOUT) Log file

(combine= average) Type of combine operation

(reject = minmax) Type of rejection

(project= no) Project highest dimension of input images?

(outtype= real) Output image pixel datatype

(outlimi= none) Output limits (x1 x2 y1 y2 ...)

(offsets= none) Input image offsets

(masktyp=) Mask type

(maskval= 0.) Mask value

(blank = 0.) Value if there are no pixels

(scale = exposure) Image scaling

(zero = none) Image zero point offset

(weight = none) Image weights

(statsec=) Image section for computing statistics

(expname= exposure) Image header exposure time keyword

(lthresh= INDEF) Lower threshold

```

(hthresh=          INDEF) Upper threshold
(nlow   =          0) minmax: Number of low pixels to reject
(nhigh  =          1) minmax: Number of high pixels to reject
(nkeep  =          1) Minimum to keep (pos) or maximum to reject (neg)
(mclip  =          yes) Use median in sigma clipping algorithms?
(lsigma =          3.) Lower sigma clipping factor
(hsigma =         10.) Upper sigma clipping factor
(rdnoise=        2.64) ccdclip: CCD readout noise (electrons)
(gain   =        2.25) ccdclip: CCD gain (electrons/DN)
(snoise =          0.) ccdclip: Sensitivity noise (fraction)
(sigscal=        0.1) Tolerance for sigma clipping scaling corrections
(pclip  =       -0.5) pclip: Percentile clipping parameter
(grow   =          0.) Radius (pixels) for neighbor rejection
(mode   =          ql)

```

:g

2.7.3 修改第三幅图像的文件头:

键入:

```
im> hedit w3com.fits ncombine 0
```

```
im> hedit w3com.fits exposure 0
```

2.7.4 合并三幅图像以去除宇宙线:

```
im> epa imco
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = immatch
```

```
    TASK = imcombine
```

```
input   = w0126012,w0126014,w3com  List of images to combine
```

```
output  =          w3caver  List of output images
```

```
(headers=          ) List of header files (optional)
```

```
(bpmasks=          ) List of bad pixel masks (optional)
```

```
(rejmask=          ) List of rejection masks (optional)
```

```
(nrejmas=          ) List of number rejected masks (optional)
```

(expmask=) List of exposure masks (optional)
 (sigmas =) List of sigma images (optional)
 (logfile= STDOUT) Log file
 (combine= average) Type of combine operation
 (reject = avsigclip) Type of rejection
 (project= no) Project highest dimension of input images?
 (outtype= real) Output image pixel datatype
 (outlimi= none) Output limits (x1 x2 y1 y2 ...)
 (offsets= none) Input image offsets
 (masktyp=) Mask type
 (maskval= 0.) Mask value
 (blank = 0.) Value if there are no pixels
 (scale = exposure) Image scaling
 (zero = none) Image zero point offset
 (weight = exposure) Image weights
 (statsec=) Image section for computing statistics
 (expname= exposure) Image header exposure time keyword
 (lthresh= INDEF) Lower threshold
 (hthresh= INDEF) Upper threshold
 (nlow = 0) minmax: Number of low pixels to reject
 (nhigh = 1) minmax: Number of high pixels to reject
 (nkeep = 1) Minimum to keep (pos) or maximum to reject
 (neg)
 (mclip = yes) Use median in sigma clipping algorithms?
 (lsigma = 3.) Lower sigma clipping factor
 (hsigma = 10.) Upper sigma clipping factor
 (rdnoise= 2.64)ccdclip: CCD readout noise (electrons)
 (gain = 2.25)ccdclip: CCD gain (electrons/DN)
 (snoise = 0.)ccdclip: Sensitivity noise (fraction)
 (sigscal= 0.1) Tolerance for sigma clipping scaling corrections
 (pclip = -0.5) pclip: Percentile clipping parameter
 (grow = 0.) Radius (pixels) for neighbor rejection
 (mode = ql)

 :g

2.7.5 对合并的图像手动去除宇宙线:

去除宇宙线是一个反复的过程,可以先粗略的去除,用 ds9 检查结果后再反复去除,以达到最佳效果;

```
im> crutil
cr> epa cosmicrays
I R A F
Image Reduction and Analysis Facility
PACKAGE = crutil
TASK = cosmicrays
input = w3caver List of images in which to detect cosmic rays
**再次去除时,要将此改为前一次去除后的输出图像,如w3caver.c
output = w3caver.c List of cosmic ray replaced output images (optional)
(crmasks= ) List of bad pixel masks (optional)
(thresho= 25.) Detection threshold above mean
(fluxrat= 2.) Flux ratio threshold (in percent)
**可先设置一个值,剔除一次宇宙线后可以再设置;
(npasses= 5) Number of detection passes
(window = 5) Size of detection window
(interac= yes) Examine parameters interactively?
(train = no) Use training objects?
(objects= ) Cursor list of training objects
(savefil= ) File to save train objects
(plotfil= ) Plot file
(graphic= stdgraph) Interactive graphics output device
(cursor = ) Graphics cursor input
answer = yes Review parameters for a particular image?
(mode = ql)
```

```
-----
:g
```

在交互式窗口中,鼠标放到某个小十字上,按 s 为查看该点的二维强度分布, d 为剔除,典型的宇宙线呈较平坦的面上突出的一个或较少得几个像素,如尖状;大多数情况需要反复剔除。

2.7.6 利用 `splot` 命令手动剔除宇宙线

在完成上述处理以及谱线抽取后，需要用`splot`命令显示抽取后的谱线，检查是否有宇宙线打在了目标上，如果有，需要进行手动剔除。

```
noao/onedspec
```

```
one> splot aobject.fits **完成光谱抽取后的图像，见第三章
```

```
: hist yes **直方图显示
```

按 `j` 删除宇宙线（宇宙线在直方图显示时，往往只占1~2像素，突然增强）

按 `i` 生成新的图像

第三章： 一维光谱处理

3.1 目标的光谱抽取

```
noao/twospec/ape/appall
```

```
cr> epa apall
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = apextract
```

```
TASK = apall
```

```

input = object.fits      List of input images  **经过图像处理的目标文件，习惯改成目标名字，方便识别，此处用object代表
(output = aobject      ) List of output spectra
(apertur=              ) Apertures
(format = multispec) Extracted spectra format
(referen=              ) List of aperture reference images
(profile=              ) List of aperture profile images
(interac=              yes) Run task interactively?
(find =                yes) Find apertures?
(recente=              yes) Recenter apertures?
(resize =              yes) Resize apertures?
(edit =                yes) Edit apertures?
(trace =               yes) Trace apertures?
(fittrac=              yes) Fit the traced points interactively?
(extract=              yes) Extract spectra?
(extras =              yes) Extract sky, sigma, etc.?
(review =              yes) Review extractions?
(line =                INDEF) Dispersion line
(nsum =                10) Number of dispersion lines to sum or median
                        # DEFAULT APERTURE PARAMETERS
(lower =                -5.) Lower aperture limit relative to center**设置目标孔径，可以在交互模式下更改
(upper =                5.) Upper aperture limit relative to center

```

```

(apidtab=                ) Aperture ID table (optional)
                        # DEFAULT BACKGROUND PARAMETERS
(b_funct=                chebyshev) Background function
(b_order=                1) Background function order
(b_sampl=                -10:-6,6:10) Background sample regions** 设置背景孔径
(b_naver=                -3) Background average or median
(b_niter=                0) Background rejection iterations
(b_low_r=                3.) Background lower rejection sigma
(b_high_=                3.) Background upper rejection sigma
(b_grow =                0.) Background rejection growing radius
                        # APERTURE CENTERING PARAMETERS
(width  =                5.) Profile centering width
(radius =                10.) Profile centering radius
(thresho=                0.) Detection threshold for profile centering
                        # AUTOMATIC FINDING AND ORDERING PARAMETERS
nfind  =                1  Number of apertures to be found automatically ** 对一个目标只需要一个孔径
(minsep =                5.) Minimum separation between spectra
(maxsep =                1000.) Maximum separation between spectra
(order  =                increasing) Order of apertures
                        # RECENTERING PARAMETERS
(aprecen=                ) Apertures for recentering calculation
(npeaks =                INDEF) Select brightest peaks
(shift  =                yes) Use average shift instead of recentering?
                        # RESIZING PARAMETERS
(llimit =                INDEF) Lower aperture limit relative to center
(ulimit =                INDEF) Upper aperture limit relative to center
(ylevel =                0.1) Fraction of peak or intensity for automatic width
(peak   =                yes) Is ylevel a fraction of the peak?
(bkg    =                yes) Subtract background in automatic width?
(r_grow =                0.) Grow limits by this factor
(avglimi=                no) Average limits over all apertures?
                        # TRACING PARAMETERS
(t_nsum =                10) Number of dispersion lines to sum
(t_step =                10) Tracing step

```

```

(t_nlost=          3) Number of consecutive times profile is lost before
quitting
(t_func=          spline3) Trace fitting function
(t_order=         3) Trace fitting function order
(t_sampl=         *) Trace sample regions
(t_naver=         1) Trace average or median
(t_niter=         0) Trace rejection iterations
(t_low_r=         3.) Trace lower rejection sigma
(t_high_=         3.) Trace upper rejection sigma
(t_grow =         0.) Trace rejection growing radius

# EXTRACTION PARAMETERS

(backgro=         fit) Background to subtract **在提取定标灯谱线时, 此处
为none
(skybox =         1) Box car smoothing length for sky
(weights=         variance) Extraction weights (none|variance)
(pfit  =         fit2d) Profile fitting type (fit1d|fit2d)
(clean  =         no) Detect and replace bad pixels?
(saturat=        INDEF) Saturation level
(readnoi=        2.64) Read out noise sigma (photons)
(gain   =        2.25) Photon gain (photons/data number)
(lsigma =        4.) Lower rejection threshold
(usigma =        4.) Upper rejection threshold
(nsubaps=        1) Number of subapertures per aperture
(mode   =        ql)

```

:g

在交互式窗口中的操作如下:

在峰的左侧按 **l**, 右侧按 **u**, 以设置孔径。要使得峰尽量被包括进孔径当中, 但尽量不要使背景被划入孔径; 见图 3.1.1.

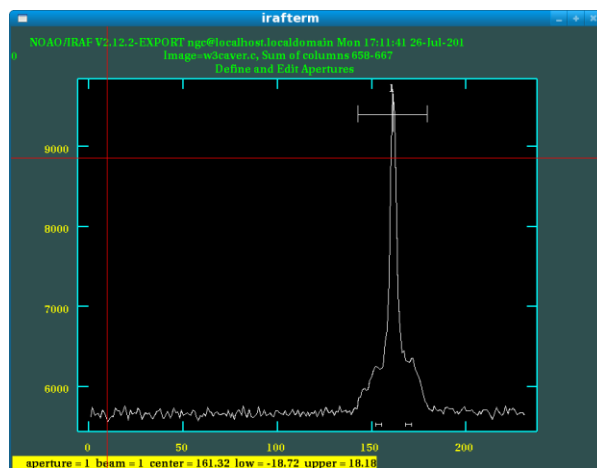


图 3. 1. 1

**：注意观察横坐标，此时横坐标为 y 方向，也就是最大 200 多大小的刻度。由于是一维光谱，所以只应该又一个峰；如果是高色散光谱，那么拍摄的原图有几个水平亮条纹，就对应几个级次，也应该有几个峰，应该选择数量相当的孔径。如果横坐标为 x 方向，说明色散轴的方向不对，需重新设置：

可用命令：

```
ap> epa apex
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = twodspec
```

```
TASK = apextract
```

```
(dispaxi=          1) Dispersion axis (1=along lines, 2=along columns) **dispaxi改为1
```

```
(databas=          database) Database
```

```
(verbose=          no) Verbose output?
```

```
(logfile=          ) Text log file
```

```
(plotfil=          ) Plot file
```

```
(version= APEXTRACT V3.0: August 1990)
```

```
(mode   =          ql)
```

```
($args =          0)
```

```
-----
```

```
:g
```

即可；

设置完 l 和 u 后，有效孔径选取完毕；

之后键入 b，来标定天光孔径，按 t 删除默认的天光孔径；随后我们要手动在 x 轴方向，对应原图的 y 方向选取背景的亮度；

具体做法是：在峰的左右两侧分别输入两次 s，即鼠标放在峰背景左侧的某处，按 s；

再移动鼠标到峰左侧另一处，再按 s，即圈定了一侧的天光孔径范围；

鼠标移到峰的右侧执行相同的操作，即可选定天光孔径；

需要按的键大致是:b-t-s-s-s-f-q。

3)在交互式窗口中，若需要确认，则按回车(yes)；

确认完成后，按 q 退出，查看拟合图；

此处的拟合参数在之前得 apall 当中已经设置过，但为了更好的拟合孔径，需要在交互式窗口中剔除坏点，并重新选择拟合方式和阶数；

具体做法是：在拟合图中，鼠标放到坏点上按 d 剔除，再键入

:f spline3 回车(yes) ****一般采取三次样条函数拟合；**

以及

:o 5 ****5为拟合阶数；**

可以选择别的阶数，但不应过低或过高，阶数过低难真实拟合曲线大致的轮廓；阶数过高则会拟合出一些精细的轮廓，每个点的影响都会加大，测量的结果会失真。

如图 3.1.2

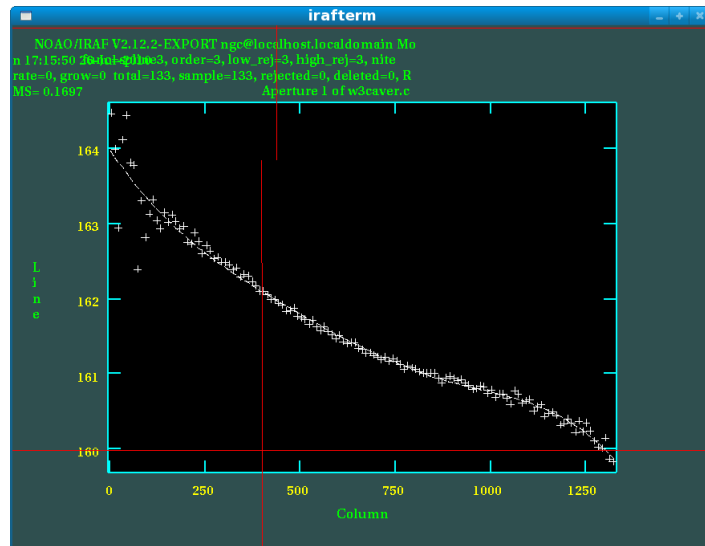


图 3.1.2

4) 如果确认孔径选取好了，拟合效果也不错，按 q 退出，对交互式窗口中弹出的问题全部回车(yes)，最后按 q 退出；

可以发现新生成了文件 aobject.fits；这就是抽取得到的目标的光谱，注意观

察横轴和纵轴坐标：此时得到的是未定标的谱。如图 3.1.3,之后要对它进行波长定标和流量定标:

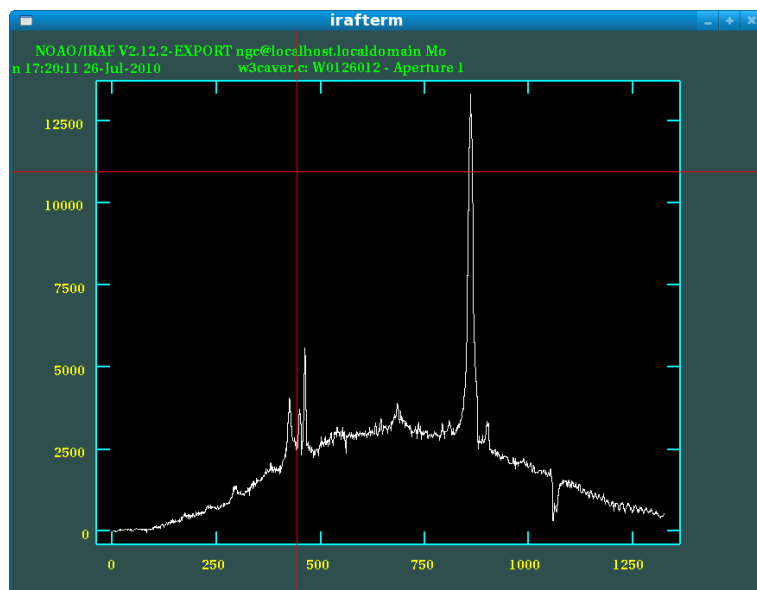


图 3. 1. 3

3.2 定标灯灯谱抽取

```
ap> epa apall
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = apextract
```

```
TASK = apall
```

```
input = Neon.fits List of input images ** 经过bias、flat处理后的Neon灯灯谱
```

```
(output = aNeon) List of output spectra ** 根据自身习惯命名, 方便以后处理
```

```
(apertur= ) Apertures
```

```
(format = onedspec) Extracted spectra format
```

```
(referen= object.fits) List of aperture reference images ** 以前面的目标为参考
```

```
(profile= ) List of aperture profile images
```

```
(interac= no) Run task interactively?
```

```
(find = no) Find apertures?
```

```
(recente= no) Recenter apertures?
```

```
(resize = no) Resize apertures?
```

```

(edit    =          no) Edit apertures?
(trace  =          no) Trace apertures?
(fittrac=         no) Fit the traced points interactively?
(extract=         yes) Extract spectra?
(extras =         no) Extract sky, sigma, etc.?
(review =         yes) Review extractions?
(line   =          INDEF) Dispersion line
(nsum   =          10) Number of dispersion lines to sum or median
                                # DEFAULT APERTURE PARAMETERS
(lower  =         -5.) Lower aperture limit relative to center
(upper  =          5.) Upper aperture limit relative to center
(apidtab=         ) Aperture ID table (optional)
                                # DEFAULT BACKGROUND PARAMETERS
(b_funct=        chebyshev) Background function
(b_order=          1) Background function order
(b_sampl=       -10:-6,6:10) Background sample regions
(b_naver=        -3) Background average or median
(b_niter=         0) Background rejection iterations
(b_low_r=         3.) Background lower rejection sigma
(b_high_=        3.) Background upper rejection sigma
(b_grow =         0.) Background rejection growing radius
                                # APERTURE CENTERING PARAMETERS
(width  =          5.) Profile centering width
(radius =         10.) Profile centering radius
(thresho=         0.) Detection threshold for profile centering
                                # AUTOMATIC FINDING AND ORDERING
PARAMETERS
nfind   =          1  Number of apertures to be found automatically
(minsep =          5.) Minimum separation between spectra
(maxsep =        1000.) Maximum separation between spectra
(order  =          increasing) Order of apertures
                                # RECENTERING PARAMETERS
(aprecen=         ) Apertures for recentering calculation
(npeaks =          INDEF) Select brightest peaks
(shift  =         yes) Use average shift instead of recentering?

```

RESIZING PARAMETERS

(llimit = INDEF) Lower aperture limit relative to center
 (ulimit = INDEF) Upper aperture limit relative to center
 (ylevel = 0.1) Fraction of peak or intensity for automatic width
 (peak = yes) Is ylevel a fraction of the peak?
 (bkg = yes) Subtract background in automatic width?
 (r_grow = 0.) Grow limits by this factor
 (avglimi= no) Average limits over all apertures?

TRACING PARAMETERS

(t_nsum = 10) Number of dispersion lines to sum
 (t_step = 10) Tracing step
 (t_nlost= 3) Number of consecutive times profile is lost before
 quitting
 (t_func= spline3) Trace fitting function
 (t_order= 3) Trace fitting function order
 (t_sampl= *) Trace sample regions
 (t_naver= 1) Trace average or median
 (t_niter= 0) Trace rejection iterations
 (t_low_r= 3.) Trace lower rejection sigma
 (t_high_= 3.) Trace upper rejection sigma
 (t_grow = 0.) Trace rejection growing radius

EXTRACTION PARAMETERS

(backgro= none) Background to subtract**此处选择none而不是fit, 因此对定标灯谱而言, 背景和目标是相同的
 (skybox = 1) Box car smoothing length for sky
 (weights= none) Extraction weights (none|variance)
 (pfit = fit1d) Profile fitting type (fit1d|fit2d)
 (clean = no) Detect and replace bad pixels?
 (saturat= INDEF) Saturation level
 (readnoi= 2.64) Read out noise sigma (photons)
 (gain = 2.25) Photon gain (photons/data number)
 (lsigma = 4.) Lower rejection threshold
 (usigma = 4.) Upper rejection threshold
 (nsubaps= 1) Number of subapertures per aperture
 (mode = ql)

```
:g
```

之后会发现生成了文件 aNeon.fits

可以用 `splot` 命令查看灯谱,

```
ap> splot aNeon.fits
```

如图 3.2.1(YFOOSC:Neon+G8+Slit1.2)

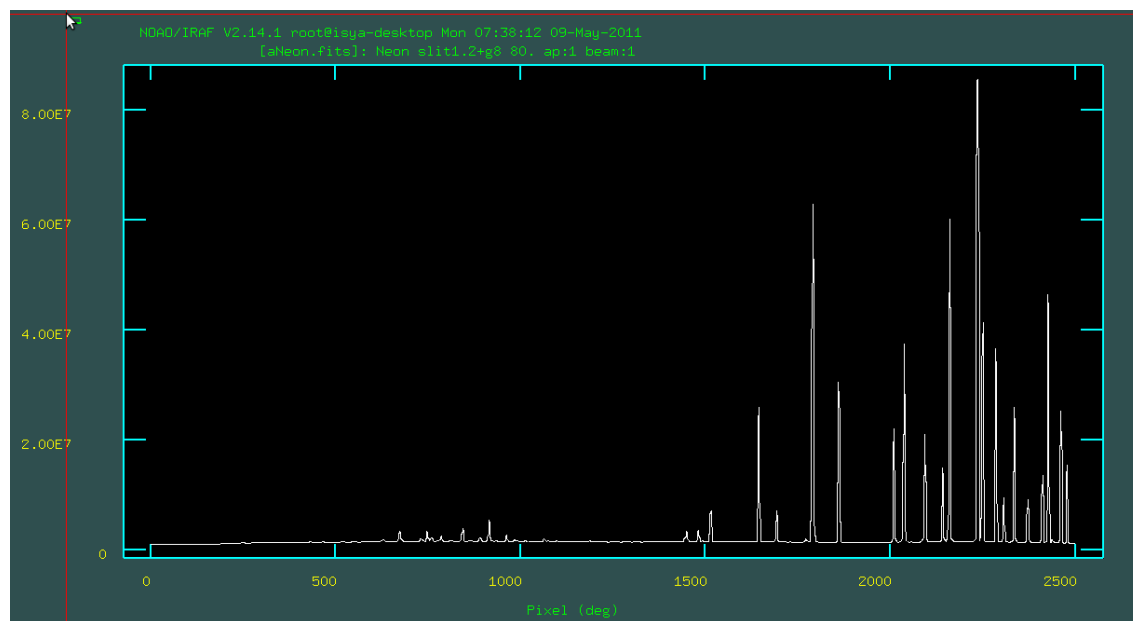


图 3.2.1

3.3 波长定标

3.3.1 波长证认

```
noao\onedspec\identify
```

```
one> epa identify
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = longslit
```

```
TASK = identify
```

```
images = aNeon Images containing features to be identified **抽出的灯谱
```

```
(section= middle line) Section to apply to two dimensional images
```

```
(databas= database) Database in which to record feature data
```

```
(coordli= linelists$henear.dat) User coordinate list **选择相对应的波长文件,此
```

处为氩氦氙灯谱文件

```

(units = ) Coordinate units
(nsum = 10) Number of lines/columns/bands to sum in 2D images
(match = -3.) Coordinate list matching limit
(maxfeat= 50) Maximum number of features for automatic
identification
(zwidth = 100.) Zoom graph width in user units
(ftype = emission) Feature type **有时候会用到吸收线, absorption
(fwidth = 4.) Feature width in pixels
(cradius= 5.) Centering radius in pixels
(thresho= 0.) Feature threshold for centering
(minsep = 2.) Minimum pixel separation
(funcio= spline3) Coordinate function
(order = 1) Order of coordinate function
(sample = *) Coordinate sample regions
(niterat= 0) Rejection iterations
(low_rej= 3.) Lower rejection sigma
(high_re= 3.) Upper rejection sigma
(grow = 0.) Rejection growing radius
(autowri= no) Automatically write to database
(graphic= stdgraph) Graphics output device
(cursor = ) Graphics cursor input
crval = Approximate coordinate (at reference pixel)
cdelt = Approximate dispersion
(aidpars= ) Automatic identification algorithm parameters
(mode = ql)

```

:g

在交互窗口按 m，输入已知谱线波长的整数部分再回车，输入几个又把握的值之后，再按 l 标谱，这时所有的谱线上面都会有一个黄色的小竖线，证明这些谱线已经标好，按 f 查看拟合情况。*对于 Prism，由于棱角分光时，色散能力随波长而变化，因此，在进行波长标定时，需要在红端、蓝端各选几条线。*也就是说对于非线性的色散，需要在不同区间进行波长定标。

对于方差过大的点，鼠标放在上面按 d 剔除，最终的拟合结果最好要有接近 30 个点，且方差小于 0.5。

3.3.3 对目标源和标准星分别加定标文件头

对目标源:

```
one> epa refspectra
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = specred
```

```
    TASK = refspectra
```

```
input   =          aobject List of input spectra
(referen=          aNeon) List of reference spectra
(apertur=          ) Input aperture selection list
(refaps =          ) Reference aperture selection list
(ignorea=          yes) Ignore input and reference apertures?
(select =          interp) Selection method for reference spectra
(sort   =          ) Sort key
(group  =          ) Group key
(time   =          no) Is sort key a time?
(timewra=          17.) Time wrap point for time sorting
(overrid=          no) Override previous assignments?
(confirm=          yes) Confirm reference spectrum assignments?
(assign =          yes) Assign the reference spectra to the input spectrum?
(logfile=          STDOUT,logfile) List of logfiles
(verbose=          no) Verbose log output?
answer  =          yes  Accept assignment?
(mode   =          ql)
```

```
-----
:g
```

也可以使用 `hedit` 命令添加 `DCLOG1 = 'REFSPEC1 = aNeon'`

对标准星的操作与目标星完全一样，在此，我们也对定标灯的头文件进行修改，即，用 `aNeon` 作为 `aNeon` 的参考。下一步对 `aNeon` 进行色散轴改正。

3.3.4 对目标源和标准星分别进行色散轴改正

对 `aNeon`，利用 `dispcor` 命令


```
one> epa dispcor
```

```
I R A F
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = onedspec
```

```
  TASK = dispcor
```

```
input   =          aNeon  List of input spectra
```

```
output  =          wNeon  List of output spectra
```

```
(lineari=          yes) Linearize (interpolate) spectra?
```

```
(databas=          database) Dispersion solution database
```

```
(table   =          ) Wavelength table for apertures
```

```
(w1      =          INDEF) Starting wavelength
```

```
(w2      =          INDEF) Ending wavelength
```

```
(dw      =          INDEF) Wavelength interval per pixel
```

```
(nw      =          INDEF) Number of output pixels
```

```
(log     =          no) Logarithmic wavelength scale?
```

```
(flux    =          yes) Conserve flux?
```

```
(samedis=          no) Same dispersion in all apertures?
```

```
(global  =          no) Apply global defaults?
```

```
(ignorea=          no) Ignore apertures?
```

```
(confirm=          no) Confirm dispersion coordinates?
```

```
(listonl=          no) List the dispersion coordinates only?
```

```
(verbose=          yes) Print linear dispersion assignments?
```

```
(logfile=          ) Log file
```

```
(mode    =          ql)
```

```
-----  
:g
```

```
得到文件 wNeon;
```

```
ecl> splot wNeon **x轴此时显示的是波长
```

```
如图 3.4.1 所示:
```

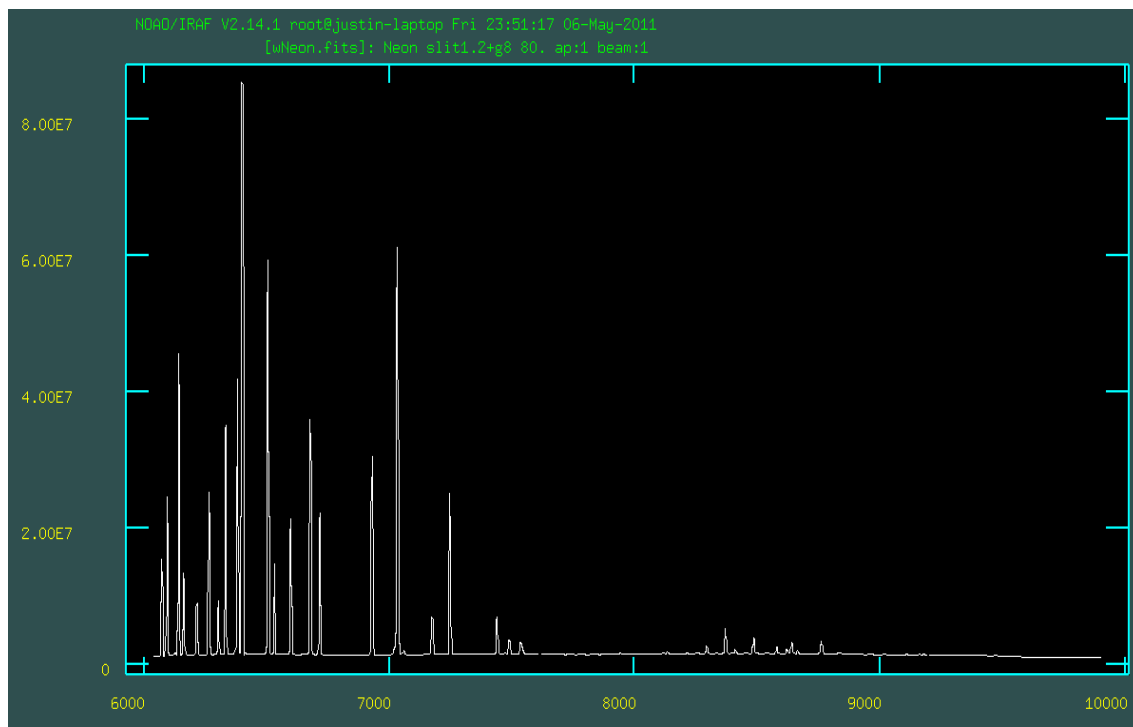


图 3.4.1

3.4 流量定标

3.4.1 对标准星加文件头，生成定标函数

```
two> longslit
lo> epa standard
I R A F
Image Reduction and Analysis Facility
PACKAGE = longslit
    TASK = standard
input   = dwstandcaver   Input image file root name **波长定标后的标准星图像
output  = std           Output flux file (used by SENSFUNC)
(samesta=                               yes) Same star in all apertures?
(beam_sw=                               no) Beam switch spectra?
(apertur=                               ) Aperture selection list
(bandwid=                               30.) Bandpass widths
(bandsep=                               20.) Bandpass separation
(fnuzero= 3.6800000000000000E-20) Absolute flux zero point
(extinct= workstation/LJextinct.dat) Extinction file **丽江观测站的大气消光文件
```

LJextinct.dat, 地址为该文件所在目录, 本人习惯有一个专门的数据处理区

```
(caldir =          )_caldir) Directory containing calibration data
(observa=          )_observatory) Observatory for data
(interac=          yes) Graphic interaction to define new bandpasses
(graphic=          stdgraph) Graphics output device
(cursor =          ) Graphics cursor input
star_nam=          Hiltner600 Star name in calibration list **标准星的星名
answer =          yes (no|yes|NO|YES|NO!|YES!)
(mode =           ql)
```

:g

如果 fits 头文件中没有 airmass, 则会出现如下错误:

```
dwstandcaver.c.ms: ERROR: parameter `airmass' not found
standard (mode=h)
```

在另一个关于脚本的文档中, 将提到如何大批量为 fits 文件计算并添加 airmass 等信息, 此处暂时用手动的方式添加, 同时也需要对目标星添加 airmass 信息。

```
lo> epa hedit
```

```
I R A F
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = imutil
```

```
  TASK = hedit
```

```
images = dwstandcaver.fits  images to be edited
```

```
fields =          airmass  fields to be edited
```

```
value =          1.16  value expression
```

```
(add =          yes) add rather than edit fields
```

```
(addonly=       yes) add only if field does not exist
```

```
(delete =      no) delete rather than edit fields
```

```
(verify =      no) verify each edit operation
```

```
(show =       yes) print record of each edit operation
```

```
(update =     yes) enable updating of the image header
```

```
(mode =       ql)
```

:g

这样就可以正常运行 standard 了;

新生成文件 std;

3.4.2 生成定标函数:

```
lo> epa sensfunc
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = longslit
```

```
TASK = sensfunc
```

```
standard=          std  Input standard star data file (from STANDARD)
```

```
sensitiv=          sens  Output root sensitivity function imagename
```

```
(apertur=          ) Aperture selection list
```

```
(ignorea=          yes) Ignore apertures and make one sensitivity function?
```

```
(logfile=          logfile) Output log for statistics information
```

```
(extinct=          )_extinction) Extinction file
```

```
(newexti=workstation/LJextinct.dat) Output revised extinction file ** 依旧为之前的 LJextinct.dat 所在目录
```

```
(observa=          bao) Observatory of data
```

```
(functio=          spline3) Fitting function
```

```
(order  =          6) Order of fit
```

```
(interac=          yes) Determine sensitivity function interactively?
```

```
(graphs =          sr) Graphs per frame
```

```
(marks  =          plus cross box) Data mark types (marks deleted added)
```

```
(colors =          2 1 3 4) Colors (lines marks deleted added)
```

```
(cursor =          ) Graphics cursor input
```

```
(device =          stdgraph) Graphics output device
```

```
answer  =          yes (no|yes|NO|YES)
```

```
(mode   =          ql)
```

```
-----  
:g
```

在交互界面按回车后, 用 d-p 消除坏点, f 为重新拟合。如图 3.4.1;

得到满意的拟合曲线后按 q 退出; 产生文件 sens.0001.fits, 如图 3.4.2;

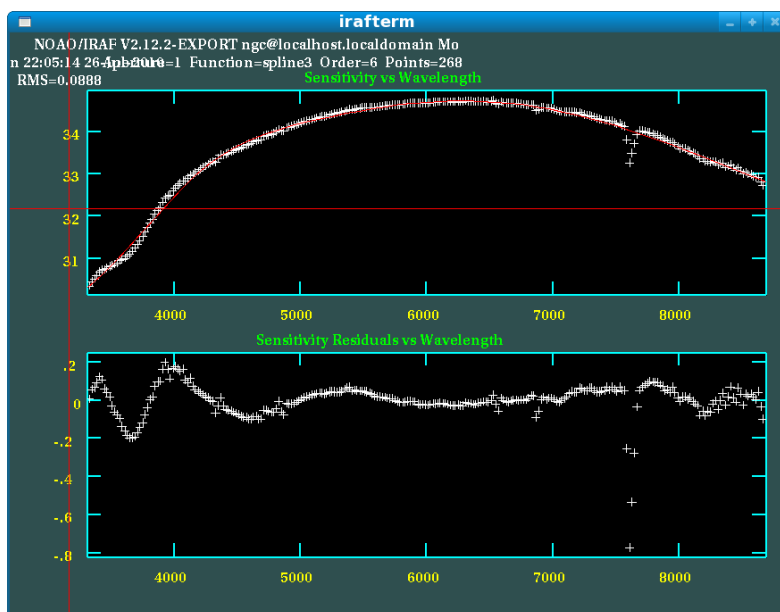


图 3.4.2

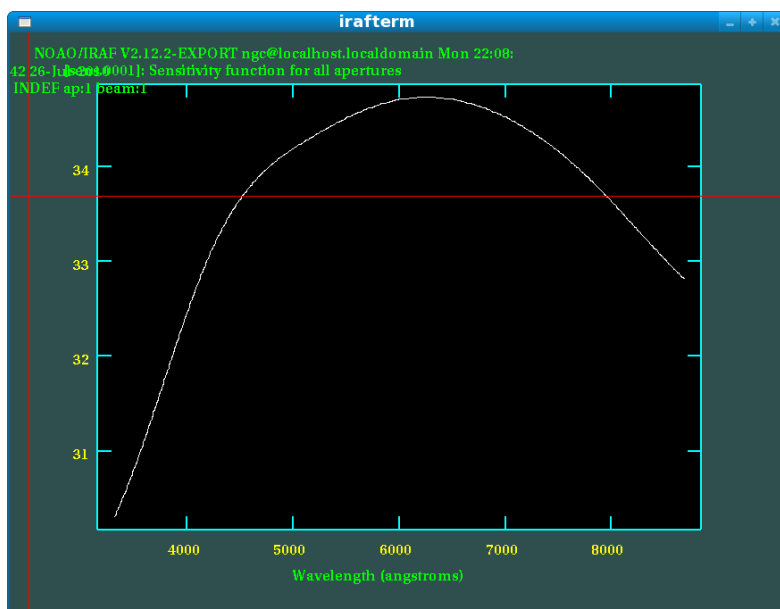


图 3.4.3

3.4.3 流量定标:

```
lo> epa calibrate
```

```
IRAF
```

```
Image Reduction and Analysis Facility
```

```
PACKAGE = longslit
```

```
TASK = calibrate
```

```
input =          wobject  Input spectra to calibrate ** 输入经过波长定标后的目标源
```

```

output =          cwobject  Output calibrated spectra
(extinct=          yes) Apply extinction correction?
(flux   =          yes) Apply flux calibration?
(extinct= workstation/LJoextinct.dat) Extinction file
(observa=          bao) Observatory of observation
(ignorea=          yes) Ignore aperture numbers in flux calibration?
(sensiti=          sens.0001) Image root name for sensitivity spectra **新生成的
sens.0001文件
(fnu    =          no) Create spectra having units of FNU?
(mode   =          ql)

```

:g

运行后，当xgterm中显示：

cwobject:object

extinction correction applied

flux calibration applied

至此，抽谱以及定标全部完成；

3.4.4 将结果转成一维文件

lo>epa scopy

IRAF

Image Reduction and Analysis Facility

PACKAGE = longslit

TASK = scopy

```

input   = cwobject      List of input spectra **定标全部完成后的输出文件
output  = markobject.fits List of output spectra
(w1     =               INDEF) Starting wavelength
(w2     =               INDEF) Ending wavelength
(apertur=               ) List of input apertures or columns/lines
(bands  =               1) List of input bands or lines/bands
(beams  =               ) List of beams or echelle orders
(apmodul=               0) Input aperture modulus (0=none)
(format =               onedspec) Output spectra format
(renumbe=               no) Renumber output apertures?

```

```

(offset = 0) Output aperture number offset
(clobber= no) Modify existing output images?
(merge = no) Merge with existing output images?
(rebin = yes) Rebin to exact wavelength region?
(verbose= no) Print operations?
(mode = ql)

```

```

:g

```

生成文件 mark618-1.fits

用 `splot` 查看:

```

lo> splot mark618-1.fits.0001.fits

```

即为最终结果，如图 3.4.3:

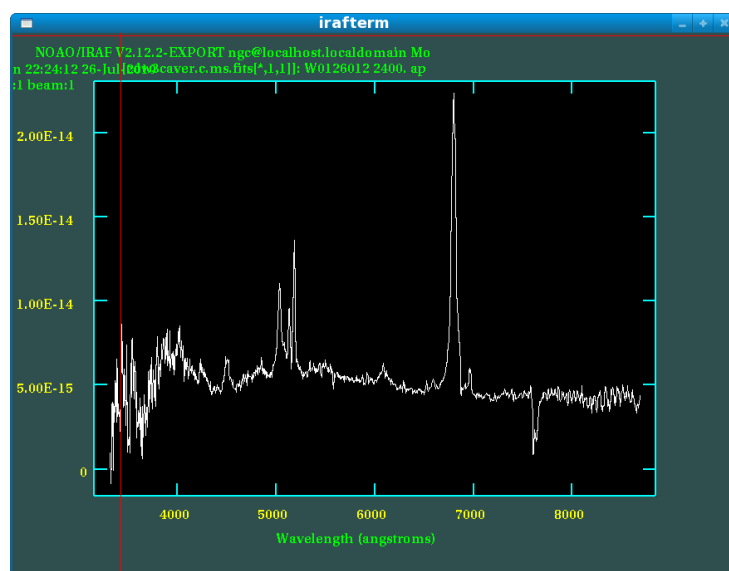


图 3.4.3

3.5 连续谱归一化

这一步并不是必须的，很多时候对光谱分析而言，连续谱的信息很重要，但是对于谱线位置等相对测量而言，去除背景的影响，能更好的标定波长，这时候需要对定标后的图像进行连续谱的归一化，用到 `noao\onedspec\continuum` 命令。

```

PACKAGE = onedspec

```

```

TASK = continuum

```

```

input = wobject Input images

```

```

output = conobject Output images

```

```

(lines = *) Image lines to be fit

```

(bands = 1) Image bands to be fit

(type = *ratio*) Type of output

(replace= no) Replace rejected points by fit?

(wavesca= yes) Scale the X axis with wavelength?

(logscal= no) Take the log (base 10) of both axes?

(overrid= no) Override previously fit lines?

(listonl= no) List fit but don't modify any images?

(logfile= logfile) List of log files

(interac= yes) Set fitting parameters interactively?

(sample = *) Sample points to use in fit

(naverag= 1) Number of points in sample averaging

(functio= spline3) Fitting function

(order = 1) Order of fitting function

(low_rej= 2.) Low rejection in sigma of fit

(high_re= 0.) High rejection in sigma of fit

(niterat= 10) Number of rejection iterations

(grow = 1.) Rejection growing radius in pixels

(markrej= yes) Mark rejected points?

(graphic= stdgraph) Graphics output device

(cursor =) Graphics cursor input

ask = yes

(mode = ql)

生成归一化的 conobject.fits

用 `plot` 命令显示，以 YFOSC 拍摄的 Hiltner600 为例

