# **Interactive Tutorial**

#### **ALMA Guides: a first look at imaging and at spectral line imaging**



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### **Goals of this tutorial**

- Learn how to create continuum and line cubes in tclean
- Understand some key parameters in tclean



# **First look at imaging**



## **From Sky Brightness to Visibility**

- 1. An interferometer measures the interference pattern observed by pairs of apertures
- 2. The interference pattern is directly related to the source brightness. In particular, for small fields of view, the complex visibility, V(u,v), is the 2D Fourier transform of the brightness on the sky, T(x,y) *image* plane

 $X \sim \sim \sim \sim \sim \sim$ 

*uv* plane

 $\boldsymbol{\mathit{u}}$ 

y

N Pole

Fourier space/domain

$$
V(u, v) = \int \int T(x, y)e^{2\pi i (ux + vy)} dx dy
$$

Image space/domain

$$
T(x,y) = \int \int V(u,v)e^{-2\pi i(ux+vy)}dudv
$$

Slide courtesy of Amanda Kepley

### **Interferometers discretely sample the uvplane.**



- Small uv-distance: **short baselines** (measures extended emission)
- Long uv-distance: **long baselines** (measures small scale emission)
- Orientation of baseline determines orientation in the uv-plane
- **Antennas can only physically be so close together leaving a**



**hole in the center of the uv-plane (missing short spacings)**

Slide courtesy of Amanda Kepley

### **Missing Short Spacings: Demonstration**



## **Gaps in uv coverage: why we clean**





"Dirty" image vs. "Clean" image of TW Hya of TW Hya

### **Clean is the most common deconvolution algorithm.**

#### Sky Model : List of delta-functions

(1) Construct the observed (dirty) image and PSF

(2) Search for the location of peak amplitude.

(3) Add a delta-function of this peak/location to the model

(4) Subtract the contribution of this component from the dirty image - a scaled/shifted copy of the PSF

Repeat steps (2), (3), (4) until a stopping criterion is reached.

(5) Restore : Smooth the model with a 'clean beam' and add residuals



#### **Choices:** what and how much PSF to subtract and when to stop



Slide courtesy of Amanda Kepley, adapted from slide by Urvashi Rau

# **Time for Hands on Tutorial**

### **First look at continuum imaging**

Following ALMA CASA Guide "First Look at Imaging" : Continuum

• [https://casaguides.nrao.edu/index.php?](https://casaguides.nrao.edu/index.php?title=First_Look_at_Imaging_CASA_6.4) title=First Look at Imaging CASA 6.4







### **Cd to data directory and start-up CASA**

\$casa



**Slides and commands (imaging\_tutorial\_commands.txt) are in [https://astrocloud.nrao.edu/s/](http://www.apple.com) [T7wzjL2tTP6Leg4](http://www.apple.com)**



### **Using CASA tasks and getting oriented with the data**

Bring up a list of the input parameters and enter them one by one:

inp listobs vis='sis14\_twhya\_calibrated\_flagged.ms' go

(commands to input to CASA are highlighted in yellow)



### **Using CASA tasks and getting oriented with the data**

Bring up a list of the input parameters and enter them one by one:

inp listobs vis='sis14\_twhya\_calibrated\_flagged.ms' go

OR All in one command (preferred!): listobs(vis='sis14\_twhya\_calibrated\_flagged.ms')



## **Using CASA tasks and getting oriented with the data**

Bring up a list of the input parameters and enter them one by one:

inp listobs

vis='sis14\_twhya\_calibrated\_flagged.ms' listfile='twhya\_listobs.txt' go OR All in one command: listobs(vis='sis14\_twhya\_calibrated\_flagged.ms',listfile='twhya\_listobs.txt') Print to file!



## **Inspecting your data:**

#### Check the uv coverage:

plotms(vis='sis14\_twhya\_calibrated\_flagged. ms', xaxis='u', yaxis='v', avgchannel='10000', avgspw=False, avgtime='1e9', avgscan=False, coloraxis='field', 300 showgui=True)





### **Inspecting your data:**

Can use plotms to check your baseline coverage: plotms(vis='sis14\_twhya\_calibrated\_flagged.ms', xaxis='UVwave', yaxis='Amp', avgchannel='10000', avgspw=False, avgtime='1e9', avgscan=False,field='2', coloraxis='antenna1', showgui=True)





We'll use phase cal as the image name for our phase calibrator image.

First, remove any old files: os.system('rm –rf phase\_cal.\*') OR rm –rf phase\_cal.\*

(not all unix commands work in CASA without a ! or os.system, but rm does)



#### Choosing tclean parameters:

tclean(vis='sis14\_twhya\_calibrated\_flagged.ms', imagename='phase\_cal\_dirty', field='3',

 $spw=$ ",

```
specmode='mfs',
```

```
deconvolver='hogbom',
```

```
gridder='standard',
```

```
imsize=[250,250],
```
cell=['0.1arcsec'],

weighting='briggs',

threshold='0.0mJy',

interactive=True)

Look in the listobs output for the intent "CALIBRATE\_PHASE#ON\_SOURCE" to get the field number for our phase calibrator.



#### Choosing tclean parameters:

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_dirty', field='3',
```
spw='',

#### Leave blank to use all spws

```
specmode='mfs',
```

```
deconvolver='hogbom',
```

```
gridder='standard',
```

```
imsize=[250,250],
```

```
cell=['0.1arcsec'],
```

```
weighting='briggs',
```

```
threshold='0.0mJy',
```

```
interactive=True)
```


#### Choosing tclean parameters:

tclean(vis='sis14\_twhya\_calibrated\_flagged.ms', imagename='phase\_cal\_dirty', field='3',

 $spw=$ ",

specmode='mfs',

deconvolver='hogbom', gridder='standard', imsize=[250,250], cell=['0.1arcsec'], weighting='briggs', threshold='0.0mJy', interactive=True)

For a continuum image, choose specmode = mfs (multifrequency synthesis)



Choosing tclean parameters: tclean(vis='sis14\_twhya\_calibrated\_flagged.ms', imagename='phase\_cal\_dirty', field='3', spw='', specmode='mfs',  $\Delta$ deconvolver='hogbom', Since - < 10%, choose hogbom gridder='standard',  $\mathcal V$ imsize=[250,250], cell=['0.1arcsec'], weighting='briggs', threshold='0.0mJy', interactive=True)



#### Choosing tclean parameters:

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_dirty', field='3',
```
 $spw=$ ",

```
specmode='mfs',
```
deconvolver='hogbom',

gridder='standard',

imsize=[250,250], cell=['0.1arcsec'], weighting='briggs', threshold='0.0mJy', interactive=True)

We're only imaging a single pointing, with 12m only, so use the standard gridder.



#### Choosing tclean parameters:

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_dirty', field='3',
```
 $spw=$ ",

```
specmode='mfs',
```
deconvolver='hogbom',

gridder='standard',

```
imsize=[250,250],
```
cell=['0.1arcsec'],

weighting='briggs', threshold='0.0mJy', interactive=True)

Want ~5 cells across minor axis of synthesized beam

Beam in arcseconds ~ 206265/(longest baseline in wavelengths)

 $Bean \sim 206265/429300 = 0.48"$ 



#### Choosing tclean parameters:

tclean(vis='sis14\_twhya\_calibrated\_flagged.ms', imagename='phase\_cal\_dirty', field='3',

 $spw=$ ",

```
specmode='mfs',
```
deconvolver='hogbom',

gridder='standard',

imsize=[250,250],

cell=['0.1arcsec'], weighting='briggs', threshold='0.0mJy', interactive=True)

Image size  $=$  # of cells needed to roughly cover primary beam (larger if non-point source)

For single fields, primary beam in arcsec is:

- $\sim 6300$  / nu[GHz] for 12m,
- $\sim$  10608 / nu[GHz] for 7m, where nu[GHz] is the sky frequency expressed in GHz.  $24$

#### Choosing tclean parameters:

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_dirty', field='3',
```
 $spw=$ ",

```
specmode='mfs',
```
deconvolver='hogbom',

gridder='standard',

imsize=[250,250],

cell=['0.1arcsec'], weighting='briggs', threshold='0.0mJy', interactive=True)

tclean will kindly tell you if your image size is inefficient, and make suggestions in the logger window!



```
Choosing tclean parameters:
```

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_dirty', field='3',
```
 $spw=$ ",

```
specmode='mfs',
```

```
deconvolver='hogbom',
```
gridder='standard',

imsize=[250,250],

cell=['0.1arcsec'], weighting='briggs', threshold='0.0mJy', interactive=True)

Weighting choice will depend on science goals.



#### Ways to get beam size and thus cell size:

• Look at alma-technical-handbook





Table 7.1: Angular resolution ( $\theta_{res}$ ) and maximum recoverable scale ( $\theta_{MRS}$ ) values for different 7-m Array and 12-m Array configurations are shown for one representative frequency in each ALMA receiver band. The value of  $\theta_{MRS}$  is computed using the 5<sup>th</sup> percentile baseline (L05) from Table 7.2 and Equation 7.6. The value of  $\theta_{res}$  is the mean size of the interferometric beam obtained through simulations with CASA. Computations were done for a source at zenith; for sources transiting at lower elevations, the North-South angular measures will increase proportional to 1/sin(elevation). Calculations are based on the notional C43-X and 7m configurations with Briggs weighting and a robust parameter of 0.5.





Ways to get beam size and thus cell size:

- Look at alma-technical-handbook
- Make dirty image (tclean with niter = 0 or 1) log tells you beam size
- Use Analysis Utils [\(https://safe.nrao.edu/wiki/bin/view/Main/](https://safe.nrao.edu/wiki/bin/view/Main/CasaExtensions) [CasaExtensions\)](https://safe.nrao.edu/wiki/bin/view/Main/CasaExtensions) 
	- pickCellSize



### **For a video overview of these concepts:**





#### <https://youtu.be/OC3IWpRRtEQ> <https://youtu.be/EVY7000zAD4>

- More videos available and under development!
- **Like and subscribe** to our Youtube channel **ALMA Primer** to get notified when new videos are uploaded.



#### Choosing tclean parameters:

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_dirty', field='3',
```
 $spw=$ ",

```
specmode='mfs',
```

```
deconvolver='hogbom',
```

```
gridder='standard',
```

```
imsize=[250,250],
```

```
cell=['0.1arcsec'],
```
weighting='briggs',

```
threshold='0.0mJy',
```
interactive=True)

The threshold is one main stopping criterion for tclean. It will stop when the residuals fall below the threshold.

Setting the number of iterations, niter, is the other stopping criterion.

By default, niter=0, so running this command will do no cleaning. Try it!



#### Run Tclean command:

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_dirty', field='3', 
spw='', 
specmode='mfs', 
deconvolver='hogbom', 
gridder='standard', 
imsize=[250,250], 
cell=['0.1arcsec'], 
weighting='briggs', 
threshold='0.0mJy', 
interactive=True)
```


#### Check the dirty image: imview('phase\_cal\_dirty.image')





Now increase niter to some large number by adding it to the previous command:

```
tclean(vis='sis14_twhya_calibrated_flagged.ms', 
imagename='phase_cal_clean', field='3', 
spw='', 
specmode='mfs', 
deconvolver='hogbom', 
gridder='standard', 
imsize=[128,128], 
cell=['0.1arcsec'], 
weighting='briggs', 
threshold='0.0mJy', 
niter=5000,
```
interactive=True)

#### Creating a mask: select add, all channels



#### Click the oval with the R, to be able to draw an oval





# **First look at Imaging: the phase**

### **calibrator**

Draw an oval around the source, and then double click to set the mask.



#### Then click the green circular arrow to run

the first cycle.





Now we see the noise looks a little brighter, relative to the source since we've removed some signal.



#### Click the green circular arrow again to run another cycle.





And we see the noise looks even brighter, relative to the source. When the residuals for the source match the noise, stop!



#### Click the red stop sign to end the cleaning.





### **First look at Imaging: the science target**

First split out only the science target data (field=5) and average over 8 channels (width='8') to reduce the size of the data set.

os.system('rm -rf twhya\_smoothed.ms')

split(vis='sis14\_twhya\_calibrated\_flagged.ms', field='5', width='8', outputvis='twhya\_smoothed.ms', datacolumn='data')

Try listobs on the new measurement set to verify. Note the field number has changed!



## **First look at Imaging: the science target**

Now clean!

```
os.system('rm -rf twhya_cont.*')
```

```
tclean(vis='twhya_smoothed.ms',
imagename='twhya_cont',
field='0', spw='', specmode='mfs',
Note the field is now '0'!.
gridder='standard',
deconvolver='hogbom',
imsize=[250,250],
cell=['0.1arcsec'],
weighting='briggs',
robust=0.5, threshold='0mJy',
niter=5000,
interactive=True)
```


### After ~68 iterations: we can see fainter parts of the disk **First look at Imaging: the science target**

now, so it looks bigger.





What if I want to adjust my mask? Change to erase, select an area surrounding the old mask, and double click to erase any mask inside.

### After 168 iterations: The brightest noise is the same color as the brightest source, so stop! **First look at Imaging: the science target**





### **First look at Imaging: the science target**

Use imview to inspect the tclean results:

#### The Clean Image





### **First look at Imaging: the science target**

Use imview to inspect the tclean results:

The PSF





### Hmm… but what about that blue arrow? **First look at Imaging: the science target**



It auto-completes an interactive session!

Here you can also adjust iterations left and the threshold before auto-completing.



## **First look at Imaging: the science target**

The last step for the continuum image: primary beam correction.

os.system('rm -rf twhya\_cont.pbcor.image')

impbcor(imagename='twhya\_cont.image', pbimage='twhya\_cont.pb', outfile='twhya\_cont.pbcor.image')

And inspect your image: imview('twhya\_cont.pbcor.image')



# **First look at spectral line imaging**

- Following ALMA CASA Guide "First Look at Line Imaging"
- [https://casaguides.nrao.edu/index.php?](https://casaguides.nrao.edu/index.php?title=First_Look_at_Line_Imaging_CASA_6.4) title=First Look at Line Imaging CASA 6.4







## **First look at Spectral Line Imaging: Removing the continuum**

Need to remove any channels with notable emission before fitting continuum:



## **First look at Spectral Line Imaging: Removing the continuum**

Fit and subtract the continuum:

```
os.system('rm -rf sis14_twhya_selfcal.ms.contsub')
uvcontsub(vis = 'sis14_twhya_selfcal.ms',
      field = '5',
       fitspw = '0:0~239;281~383',
       excludechans = False,
      fitorder = 0,
       solint='int')
```
For large datasets this might take quite some time!



## **First look at Spectral Line Imaging: Removing the continuum**

Use plotms to confirm:



### **First look at Spectral Line Imaging: Specmode=cube now**

restfreq = '372.67249GHz'

os.system('rm -rf twhya\_n2hp.\*')  $tclean(vis = 'sis14_twhya_selfcal.ms.contsub', imagename =$ 'twhya\_n2hp',field =  $'0$ ',spw =  $'0'$ ,specmode = 'cube',perchanweightdensity=True, nchan =  $15$ , start =  $'0.0$ km/s', width =  $'0.5$ km/s',outframe = 'LSRK', restfreq = restfreq,deconvolver= 'hogbom',gridder = 'standard',imsize = [250, 250],cell = '0.1arcsec',phasecenter = 0,weighting = 'briggsbwtaper',robust = 0.5,restoringbeam='common',interactive = True,niter=5000)

(note: no threshold set, so defaults to 0.0)



## **First look at Spectral Line Imaging: Masking options**

Can do channel by channel mask, or select "all channels" and do single mask for all channels: here ch 5,6,7.



Dirty images with masks.



### **First look at Spectral Line Imaging: specmode = cube**

Residuals for channels 5,6,7 after 2 major cycles:





## **First look at Spectral Line Imaging: specmode = cube**

Channel 6: Dirty vs Clean







### **First look at Spectral Line Imaging: spectral profile tool**





### **First look at Spectral Line Imaging: primary beam correction**

os.system('rm -rf twhya\_n2hp.pbcor.image')

impbcor(imagename='twhya\_n2hp.image', pbimage='twhya\_n2hp.pb', outfile='twhya\_n2hp.pbcor.image')



### **Image Analysis: Moment Maps**

### https://casaguides.nrao.edu/index.php? title=First Look at Image Analysis CASA 6.4

immoments(imagename, moments=[0], axis='spectral', region=", box=", chans=", stokes=", mask=", includepix=- 1, excludepix=- 1, outfile=", stretch=False)

- moments =  $-1$  mean value of the spectrum
- moments =  $0$  integrated value of the spectrum
- moments = 1 intensity weighted coordinate; traditionally used to get "velocity fields"
- moments = 2 intensity weighted dispersion of the coordinate; traditionally used to get "velocity dispersion"
- moments =  $3$  median value of the spectrum
- moments =  $4$  median coordinate
- moments = 5 standard deviation about the mean of the spectrum
- moments =  $6$  root mean square of the spectrum
- moments  $= 7$  absolute mean deviation of the spectrum
- moments =  $8 -$  maximum value of the spectrum
- moments =  $9$  coordinate of the maximum value of the spectrum
- moments =  $10 minimum$  value of the spectrum
- moments = 11 coordinate of the minimum value of the spectrum

