

The University of Manchester

Introduction to Imaging: General Principles

Adam Avison





This presentation has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 730562 [RadioNet]

Further Reading

The slides from this talk are based on the fundamentals of interferometry which are explained in detail across:

- *"Interferometry and Synthesis in Radio Astronomy" -* Thompson, Moran & Swenson
- *"Synthesis Imaging in Radio Astronomy II"* NRAO
- *"An introduction to Radio Astronomy"* Burke and Graham-Smith (4th edition out soon Burke, Graham-Smith (& Wilkinson?))
- "Tools of Radio Astronomy" Wilson, Rohfls & Hüttemeister
- *"The CASA Cookbook"* Ott & Kern et al.

Outline

- What we measure
- Imaging the data
- The CLEANing process

An advanced warning, for a talk about imaging it is rather wordy to begin with.

What we measure

Correlator output

$$R_{x,y} = A_0 |V| \Delta v \cos(2\pi v \mathbf{b}_{\lambda} \cdot \mathbf{s}_0 - \phi_V) \qquad \text{Eq. 1}$$

Visibility equation

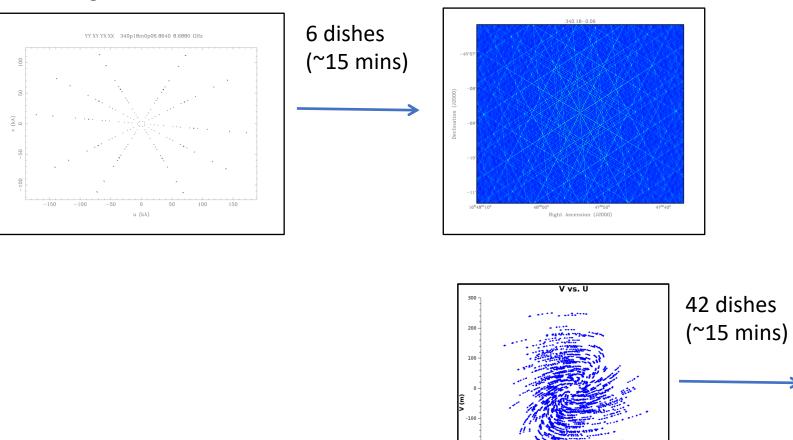
$$V(u,v) = \int A(l,m)I(l,m) e^{-i2\pi(ul+vm)} \frac{dldm}{\sqrt{1-l^2-m^2}}$$
 Eq. 2

Dirty image

$$I^{D}(l,m) = \iint_{-\infty}^{\infty} S(u,v)V(u,v)e^{2\pi i(ul+vm)}dudv$$
 Eq. 3

Filling the *uv*-plane

We want to fill the *uv*-plane because the *uv*-coverage is the **FT** of the synthesised beam, *B*. The greater the *uv*-coverage the better behaved the sidelobes are.

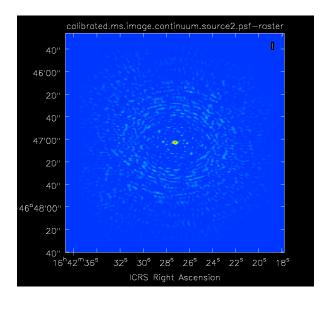


-200

-300

50 100 150

11 (m)



Sampling function

Let us call the sampling of the *uv*-plane (aka *uv*-coverage), *S*, the sampling function

$$S(u,v) = \sum_{k=1}^{M} \delta(u - u_k, v - v_k)$$

Given this the synthesized beam, B, is B = FT(S).

And for each uv-point we have an observed visibility, V(u,v), so we can define the sampled visibility function as:

$$V^{S}(u,v) \equiv \sum_{k=1}^{M} \delta(u-u_{k},v-v_{k}) V(u_{k},v_{k})$$

So $V^{s}=SV$ and from earlier (eq. 3) $I^{D}=FT(V^{s})=FT(SV)$. From which it follows that I^{D} is the measured sky brightness convolved with the synthesis beam, *B*.

This is a truncated version of Chapter 7 (Briggs, Schwab & Sramek) of "Synthesis Imaging in Radio Astronomy II"

Weighting

Given this we can introduce weighting functions to control the shape of the synthesised beam.

$$W(u,v) = \sum_{k=1}^{M} R_k D_k T_k \delta(u - u_k, v - v_k)$$

 R_k = Weights relating to data quality, i.e. down weight bad data. This is observation dependent and we have no post observation control over it (so ignore).

 T_k = Tapering function. Apply a tapering function (i.e. Gaussian), to the *uv*-coverage to for example downweight the outer *uv*-points lowering resolution.

 D_k = Density weighting... Next slide.

And as per the previous slide we can define the weighted and sampled visibility function as $V^{W} = WV$.

This is a truncated version of Chapter 7 (Briggs, Schwab & Sramek) of "Synthesis Imaging in Radio Astronomy II"

Density weighting description

Due to the nature of how arrays are typically built it the *uv*-coverage density is typically higher toward the *uv* origin.

Two "extremes" of density weighting are typically used:

Natural weighting: D_k = 1. All visibilities are treated the same.

- This gives the highest signal to noise possible within the final image
- The poorest angular resolution and higher sidelobe effects.

Uniform weighting: $D_k = 1/N_s(k)$. Weight visibilities by the number of data points in a symmetric region, s.

- Downweights data in dense regions.
- Higher resolution, lower sidelobes, but worse SNR.

There are then super and sub-uniform (see e.g. CASA cookbook), and...

Briggs weighting: A 'sliding scale' between Natural and Uniform controlled by the 'robust' parameter. With (in CASA) +2 = nearly Natural and -2 = nearly Uniform.

We'll come back to these later as we define these during CLEAN.

This is a truncated version of Chapter 7 (Briggs, Schwab & Sramek) of *"Synthesis Imaging in Radio Astronomy II"*

Imaging the data

We now have the sampled and weighted visibilities, V^{W} .

In order to efficiently make an image of our target sky brightness distribution, *I*, we need to take the Fourier transform of this using Fast Fourier Transforms (FFTs).

This requires the V^{W} data to be gridded on to a regular grid.

This is done by convolution with some suitable gridding function^{*}. Leaving us ultimately with some weighted and gridded visibilities which can be FFT'd to give us our dirty image I^{p} .

CLEAN-ing

We've seen that our dirty image I^{p} is the sky brightness distribution convolved with the synthesised beam B_{r} .

To get a better representation of the sky brightness distribution we need to remove the artefacts introduced by ${\it B}$ To achieve this we use CLEAN*

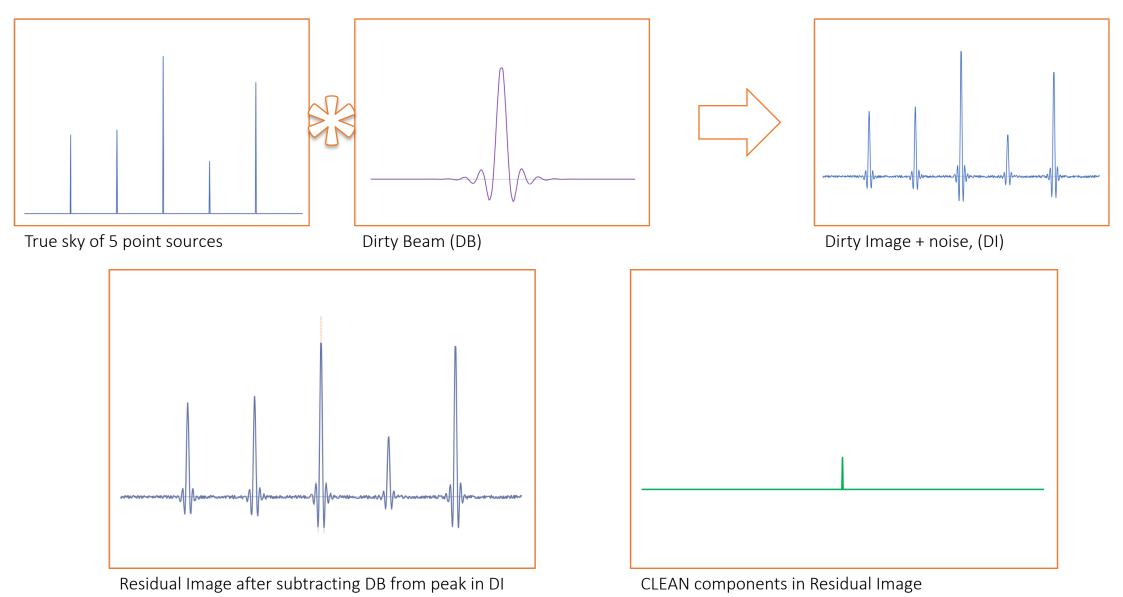
Simple CLEAN overview

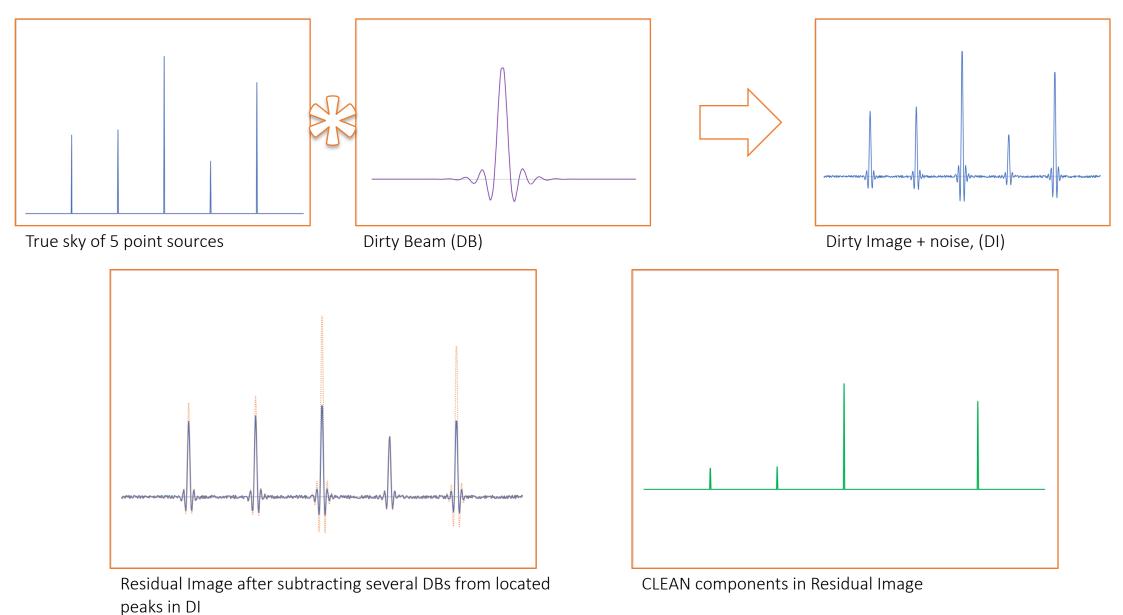
The Högbom (1974) CLEANing algorithm is the simplest CLEAN algorithm and is very illustrative of how CLEAN works in general.

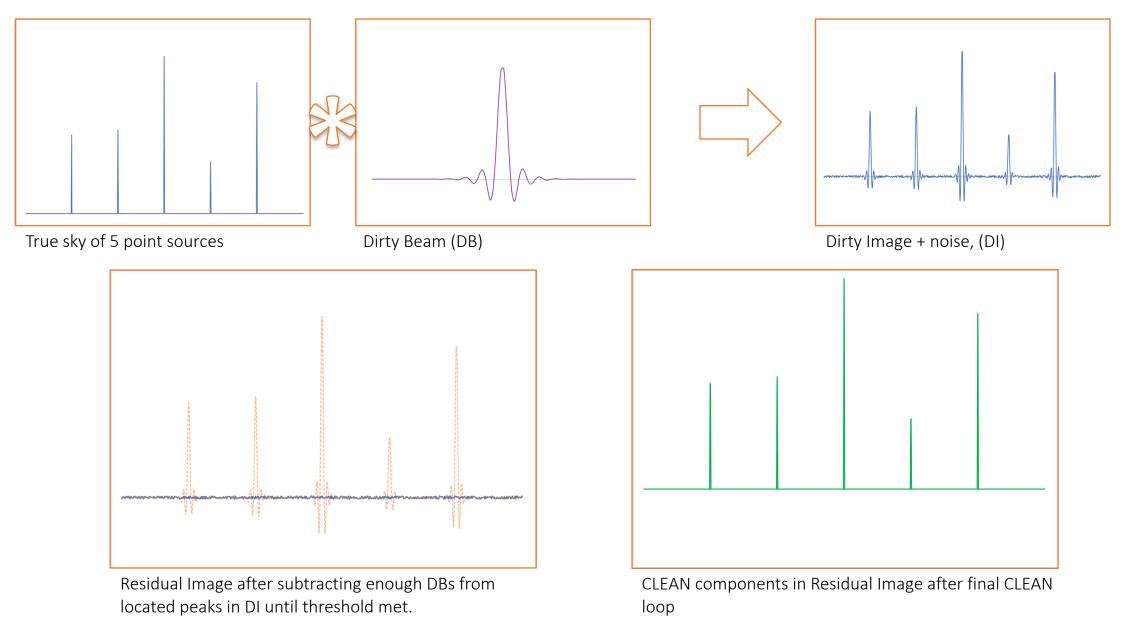
In words the Högbom algorithm works as follows:

- 1) Find the magnitude and position of peak emission in the dirty image.
- 2) Subtract from the dirty image the dirty beam, *B*, scaled by some gain value (i.e. 0.1). Creating a 'residual' image.
- 3) Note the position and magnitude subtracted as a point in a model.
- 4) Repeat 1-3 until a user defined threshold is reached, either some noise limit (in the residual) or a given number of iterations.
- 5) Convolve the final model with an idealised beam. I.e. a beam based on the interferometer if it was a huge single dish.

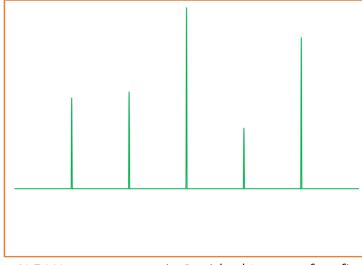
Or in a 2D example on the next page



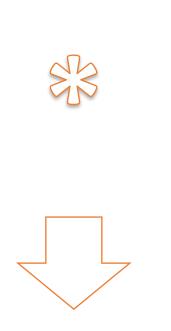


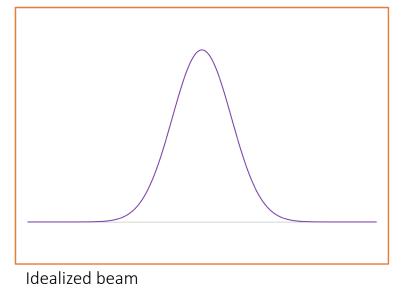


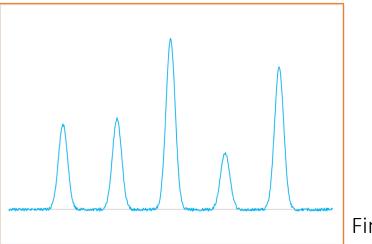
Iteration FINAL



CLEAN components in Residual Image after final CLEAN loop

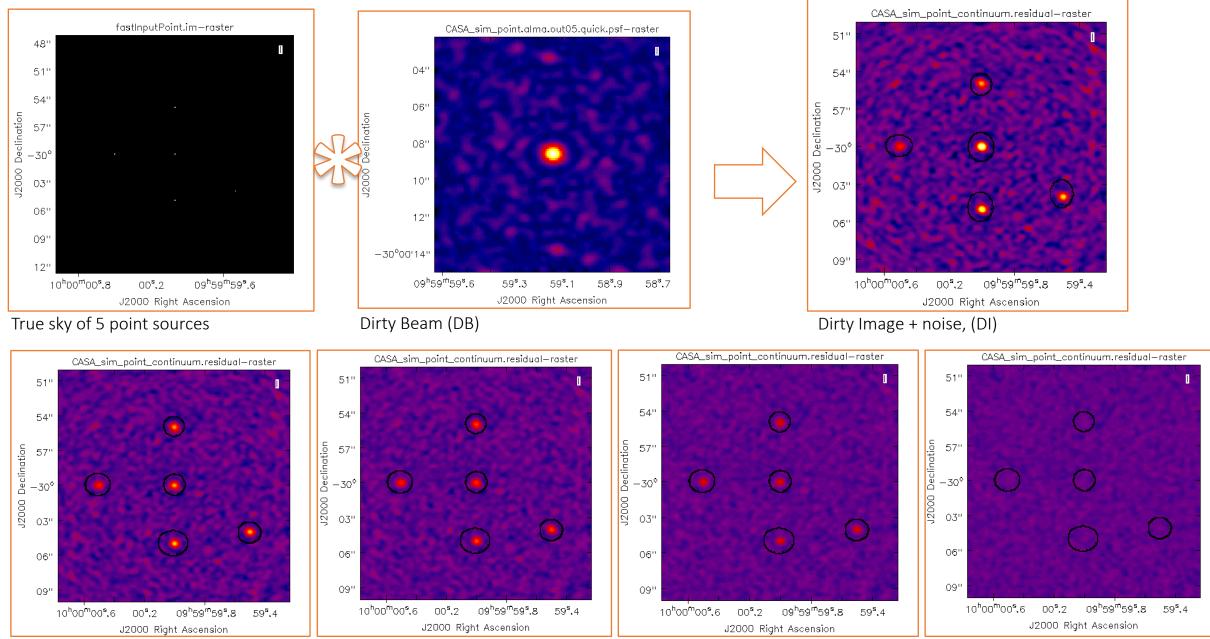






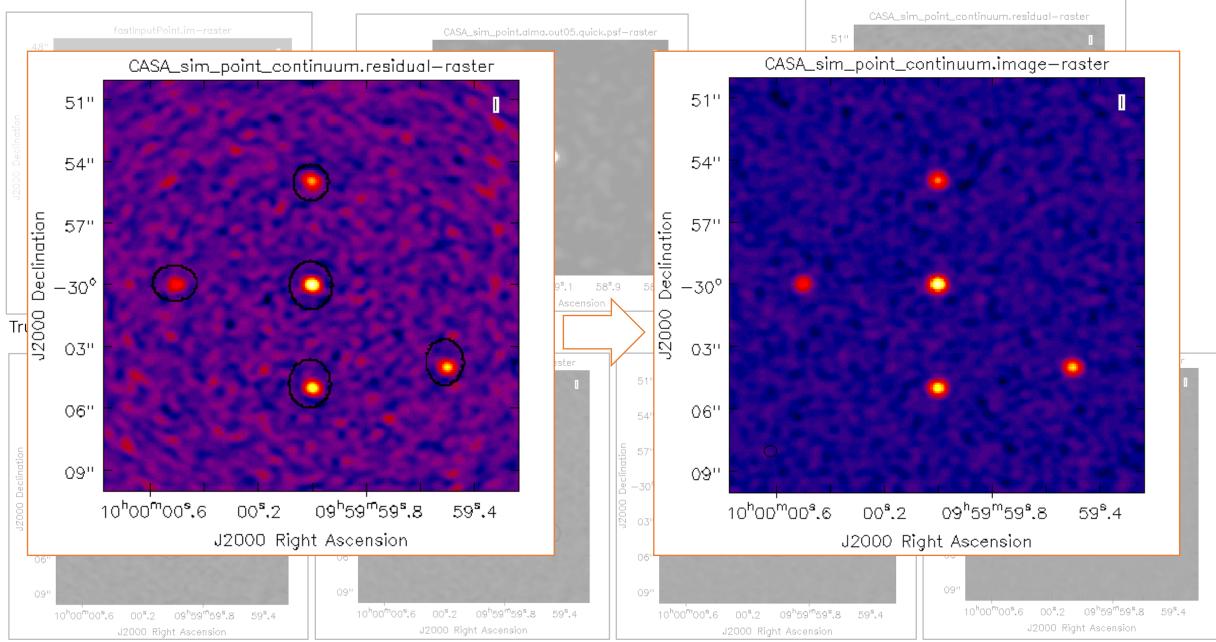
Final reconstructed image

... and in 3D



ightarrow Increasing CLEAN cycles ightarrow

... and in 3D



Other CLEAN algorithm

- Clarke/Cotton-Schwab/ClarkeStokes: Improvements and refinements on Högbom. In general these are preferred to Högbom. They rely on major and minor CLEAN cycles. (Next slide).
- **Multiscale:** Searches for clean components based on user defined size scales. Better for cleaning extended, diffuse, none point-like sources.
- These are only a couple

Major and Minor CLEAN cycles

- For the more 'advanced' CLEAN algorithms (and infact CASA's version of Högbom). CLEAN runs what are termed Major and Minor cycles.
- During a Major Cycle the current model is used to generate model visibilities. Subtract these from the data generate residual visibilities. Convert the residual visibilities into a residual image. (In the visibility domain).
- During a Minor Cycle peak flux components are found and the model and residual image are updated. (All in the image domain).

Basic Sequence of Imaging Logic:

Data : Calibrated visibilities, data weights, UV sampling function Input : Algorithm and iteration controls (stopping threshold, loop gain,...) Output : Model Image, Restored Image, Residual Image,...

```
Initialize the model image

Compute the point spread function

Compute the initial residual image

While ( not reached global stopping criterion ) /* Major Cycle */

{

While ( not reached minor-cycle stopping criterion ) /* Minor Cycle */

{

Find the parameters of a new flux component

Update the model and residual images

}

Use current model image to predict model visibilities

Calculate residual visibilities (data - model)

Compute a new residual image from residual visibilities

}

Convolve the final model image with the fitted beam and add to the residual image
```

				construction
/is	=		#	Name of input visibility file(s)
electdata	=	True	#	Enable data selection parameters
field				field(s) to select
spw				spw(s)/channels to select
timerange				Range of time to select from data
uvrange				Select data within uvrange
antenna				Select data based on
				antenna/baseline
scan observation				Scan number range
intent				Observation ID range Scan Intent(s)
Incenc			#	Scall Incent(S)
latacolumn	= 'co	rrected'	#	Data column to image(data,corrected)
Imagename	=	11		Pre-name of output images
msize		[100]		Number of pixels
ell		arcsec']		Cell size
hasecenter				Phase center of the image
tokes		'I'		Stokes Planes to make
rojection		'SIN'		Coordinate projection (SIN, HPX)
tartmodel				Name of starting model image
pecmode	=	'mfs'		Spectral definition mode
				(mfs,cube,cubedata)
reffreq			#	Reference frequency
ridder	= 'sta	andard'	#	Gridding options (standard,
			#	wproject, widefield, mosaic,
			#	awproject)
vptable				Name of Voltage Pattern table
pblimit		0.2		>PB gain level at which to cut off
			#	normalizations
leconvolver	= '	hogbom'	#	Minor cycle algorithm (hogbom,clark,
and a section		T		multiscale, mtmfs, mem, clarkstokes)
restoration	=	True		Do restoration steps (or not)
restoringbea	am =	0		Restoring beam shape to use. Default
abcar		False		is the PSF main lobe
pbcor		raise	#	Apply PB correction on the output restored image
			#	restored image
outlierfile			#	Name of outlier-field image
			#	definitions
eighting	= 'n;	atural'		
eighting	= 'na	atural'	#	Weighting scheme
	= 'n: =		# #	Weighting scheme (natural,uniform,briggs)
v eighting uvtaper	= 'na =	atural' []	# #	Weighting scheme
	= 'n: =		# # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane
	= 'n: =	[]	# ' # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations
uvtaper	= 'na = =	П	# ' # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution
uvtaper Diter	= 'n; = = =	[]	# # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto-
uvtaper iter semask	= 'n: = =	[] 0 'user'	# # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto-
uvtaper Diter	= 'n; = = =	[]	# # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or
uvtaper iter semask	= 'n = = =	[] 0 'user'	# # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)
uvtaper iter isemask mask	=	[] 0 'user' ''	# # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s))
uvtaper iter semask	= 'n = = =	[] 0 'user'	# # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)
uvtaper isemask mask pbmask	=	[] 0 'user' 0.0	# # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)) primary beam mask
uvtaper iter isemask mask	=	[] 0 'user' ''	# # # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)) primary beam mask True : Re-use existing images. False
uvtaper semask mask pbmask restart	=	[] 'user' '' 0.0 True	# # # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)) primary beam mask True : Re-use existing images. False : Increment imagename
uvtaper isemask mask pbmask	=	[] 0 'user' 0.0	# # # # # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)) primary beam mask True : Re-use existing images. False : Increment imagename Options to save model visibilities
uvtaper semask mask pbmask restart avemodel	=	[] 'user' '' 0.0 True	# # # # # # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)) primary beam mask True : Re-use existing images. False : Increment imagename Options to save model visibilities (none, virtual, modelcolumn)
uvtaper semask mask pbmask restart aavemodel calcres	=	[] 0 'user' '' 0.0 True 'none' True	# # # # # # # # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)) primary beam mask True : Re-use existing images. False : Increment imagename Options to save model visibilities (none, virtual, modelcolumn) Calculate initial residual image
uvtaper semask mask pbmask restart avemodel		[] 'user' '' 0.0 True 'none'	# # # # # # # # # # # # # # #	Weighting scheme (natural,uniform,briggs) uv-taper on outer baselines in uv- plane Maximum number of iterations Type of mask(s) for deconvolution (user, pb, auto-thresh, auto- thresh2, or auto-multithresh) Mask (a list of image name(s) or region file(s) or region string(s)) primary beam mask True : Re-use existing images. False : Increment imagename Options to save model visibilities (none, virtual, modelcolumn)

Setting up CLEAN in CASA

In CASA the task to use for CLEANing is **'tclean'**, which is the current state of the art CLEAN.

It is a refactored replacement to the task 'clean'.

There are a *LOT* of parameters in this task.

At the beginner level the most important to define are:

- 1. The measurement set to use
- 2. The field you want to image
- 3. Which SPW to use
- 4. The size of the image you want to make (next slide)
- 5. The pixel (cell) size you want in your image (next slide)
- 6. The image weighting
- 7. The cleaning threshold value

Rules of thumb:

• Cell size = At least 3 pixels across the synthesised beam. Defined in arcsec.

$$cell \sim \frac{\lambda}{b_{max}} \times \frac{1}{n}$$
 (where $n \sim 3-6$)

• Image size = Cover at least the primary beam/field of view. Defined in pixels of size=*cell*.

$$imsize \sim \frac{\lambda}{D} \times \frac{1}{cell}$$
 (where D is the dish diameter)

• Your threshold should be roughly:

threshold ~
$$\frac{2kT_{sys}}{A_e\sqrt{(N(N-1)\Delta\nu\Delta t)}}$$

That's some theory... Time for the practice!