Data Processing: Imaging

If multiple measurement sets are available for a target, it is necessary to concatenate the data before imaging the data. (This is not the case for the example.)

Concatenation can be performed with the following command:

concat(vis=['uid1.ms,uid2.ms'],concatvis='calibrated.ms')

Any old calibrated.ms files should be deleted before running this command.

Interferometry data are converted into images using a process called cleaning. This serves two purposes:

 It converts the visibility (phase and amplitude data) into signal on the sky plane.

• It restructures the beam (point spread function) of the data so that the beam is Gaussian in shape.

Cleaning is done using either the old clean command or the new tclean command. The example tclean command for the continuum imaging looks like the following:

tclean(

vis='uid____A002_xb4264b_x946.ms.split.cal', #Inpu imagename='NGC3169.continuum', #Outp field='2', #Fiel gridder='standard', #Proj phasecenter=2, #Cent imsize=[500, 500], #Imag cell='0.1arcsec', #Pixe pblimit=0.0, #PB g

spw='0,1,2,3:0~1200;2500~3839'
specmode='mfs',
outframe='lsrk',
deconvolver='hogbom',
nterms=1,

chanchunks=-1, channels niter=100, threshold='0.1mJy', weighting='natural' interactive=F) **#Input** filename **#Output filename** #Fields to be imaged **#Projection method** #Centre position of image #Image size in pixels #Pixel scale pbcor=True, **#PB** gain level at which to cut # off normalization #Spectral windows to image #Imaging mode (continuum) #velocity frame of image #Cleaning algorithm #Number of Taylor coefficients # in spectral slope #Chunking for gridding

#Maximum number of iterations
#Stopping threshold
#Cleaning weights
#Interactive mode setting

The various options for tclean in this example have been reorganized into rough groups:

- Input/output information
- Image plane settings
- Spectral dimension settings
- Deconvolution options

Input/Output Information

vis imagename Input filename Output filename

Image plane settings

field gridder phasecenter imsize cell pbcor pblimit Fields to be imaged Projection method (tclean only) Centre position of image Image size in pixels Pixel scale Setting to apply PB correction PB gain level at which to cut off normalization (tclean only)

Important points on image plane settings

- The gridder should be set to "standard" for single pointings and "mosaic" for multiple pointings.
- The phasecenter can be specified either by a field number or by coordinates (which may be necessary when working with multiple pointings).
- The cell value should be at least 2× (and preferable 3-4×) the size of the beam.
- It is recommended to apply the primary beam (PB) correction.

Spectral dimension settings

Spw
specmode
start
nchan
width
outframe
restfreq

Spectral windows to image Spectral imaging mode (mode in clean) Staring point of spectral cube Number of spectral cube channels Width of spectral cube channels Velocity frame Rest frequency of reference line (optional)

Important points on spectral dimension settings

 The specmode should be set to "mfs" for continuum images and "cube" for spectral cube. The older clean command had multiple options instead of "cube", such as "frequency", "velocity", and "channel".

- The start, nchan, and width options should be set for spectral cubes but not for continuum images.
- The outframe should be set just to avoid confusion. Currently, tclean only supports "lsrk", but for clean, "lsrk" is recommended for galactic objects and "bary" for extragalactic objects.

Important points on spectral dimension settings

 The restfreq is optional. The CASA software will be able to perform some additional meaningful unit conversions between frequency and velocity if this is given.

Deconvolution options

deconvolver

nterms

chanchunks

niter threshold weighting interactive Cleaning algorithm (imagermode in clean) Number of Taylor coefficients in spectral slope (tclean only) Chunking for gridding channels (tclean only) Maximum number of iterations Stopping threshold **Cleaning** weights Interactive mode setting

Important points on deconvolution options

- The weighting is very important. Three standard options are used in radio interferometry.
 - Natural weighting is based on not altering the weights of data points in the uv plane. This results in images with more large-scale structure
 - Uniform weighting is based on altering the weights to account for the lack of data on long baselines in the uv plane (thus making the uv plane appear "uniform"). This results in images with more small-scale structure, but ALMA image with uniform weights tend to look too noisy.
 - Briggs weighting allows for adjusting between these two extremes. The robust parameter can be used to adjust between these extremes, with "2" equivalent to natural and "-2" equivalent to uniform. A robust value of "0.5" is used in most ALMA QA2 imaging.

Cleaning is an iterative process in which the following is done in each iteration:

- An image is displayed.
- Either the user or the program identifies sources and masks them.
- Using the identified sources, the algorithm models and removes them from the image, producing a residual image that is used at the beginning of the cycle.











Multiple files are created after cleaning:

- image
- image.pbcor
- mask
- model
- pb
- psf
- residual

Science image Science image with a PB correction Image mask created while cleaning Model of the emission used for deconvolution Primary beam (area where the telescope was sensitive) Beam before cleaning Residuals after the final iteration of cleaning



















Imaging tips

- Try using different weights (natural or briggs with different robust values) to bring out emission on different spatial scales.
- Attempt changing the channel width to improve the S/N of line emission.
- To identify artefacts (such as ripples across the map), try producing images using only subsets of the data.
- To check the reliability of the flux densities, try one of the following:
 - Produce images using subsets of the data.
 - Image the calibration sources and check the flux densities using aU.planetFlux or aU.getALMAflux.