

Restart the Maple server to clear old definitions, etc.

```
> restart;
```

Load the special Maple tools packages needed for operations within the *MCMC\_SPIFI* procedure.

```
> with(plots): with(plottools): with(stats): with(statplots): with
  (StringTools):
```

Warning, the name changecoords has been redefined

Warning, the assigned name arrow now has a global binding

Warning, the previous binding of the name transform has been removed and it now has an assigned value

Warning, the assigned name Group now has a global binding

\*\*\*\*\*

Initialize the procedure

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The *MCMC\_SPIFI\_manualXY* procedure takes the following arguments:

*Description*: a string that will appear in the final spectrum title.

*X*: data array of velocity values (in km/s).

*Y*: data array of main beam temperature values (in K).

*RasterPosition*: the raster position of the current data, in the form: (delta "row pixels", delta "column pixels").

*BadBins*: a list (enclosed in square brackets and separated by commas) of bins to be removed from the spectrum before the fit is attempted. If no bins are to be removed, an empty list must be entered: [ ].

*CParam*: a list of the minimum, maximum, step size, and initial value to use when fitting the Lorentzian centroid.

*FParam*: a list of the minimum, maximum, step size, and initial value to use when fitting the Lorentzian FWHM.

*OParam*: a list of the minimum, maximum, step size, and initial value to use when fitting the baseline offset. In this program, I have defined the baseline as  $y=m(x-\text{average}(x[i]))+b$ , such that the offset,  $b$ , is the offset at the center of the spectrum, not at  $x=0$ .

*SParam*: a list of the minimum, maximum, step size, and initial value to use when fitting the baseline slope.

*HParam*: a list of the minimum, maximum, step size, and initial value to use when fitting the Lorentzian height.

*N*: total number of steps to attempt in your Markov Chain random walk.

*BadN*: number of steps to throw out from the beginning of the Markov Chain (start-up iterations). This is only necessary if you choose an initial point far from the  $\chi^2$  minimum.

*sigma*: a parameter that controls the size of proposed steps in the Markov Chain. More precisely, *sigma* is standard deviation of the random normal distribution which is used to determine the Markov step size.

*r2*: a parameter that controls the acceptance rate of proposed steps in the Markov Chain. However, in principle the acceptance rate is best controlled with *sigma*: while increasing *r2* can lower the standard for acceptance, lowering *sigma* can raise the actual number of points with good fits to the data, which is a much more desirable means of increasing the acceptance rate.

*Nvar*: Number of unconstrained variables. (Degrees of Freedom (d.o.f.) = *NumBins* - *Nvar*).

*RandSeed*: the seed to use when generating random numbers. It can be set to either an integer or to 'random'.

*AdditionalError*: sometimes the  $\chi^2$  of a fit is so large that the significance,  $\alpha$ , is too small to be computed to the digits of precision in the program (currently Digits=50). This means that we have grossly underestimated the real error of the data, and must add some additional systematic error. This parameter allows the user to add some error (in units of K), which will add in quadrature to the previously calculated error of the data.

```
> MCMC_SPIFI_manualXY:=proc(Description,RasterPosition,X,Y,
  AvgError,BadBins,CParam,FParam,SParam,OParam,HParam,N,BadN,sigma,
  r2,Nvar,RandSeed,AdditionalError)
```

Define the local variables. These variables will only retain their definitions within the *MCMCGridSPIFI* procedure.

```
> local
```

```
  i, j, k,
```

```
  CMin, CMax, dC, C0, Cmean, Csigma, Cgrid, Cfit,
```

```
  FMin, FMax, dF, F0, Fmean, Fsigma, Fgrid, Ffit,
```

```
  SMin, SMax, dS, S0, Smean, Ssigma, Sgrid, Sfit,
```

```
  OMin, OMax, dO, O0, Omean, Osigma, Ogrid, Ofit,
```

```
HMin, HMax, dH, H0, Hmean, Hsigma, Hgrid, Hfit,
XBadBins, YBadBins,
Mins, Maxs, ds,
Mnew, Seed, Rand1, Rand2, r1, r, Naccept, AcceptRate, Nbestfit,
pegged,
M, L1, L2, chi1, chi2, chif, a, a1, a2, af, amax, Baseline,
SysErrorSqd, TotError, Signal, Noise, SNR, Intensity, Isigma:
```

Define the global variables. These are variables that retain their definitions outside of the *MCMCGridSPIFI* procedure (including its sub-procedures).

```
> global
XBins, YBins, XScaled, YScaled, XFinal, YFinal, Lf,
graph1a, graph1b, graph2a, graph2b, graph3a, graph3b, graph3c,
title1, title2, title3,
AvgErrorScaled, NumBins, BinLength, BinLengthScaled, Histogram,
Cpdf, Fpdf, Spdf, Opdf, Hpdf:
```

Rename and reorganize the fitting parameters:

```
> CMin:=CParam[1]: CMax:=CParam[2]: dC:=CParam[3]: C0:=CParam[4]:
> FMin:=FParam[1]: FMax:=FParam[2]: dF:=FParam[3]: F0:=FParam[4]:
> SMin:=SParam[1]: SMax:=SParam[2]: dS:=SParam[3]: S0:=SParam[4]:
> OMin:=OParam[1]: OMax:=OParam[2]: dO:=OParam[3]: O0:=OParam[4]:
> HMin:=HParam[1]: HMax:=HParam[2]: dH:=HParam[3]: H0:=HParam[4]:
> Mins:=[CMin,FMin,SMin,OMin,HMin]: Maxs:=[CMax,FMax,SMax,OMax,
HMax]: ds:=[dC,dF,dS,dO,dH]:
```

Specify the number of decimal places to carry through calculations.

```
> Digits:=50:
> interface(prettyprint=0): interface(displayprecision=20):
```

```
*****
```

Read and display raw data

```
*****
```

Call a sub-procedure, named *HistDiscont*, that creates histogram-style displays of data. The suffix "Discont" refers to the fact that it can create histograms of discontinuous data sets (i.e. where bins have been removed).

```
> read("C://Documents and Settings/Tom Oberst/Desktop/SPIFI &
ZEUS/SPIFI observing plan/Maple Worksheets/SPIFI/SPIFI histogram
discontinuous v6.mpl"):
> graph1a:=pointplot({seq([X[k],Y[k]], k=1..nops(X))}):
> HistDiscont(X,Y): graph1b:=Histogram:
> title1:=cat(`[NII] 205 micron scaled data`,`\n`,` , SPIFI Car II
Raster Position `,RasterPosition):
```

```
*****
```

Scale data and display scaled data

\*\*\*\*\*

Remove bins:

```
> NumBins:=nops(X)-nops(BadBins): BinLength:=(X[1]-X[nops(X)])/(nops(X)-1):
> for i from 1 to nops(BadBins) do XBadBins[BadBins[i]]:=DEL: YBadBins[BadBins[i]]:=DEL: end do:
> j:=1:
> for i from 1 to nops(X) do
> if XBadBins[i]<>DEL then XBins[j]:=X[i]: YBins[j]:=Y[i]: j:=j+1: end if:
> end do:
```

Scale Data (because X and Y values are entered manually already in units of K and km/s, no scaling is necessary):

```
> for i from 1 to NumBins do
> YScaled[i]:=YBins[i]:
> XScaled[i]:=XBins[i]:
> end do:
> AvgErrorScaled:=sqrt(AdditionalError^2+AvgError^2):
> BinLengthScaled:=BinLength:
```

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Output "header"

\*\*\*\*\*

User Inputs:

Description

Raster Position

X1 (km/s)

Y1 (K)

Average Error "AvgError" (K)

Bins Removed; "BadBins"

Centroid Min (km/s)

Centroid Max (km/s)

Centroid s (km/s)

Centroid0 (km/s)

FWHM Min (km/s)

FWHM Max (km/s)

FWHM s (km/s)

FWHM0 (km/s)

Slope Min (K/km/s)

Slope Max (K/km/s)

Slope s (K/km/s)

Slope0 (K/km/s)

Offset Min (K)

Offset Max (K)

Offset s (K)

Offset0 (K)

Height Min (K)  
Height Max (K)  
Height s (K)  
Height0 (K)  
MCMC steps; "N"  
MCMC initial steps to throw out; "BadN"  
sigma of MCMC random gaussian  
acceptance aggression; "r2"  
unconstrained variables; "Nvar"  
Random seed; "RandSeed"  
Additional User Defined Error (K); "AdditionalError"

Read-in Data:

Transmission (%)  
Antenna Velocity (km/s)  
Flatfielding  
Gain (mV/K)  
Efficiency (%)  
"Flat Velocity" (counts)  
HOFPI offset (counts)  
HOFPI slope (km/s per count)

```
print(Description):  
print(RasterPosition):  
print(X[1]):  
print(Y[1]):  
print(AvgError):  
print(BadBins):  
print(CMin):  
print(CMax):  
print(dC):  
print(C0):  
print(FMin):  
print(FMax):  
print(dF):  
print(F0):  
print(SMin):  
print(SMax):  
print(dS):  
print(S0):  
print(OMin):  
print(OMax):  
print(dO):  
print(O0):  
print(HMin):  
print(HMax):
```

```

print(dH):
print(H0):
print(N):
print(BadN):
print(sigma):
print(r2):
print(Nvar):
print(`see User Outputs below`):
print(AdditionalError):

```

```

print(Transmission):
print(AntennaVelocity):
print(Flatfield[Pixel]):
print(Gain):
print(Efficiency):
print(FlatVelocity[Pixel]):
print(HOFPIOffset):
print(HOFPISlope):

```

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Markov Chain Monte Carlo

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Calculate the figures of merit for the first point in the Markov Chain,  $M_0$ :

```

> M[0]:=[C0,F0,S0,O0,H0]:
Lorentzian (the baseline is defined such that the offset occurs at the spectrum's center, not at  $x=0$ ):
> L1:=unapply(H0/(1+((x-C0)/(F0/2))^2)+(S0*(x-sum(XScaled[k],k=1..
    NumBins)/NumBins)+O0), x):

```

Chi squared:

```

> ch1:=sum((L1(XScaled[k])-YScaled[k])^2, k=1..NumBins)/
    (AvgErrorScaled^2):

```

Alpha ("confidence" or "significance level"):

```

> a1:=1-statevalf[cdf,chisquare[NumBins-Nvar]](ch1):

```

The following variables need initialized before starting the Markov Chain. Naccept is a tally of the number of times a proposed point Mnew is accepted, a is an array of the confidences of each point visited in the Markov Chain, Nbestfit is the position (i.e. iteration number) of the maximum value in a, Mnew is the first proposed step in the MCMC, and Seed is used to generate random numbers based on a user-supplied integer or your computer's clock:

```

> Naccept:=0: a[0]:=a1: Nbestfit:=0: Mnew:=[C0,F0,S0,O0,H0]:
> if type(RandSeed, integer)=true then Seed:=randomize(RandSeed):
    else Seed:=randomize(): end if:

```

Do the Markov Chain loop:

```

> for i from 1 to N do
Generate a new proposed point, Mnew:
> Rand1:=[random[normald[0,sigma]](5)]: pegged:=false:
> for j from 1 to 5 do Mnew[j]:=M[i-1][j]+ds[j]*Rand1[j]:

```

```

> if Mnew[j]>Maxs[j] then Mnew[j]:=Maxs[j]: pegged:=true:
> elif Mnew[j]<Mins[j] then Mnew[j]:=Mins[j]: pegged:=true:
> end if: end do:
Calculate the figures of merit for the proposed point, Mnew:
> L2:=unapply(Mnew[5]/(1+((x-Mnew[1])/(Mnew[2]/2))^2)+(Mnew[3]*(x-
  sum(XScaled[k],k=1..NumBins)/NumBins)+Mnew[4]), x):
> chi2:=sum((L2(XScaled[k])-YScaled[k])^2, k=1..NumBins)/
  (AvgErrorScaled^2):
> a2:=1-statevalf[cdf,chisquare[NumBins-Nvar]](chi2):
Compare Mnew with the last point in the Markov Chain, M[i-1], and decide whether to accept or reject it:
> r1:=a2/a1: r:=r1*r2: print(chi1, chi2, a1, a2, r):
> Rand2:=evalf(rand(0..1000())/1000):
> if Rand2 <= r then
> M[i]:=Mnew:
> a[i]:=a2:
> if a[i]>max(seq(a[k],k=0..i-1)) and pegged=false then Nbestfit:=i
  end if:
> a1:=a2:
> Naccept:=Naccept+1:
> else M[i]:=M[i-1]:
> a[i]:=a1:
> end if:
> end do:

```

Calculate the acceptance rate of your MCMC. This can be adjusted using r2 (raising r2 lowers the standard for accepting a proposed point), or, preferably, using sigma (lowering sigma increases the number of proposed points that are good fits to the data). Empirical data suggests aiming for *AcceptRate* ~ 25% for data with >2 d.o.f. (Gregory 2005) .

```

> AcceptRate:=100*evalf(Naccept/N):

```

Calculate parameter pdf's, means, and sigmas:

```

> Cpdf:=histogram([seq(M[i][1], i=BadN..N)], area=N):
> Cmean:=describe[mean]([seq(M[i][1], i=BadN..N)]):
> Csigma:=describe[standarddeviation]([seq(M[i][1], i=BadN..N)]):

> Fpdf:=histogram([seq(M[i][2], i=BadN..N)], area=N):
> Fmean:=evalf(describe[mean]([seq(M[i][2], i=BadN..N)])):
> Fsigma:=describe[standarddeviation]([seq(M[i][2], i=BadN..N)]):

> Spdf:=histogram([seq(M[i][3], i=BadN..N)], area=N):
> Smean:=describe[mean]([seq(M[i][3], i=BadN..N)]):
> Ssigma:=describe[standarddeviation]([seq(M[i][3], i=BadN..N)]):

> Opdf:=histogram([seq(M[i][4], i=BadN..N)], area=N):
> Omean:=describe[mean]([seq(M[i][4], i=BadN..N)]):
> Osigma:=describe[standarddeviation]([seq(M[i][4], i=BadN..N)]):

```

```
> Hpdf:=histogram([seq(M[i][5], i=BadN..N)], area=N):
> Hmean:=describe[mean]([seq(M[i][5], i=BadN..N)]):
> Hsigma:=describe[standarddeviation]([seq(M[i][5], i=BadN..N)]):
```

\*\*\*\*\*

Use the point in the MCMC with maximum alpha (*Nbestfit*) to do final fit

\*\*\*\*\*

Compute the baseline and subtract it from the data points. (The baseline is defined such that the offset is at the center of the spectrum, not at  $x=0$ ):

```
> Baseline:=unapply(M[Nbestfit][3]*(x-sum(XScaled[k],k=1..NumBins)
/NumBins)+M[Nbestfit][4], x):
> for i from 1 to NumBins do YFinal[i]:=YScaled[i]-Baseline(XScaled
[i]): XFinal[i]:=XScaled[i]: end do:
```

Compute the Lorentzian (without the baseline, since the baseline has now been subtracted from the data):

```
> Lf:=unapply(M[Nbestfit][5]/(1+((x-M[Nbestfit][1])/(M[Nbestfit][2]
/2))^2), x):
> chif:=sum((Lf(XFinal[k])-YFinal[k])^2, k=1..NumBins)/
(AvgErrorScaled^2):
> af:=1-statevalf[cdf,chisquare[NumBins-Nvar]](chif):
```

If  $\chi^2$  is not equal to 1, then there is some systematic error that has been underestimated or overestimated. The total error (TotError) is the sum of the calculated error (AvgErrorScaled) and the systematic error (SysError):

```
> SysErrorSqd:=fsolve(sum((Lf(XFinal[k])-YFinal[k])^2,k=1..NumBins)
/(AvgErrorScaled^2+x)/(NumBins-Nvar)=1,x):
> TotError:=sqrt(AvgErrorScaled^2+SysErrorSqd):
```

Compute the Signal, Noise, SNR, Intensity, and Intensity error:

```
> Signal:=evalf((Pi/2)*M[Nbestfit][5]*(M[Nbestfit][2]
/BinLengthScaled)):
> Noise:=evalf(sqrt((TotError^2)*((Pi/2)*M[Nbestfit][2]
/BinLengthScaled))):
> SNR:=Signal/Noise:
> Intensity:=Signal*BinLengthScaled:
> Isigma:=Noise*BinLengthScaled:
```

Produce a graph of the final fit to the data:

```
> graph3a:=pointplot({seq([XFinal[k],YFinal[k]], k=1..NumBins)}):
> HistDiscont(convert(XFinal,list),convert(YFinal,list)): graph3b:=
Histogram:
> graph3c:=plot(Lf(x), x=XFinal[NumBins]-BinLengthScaled/2..XFinal
[1]+BinLengthScaled/2):
> title3:=cat(Description,`\n`,`SPIFI Raster Position `,
RasterPosition):
```

\*\*\*\*\*

Print outputs

\*\*\*\*\*

fitting method  
random number generator seed; "Seed"  
MCMC acceptance rate (%)  
final reduced  $\chi^2$ ; "chif/(NumBins-Nvar)"  
final confidence; "af" (%)  
Known Error; "AvgErrorScaled" (K)  
Unknown Error Squared; "SysErrorSqd" ( $K^2$ )  
Total Error; "TotError" (K)  
Signal ( $K \cdot N_{\text{bins}}$ )  
Noise ( $K \cdot \sqrt{N_{\text{bins}}}$ )  
SNR  
Intensity ( $K \cdot \text{km/s}$ )  
Intensity sigma ( $K \cdot \text{km/s}$ )  
Centroid (km/s)  
Centroid sigma (km/s)  
FWHM (km/s)  
FWHM sigma (km/s)  
Slope ( $K/\text{km/s}$ )  
Slope sigma ( $K/\text{km/s}$ )  
Offset (K)  
Offset sigma (K)  
Height (K)  
Height sigma (K)  
Height sigma alternative (non-Gauss pdf) (K)  
Final list of X values (km/s)  
Final list of Y values (K)

```
> print(`MCMC min fit`):  
> print(Seed):  
> print(AcceptRate):  
> print(chif/(NumBins-Nvar)):  
> print(100*af):  
> print(AvgErrorScaled):  
> print(SysErrorSqd):  
> print(TotError):  
> print(Signal):  
> print(Noise):  
> print(SNR):  
> print(Intensity):  
> print(Isigma):  
> print(M[Nbestfit][1]):  
> print(Csigma):  
> print(M[Nbestfit][2]):  
> print(Fsigma):  
> print(M[Nbestfit][3]):  
> print(Ssigma):  
> print(M[Nbestfit][4]):
```



```

> print(Osigma):
> print(M[Nbestfit][5]):
> print(Hsigma):
> print(TotError*sqrt(chif/(NumBins-Nvar))):
> for i from 1 to nops(X) do print(XFinal[i]) od:
> for i from 1 to nops(X) do print(YFinal[i]) od:

*****

> end proc:
> MCMC_SPIFI_manualXY(`[NII] 205 micron emission in Car I`, `(6,7),
  LBC`, [236.259,205.586,174.914,144.242,113.570,82.898,52.226,
  21.553,-9.119,-39.791,-70.463,-101.135,-131.808,-162.480,
  -193.152,-223.824],[0.1389222,-0.1103345,-0.0928093,0.0859553,
  -0.2711668,-0.0895290,0.0311870,-0.0381949,0.0382353,0.1664311,
  0.5845062,0.4836142,-0.2317817,0.0199565,0.0130529,0.3970861],
  0.205363363,[],[-200,150,15,-30],[61,185,15,90],[-0.005,-0.00025,
  5.0*10^(-4),-3.0*10^(-4)],[-6,6,.08,0.0],[0,2,.15,.35],5000,0,1,
  1,5,random,0):
`[NII] 205 micron emission in Car I`
`(6,7), LBC`
236.259
.1389222
.205363363
[]
-200
150
15
-30
61
185
15
90
-0.5e-2
-0.25e-3
0.5000000000e-3
-0.3000000000e-3
-6
6
0.8e-1
0.
0
2

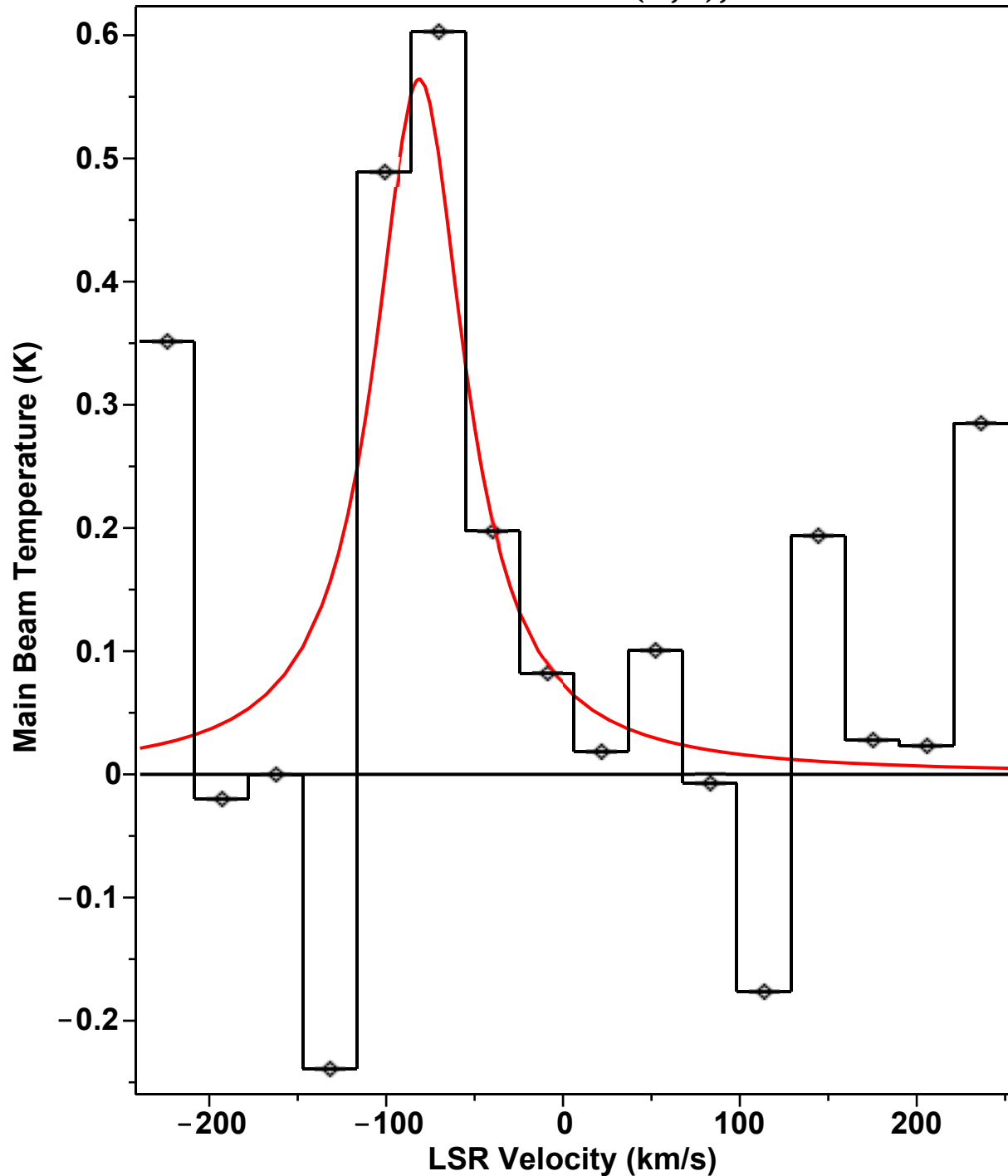
```

.19635533951553301226850254831104637943134367881607

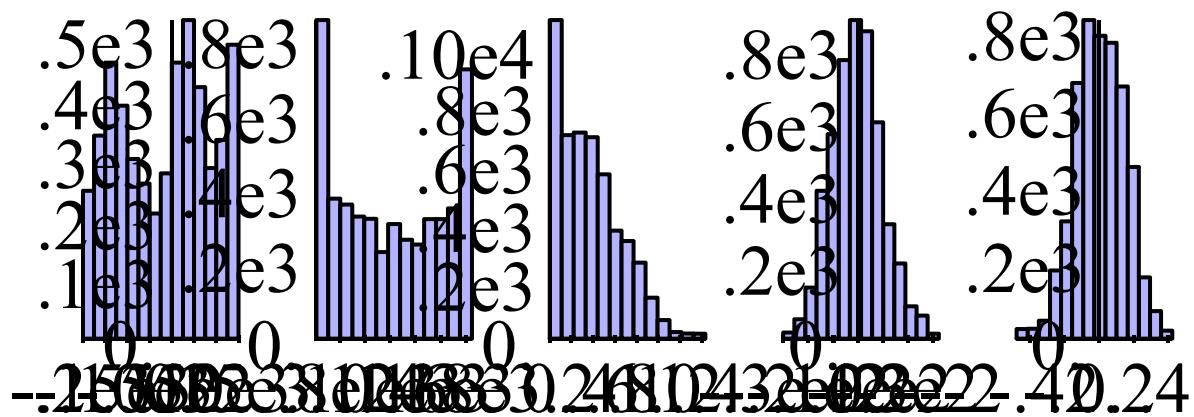
236.259  
205.586  
174.914  
144.242  
113.570  
82.898  
52.226  
21.553  
-9.119  
-39.791  
-70.463  
-101.135  
-131.808  
-162.480  
-193.152  
-223.824  
.28508262699949313936880477559752678310672668195186  
0.2303730288458049610945051243259460875369096232446e-1  
0.2777429570391344058664824464676343306965902568925e-1  
.19375068852324638506384597686093225738562708905404  
-.17615961865742067045895629092489891829840484758118  
-0.7310025838087725981758558710730093982436784216386e-2  
.10061776698124521849543917350343873033353127914840  
0.18447242866332575236084910338506555980495559520997e-1  
0.82089235685665519713282642552675380296463622885786e-1  
.19749682850499846419048037476684420461243168625058  
.60278372132433140866767810698101302892839974961536  
.48910351414366435314487583919518185324436781298015  
-.23908100997124829011447842396975032110866790664725  
-0.131017151915345637280691755581496792699843282464e-3  
-0.19822824332582401160082959541412672476731779917675e-1  
.35142216848675054331711477267275615183923628344711

```
> display(graph3a,{op(graph3b),graph3c,plot(0, x=XFinal[1]+
BinLengthScaled/2..XFinal[NumBins]-BinLengthScaled/2,color=black)}, axes=
boxed, title=title3, titlefont=[Times,14,bold], labels=["LSR Velocity
(km/s)","Main Beam Temperature (K)"], labelfont=[Arial,11,bold], font=[Arial,
11,bold], labeldirections=[horizontal,vertical]);
```

### [NII] 205 micron emission in Car I SPIFI Raster Position (6,7), LBC



```
> display(array([Cpdf,Fpdf,Hpdf,Spdf,Opdf]));
```



```
> display(graph1a,{op(graph1b)});
```

