

Introduction to imaging*

Cristiana Spingola

INAF Istituto di Radioastronomia (Bologna, Italy)



*Strongly based on previous CASA-VLBI workshops and ERIS schools -Thanks to all previous lecturers!

CASA VLBI Worskhop 2023, JIVE June 5-9, 2023

This event has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 101004719.

Intro

The output of an interferometer is basically a table of the correlation (amp. & phase) measured on each baseline every few seconds.

To get the final image out of our visibilities the steps are:

Calibration and data editing (all lectures so far)

Deconvolution = making a CLEANed image and a model of your source (this lecture!)

Refining calibration = self-calibration (next lecture)

This lecture + material

Deconvolution = making a CLEANed image and a model of your source

1) For experimenting with EVN data and for learning how to script CASA commands,

Follow this:

https://www.jb.man.ac.uk/DARA/unit4/Workshops/EVN_continuum.html

by Jack Radcliffe, Anita Richards and Des Small

2) Test data associated to this lecture are from VLBA, to have an alternative dataset

Imaging intro: Fourier Transform imaging and sampling function

S = sampling function

- = 1 where there is a measurement in the uv plane
- = 0 otherwise

B = Intrinsic source brightness distribution

D = dirty beam = point spread function (PSF)

S = Sampling function

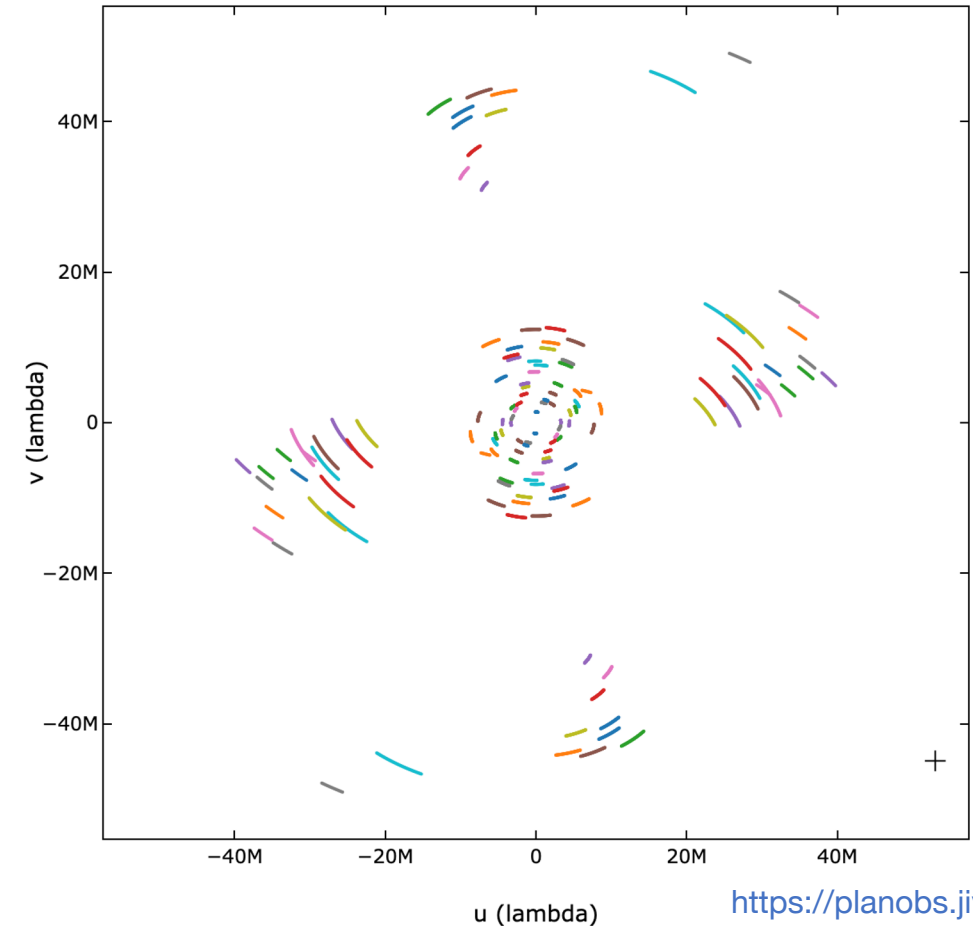
Convolution

Observed visibilities = $B(l, m) * D(l, m) \approx \iint_{uv} S(u, v) V(u, v) e^{2\pi i(ul+vm)} dudv$

$$D(l, m) = \iint_{uv} S(u, v) e^{2\pi i(ul+vm)} dudv$$

Dirty beam D(l,m) = Fourier transform of the sampling function
We know D(l,m) !!!

We need to **deconvolve** B(l,m) from the dirty beam D(l,m)



1 h EVN

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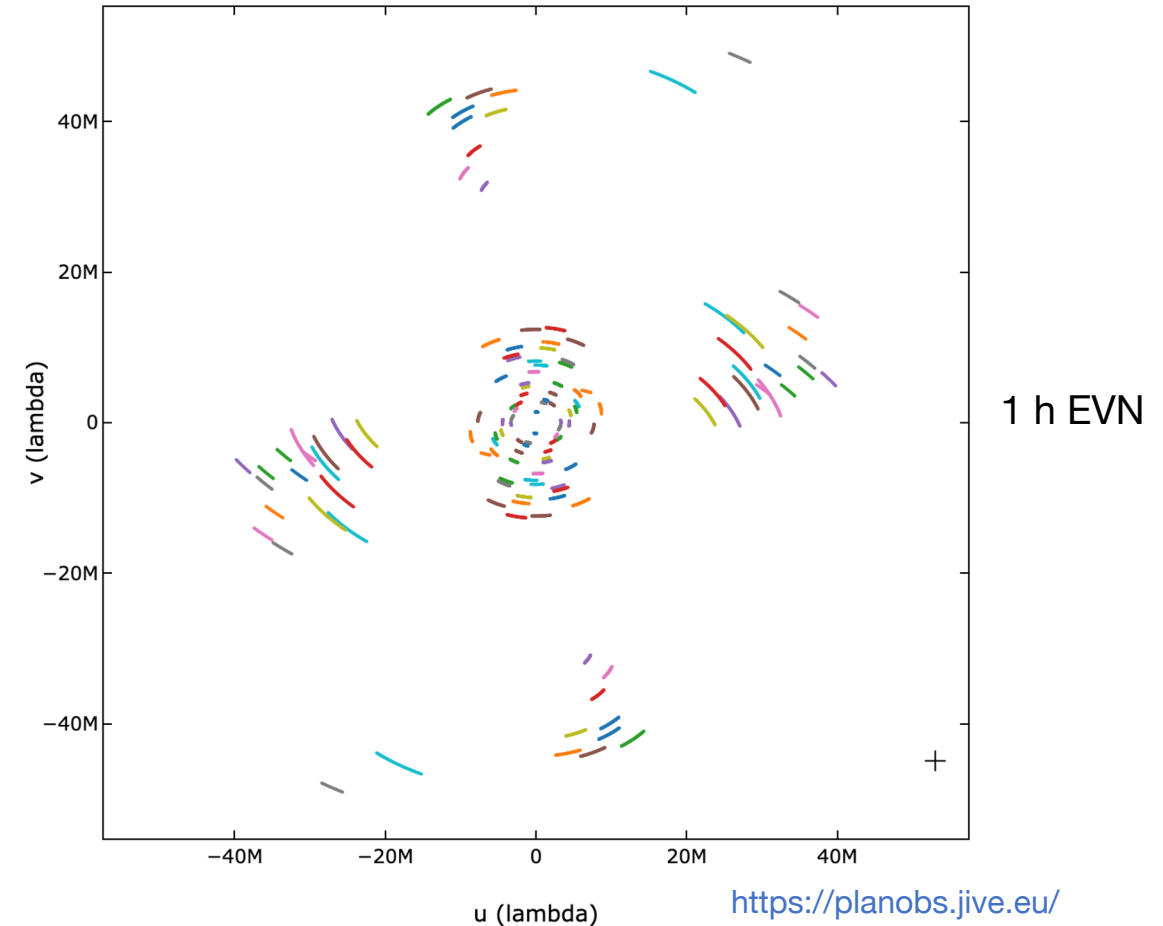
= 1 where there is a measurement in the uv plane

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An ideal interferometer would deliver on a regularly highly sampled rectangular grid. An image would then be made by simply applying a Fourier transform

But, arrays provide (poorly) sampled Fourier Transform of the radio brightness region of sky

You need as many $V(u,v)$ points as possible to reconstruct as robustly as possible the surface brightness distribution of the source



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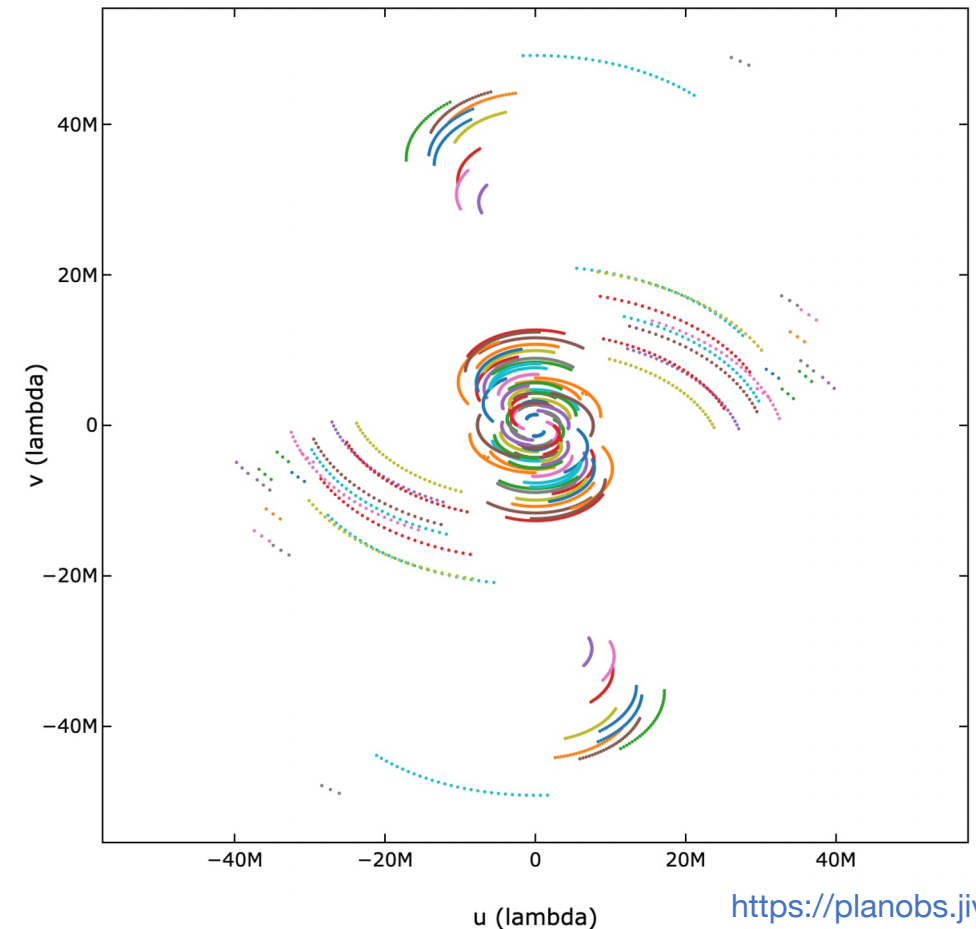
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6 h EVN

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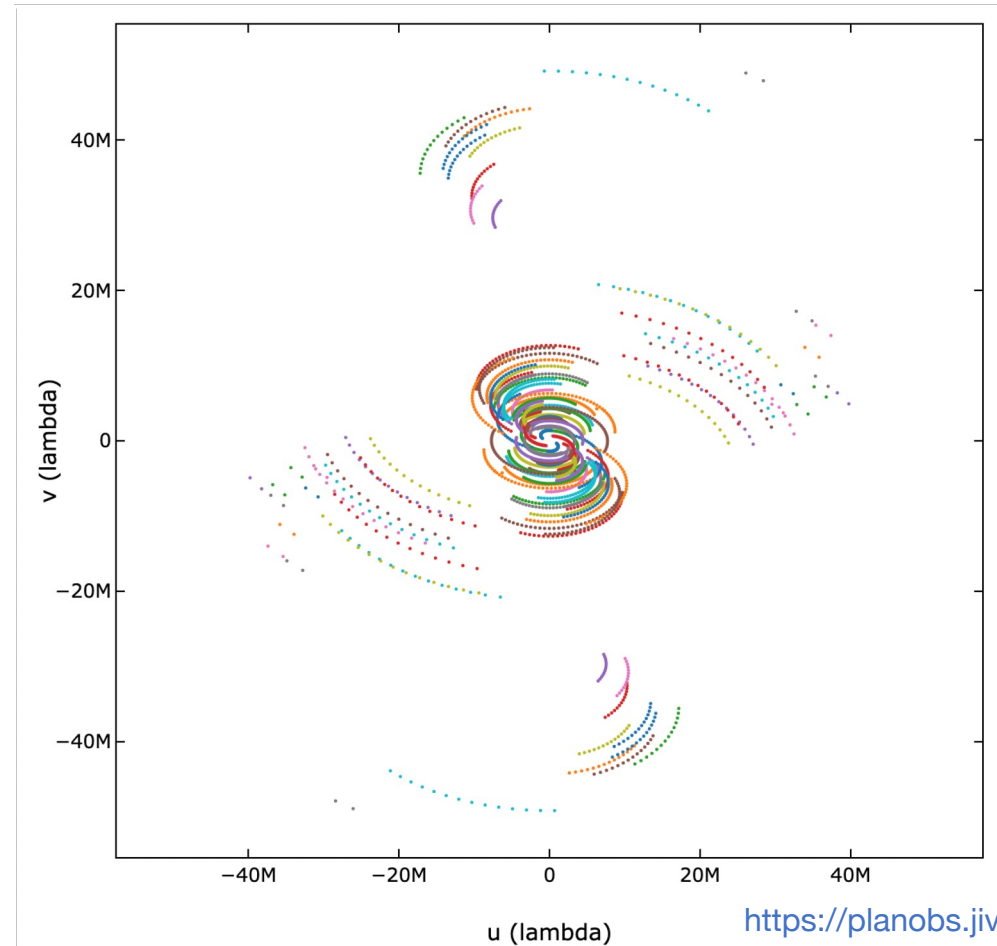
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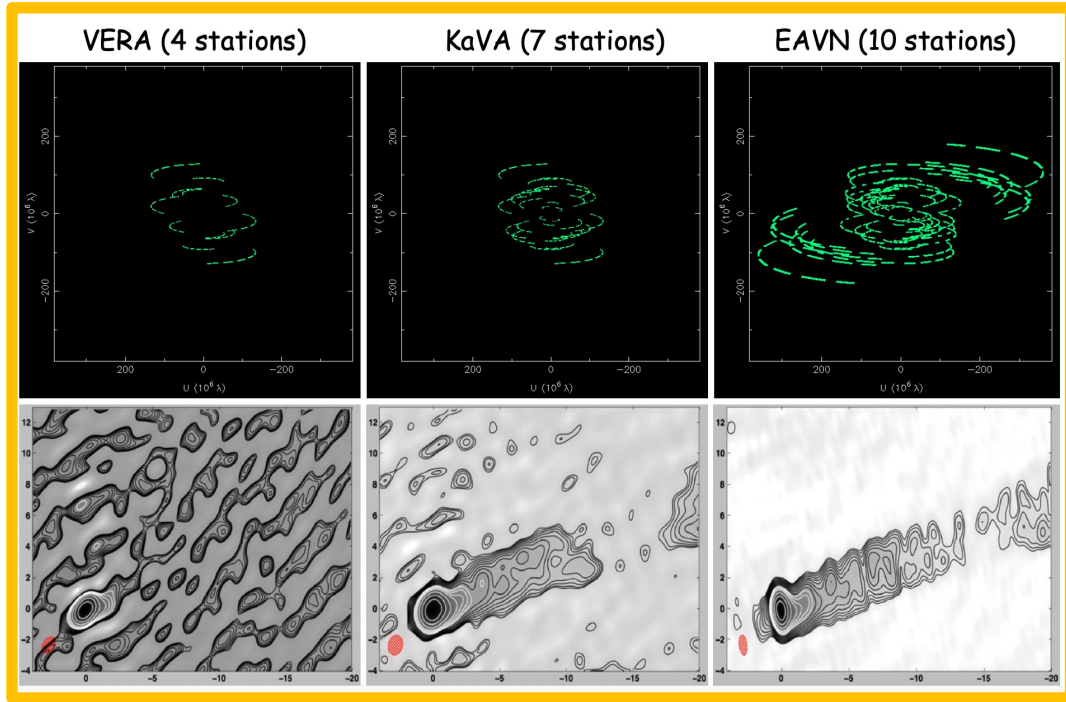
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You need as many $V(u,v)$ points as possible to reconstruct as robustly as possible the surface brightness distribution of the source



Imaging intro: Fourier Transform imaging and sampling function

Credits: Prof. Kazuhiro Hada



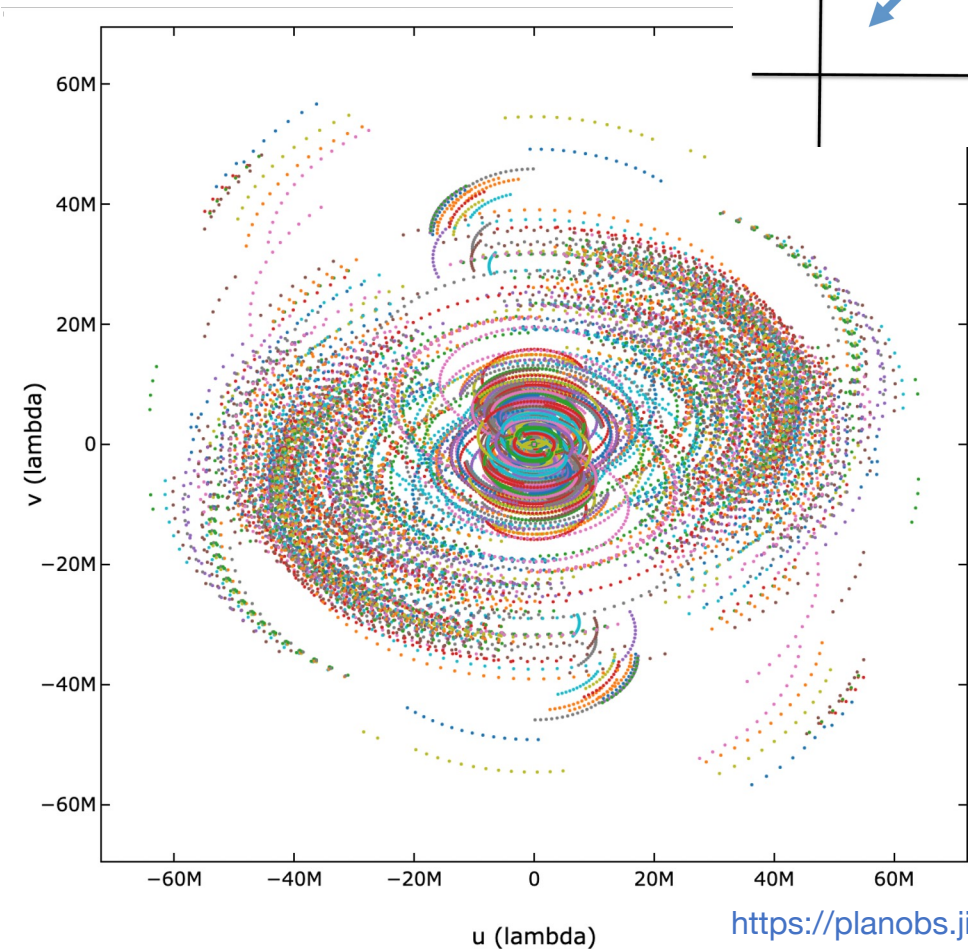
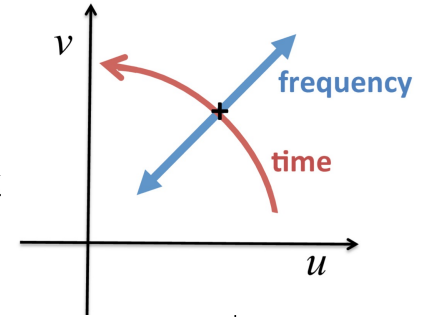
An ideal interferometer = measurements on a regularly highly sampled rectangular grid.

An image of would then be made by simply applying a Fourier transform

But, arrays provide (poorly) sampled Fourier Transform of the radio brightness region of sky

You need as many $V(u,v)$ points as possible to reconstruct as robustly as possible the surface brightness distribution of the source

S = sampling function
 = 1 where there is a measurement in the uv plane
 = 0 otherwise



12 h
>30 antennas

<https://planobs.jive.eu/>

Gridding

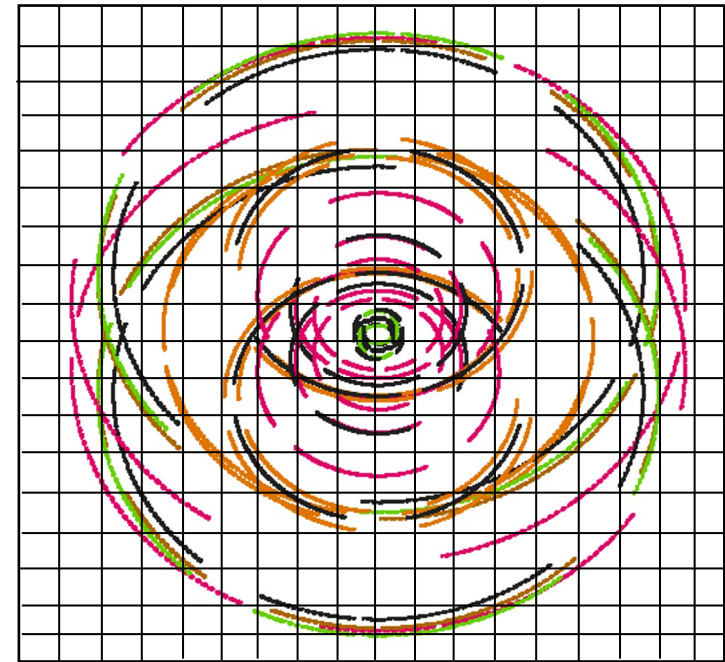
**But, arrays provide (poorly) sampled Fourier Transform of the radio brightness region of sky
AND
There will always be gaps in the u-v plane!**

Two approaches

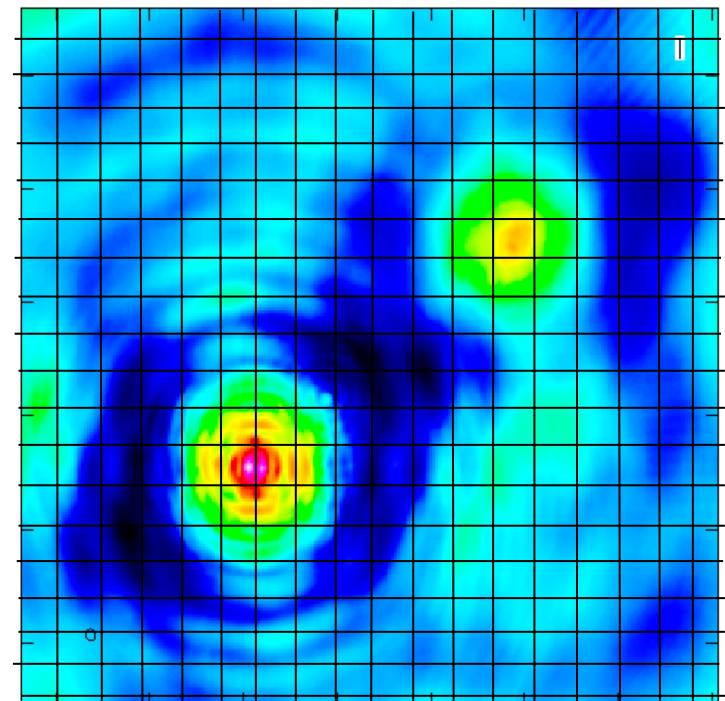
Direct Fourier Transform (DFT) = FT evaluated at every point of a rectangular grid – $O(N^2)$ operations
impractical for large number of visibilities

Fast Fourier Transform (FFT) = interpolate the data onto a rectangular grid – $O(N \log N)$ operations
It saves a lot of computing time!!

This FFT method requires the observed visibilities to be interpolated on a regular grid
Usually we define the grid in the image plane, where
grid spacing = cell size

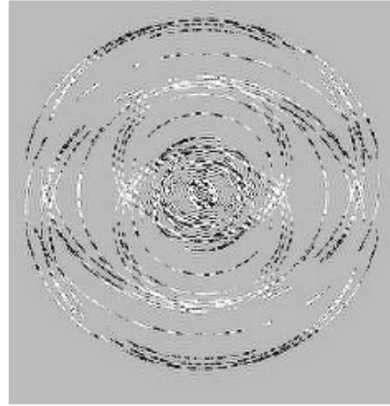


NxN grid



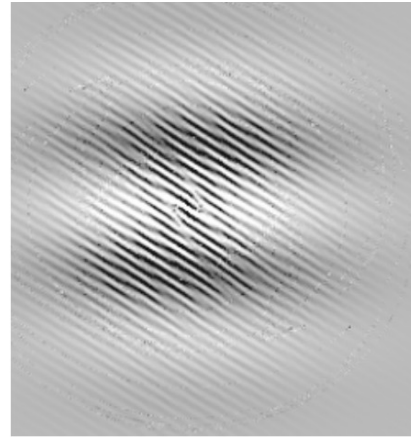
The need for deconvolution

Sampled visibilities $V'(u,v)$

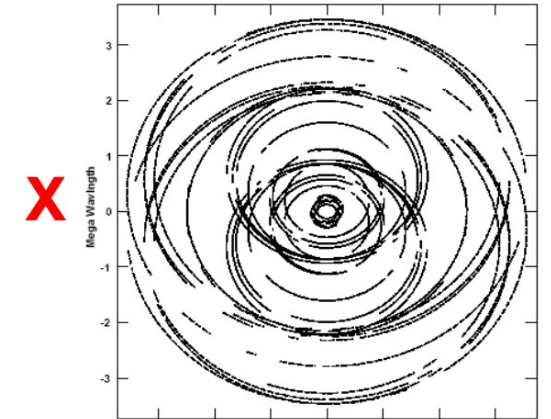


=

True visibilities $V(u,v)$



Sampling function $S(u,v)$



X

Gaps in the uv-plane will make sidelobes in the dirty beam

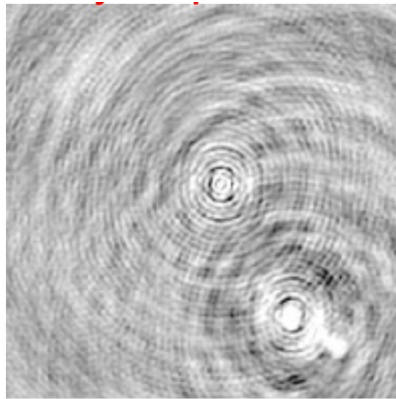
«Visibility domain»

FT



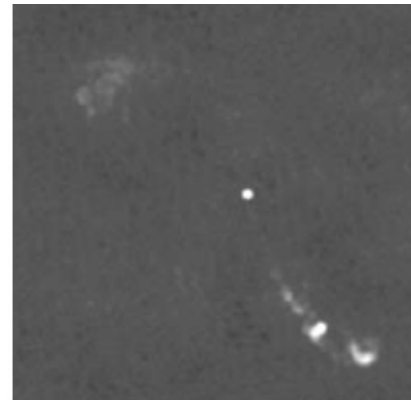
«Image domain»

Dirty image $B'(l,m)$



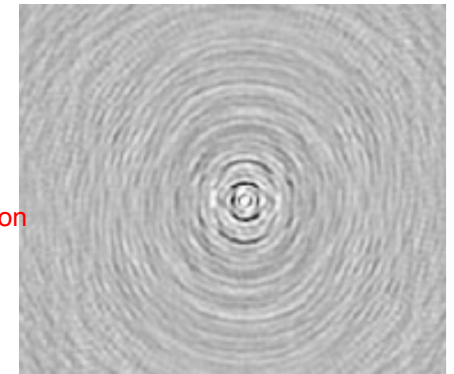
=

True sky $B(l,m)$



* convolution

Dirty beam (l,m)

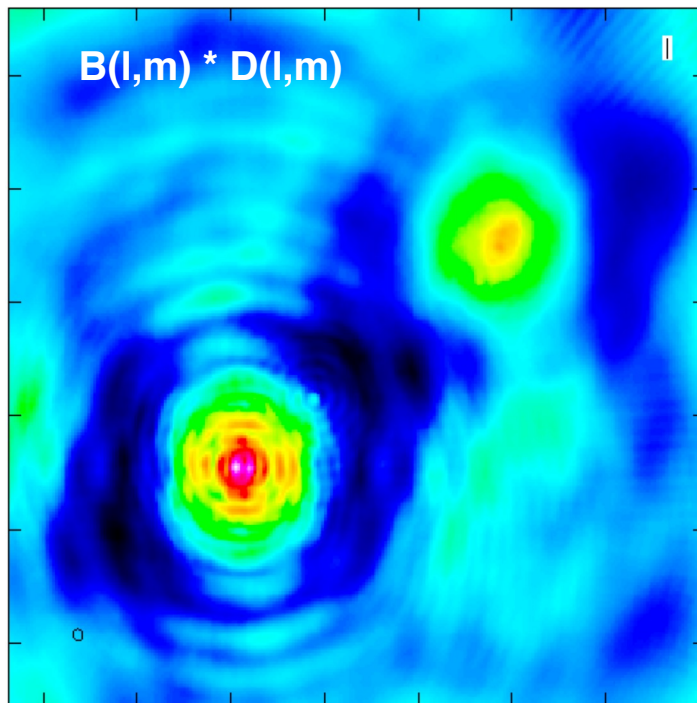


The dirty image is not the true image of the source, since the sampled visibilities are not the true visibilities

Corrections of the effect of Fourier sampling deficiencies on the dirty image = CLEAN algorithms

Deconvolution

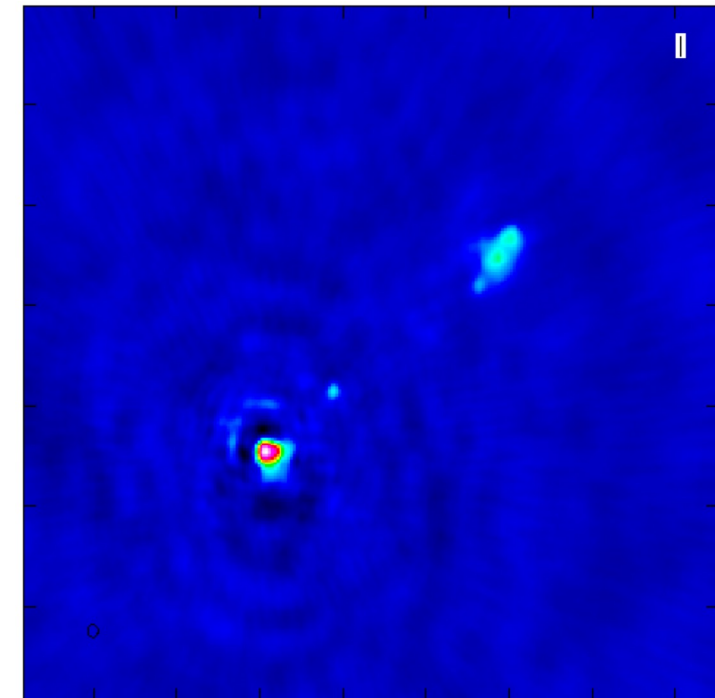
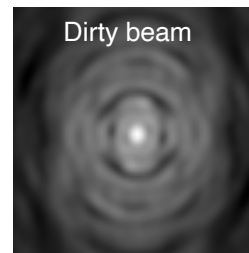
....Why do we need all of this again?



From «dirty image»



Deconvolve the intrinsic source
brightness distribution $B(l,m)$ from the
dirty beam $D(l,m)$



To «CLEAN image»

Deconvolution

Since only a finite number of (noisy) samples are measured, to recover $B(l,m)$ we need **some stable non-linear approach + *a priori* information**:

- $B(l,m)$ must be positive
- Radio sources do not resemble the dirty beam (i.e. sidelobes-like patterns)
- Sky is basically empty with just a few localised sources

B = Intrinsic source brightness distribution

D = dirty beam = point spread function (PSF)

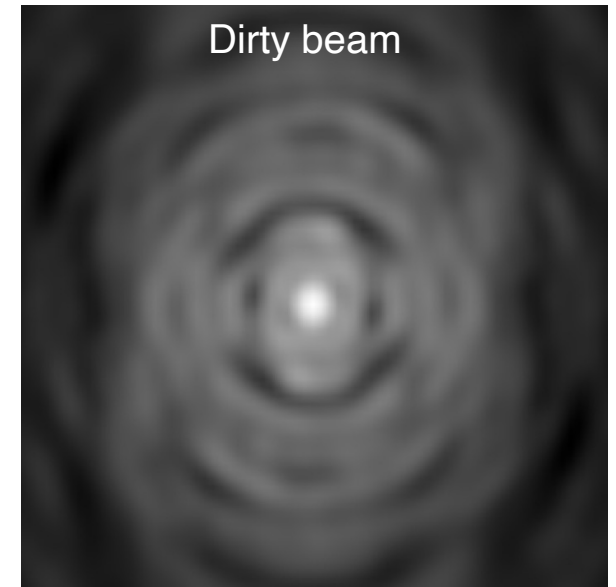
S = Sampling function

Convolution

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We know this!

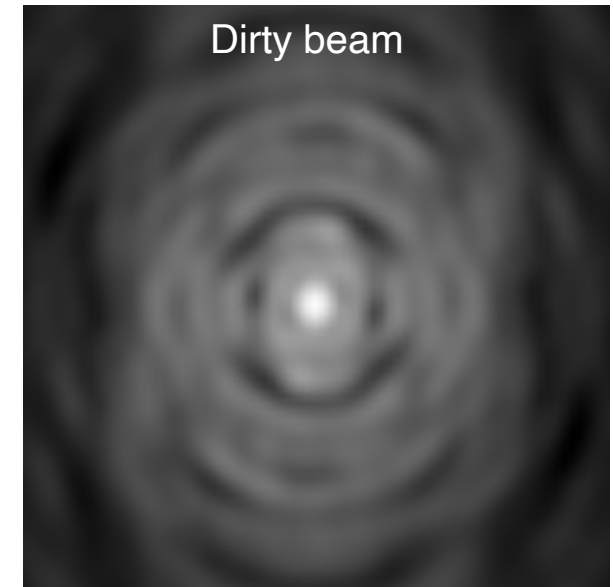
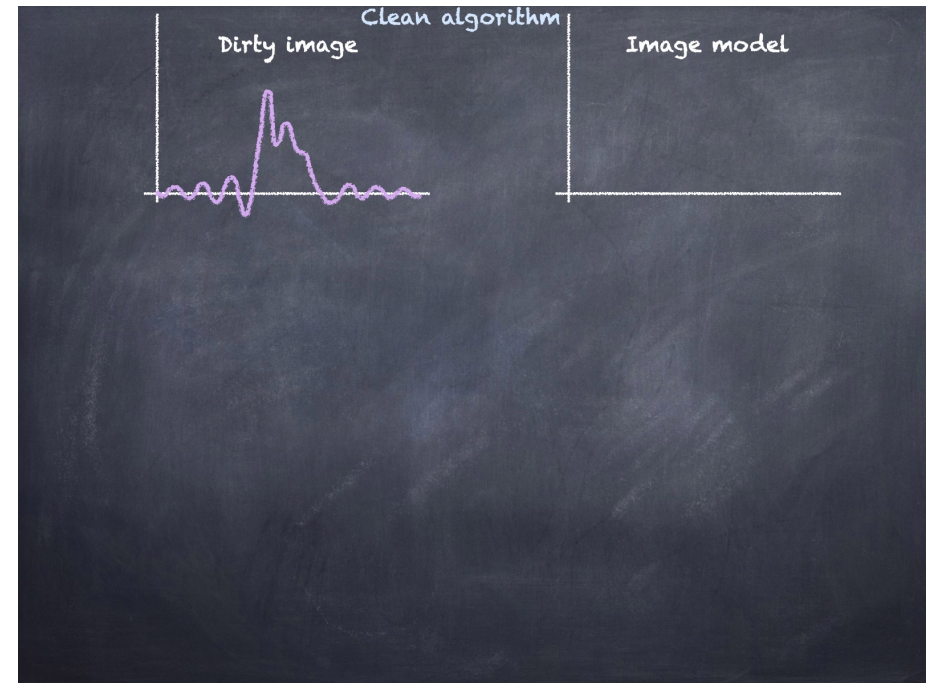
To recover B we have **“just” to deconvolve** the $D(l,m)$ term



Deconvolution

CLEAN method principal steps (Clark's algorithm):

- 1) **Initialize a residual map (first image = dirty image)**
- 2) Identify strongest peak as a delta component
- 3) Record the position and magnitude in a model (clean components), subtract it from the dirty image
- 4) Go to 1) unless you reach the stopping criterion
- 5) Convolve the model (clean components) with an idealized CLEAN beam (elliptical Gaussian fit of the main lobe of the dirty beam)
- 6) Add the residual of the dirty image to the CLEAN image



Deconvolution

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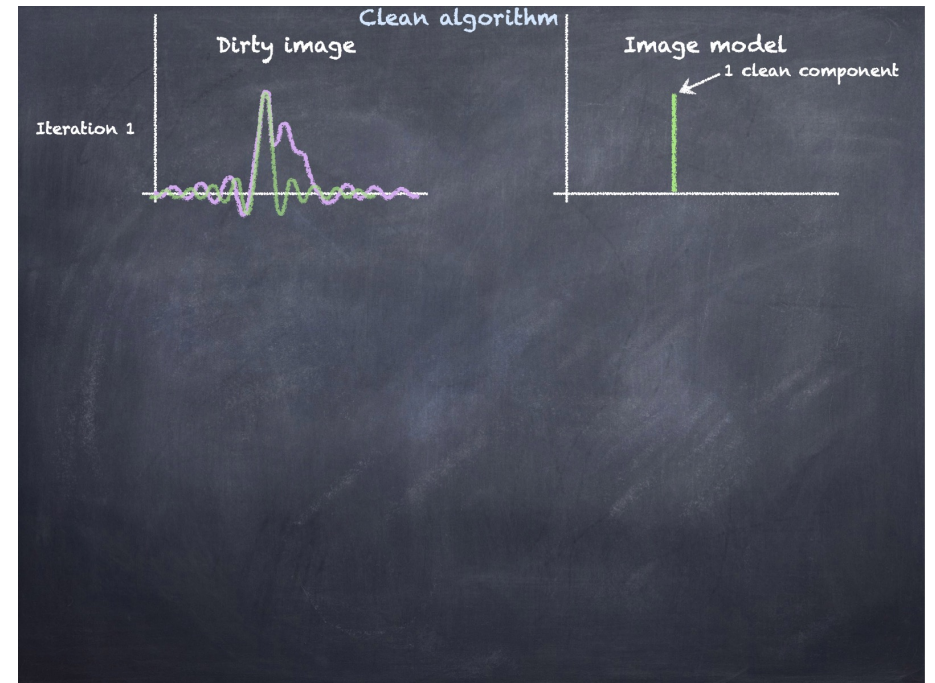
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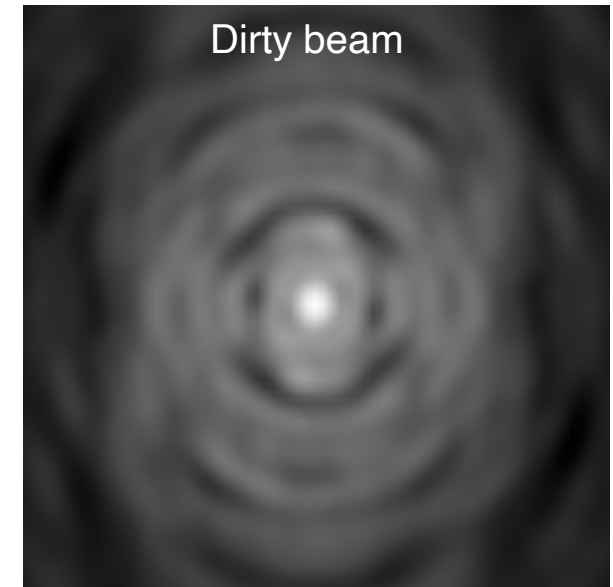
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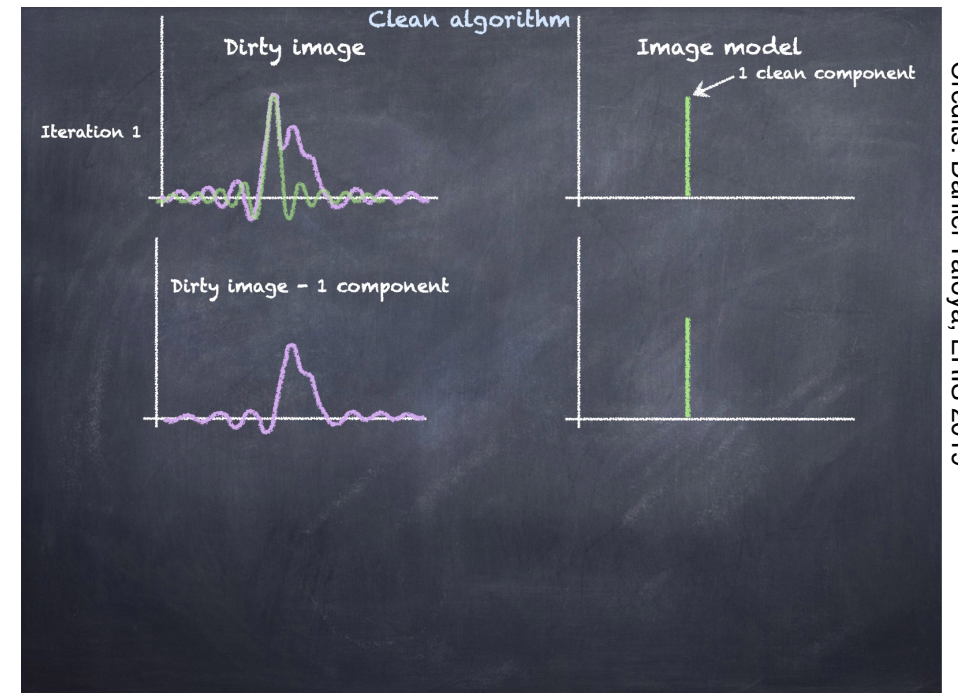
Credits: Daniel Tafuya, ERIIS 2019



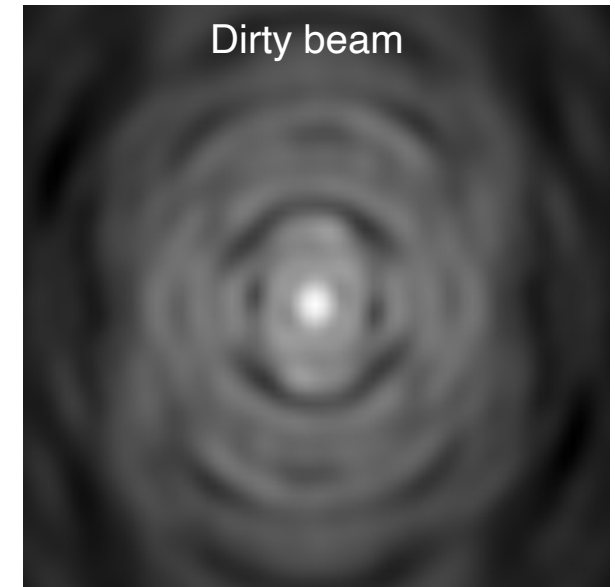
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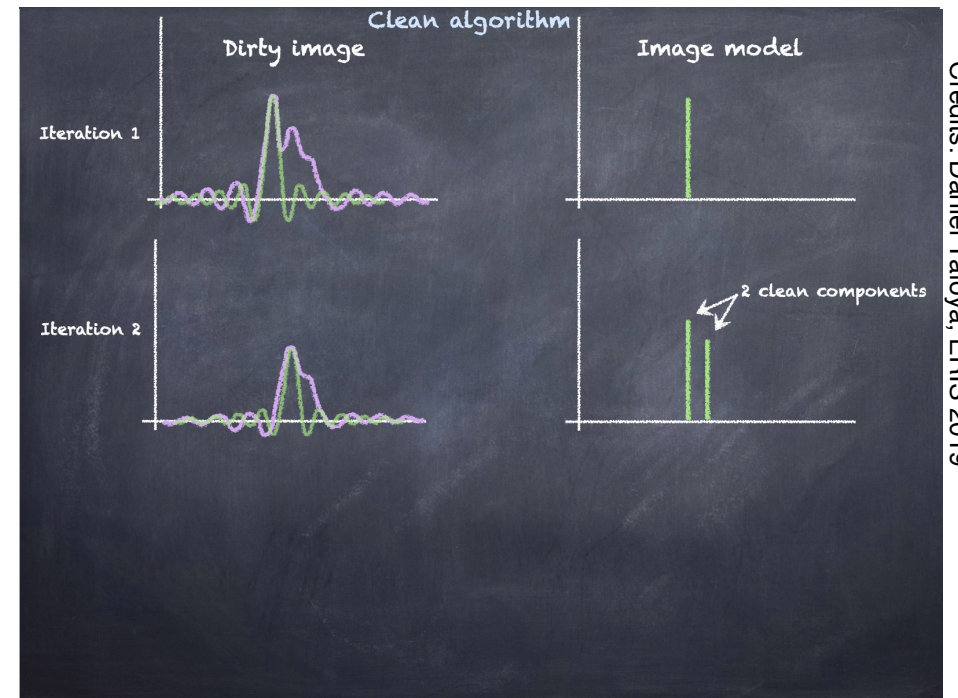
Credits: Daniel Tafuya, ERLS 2019



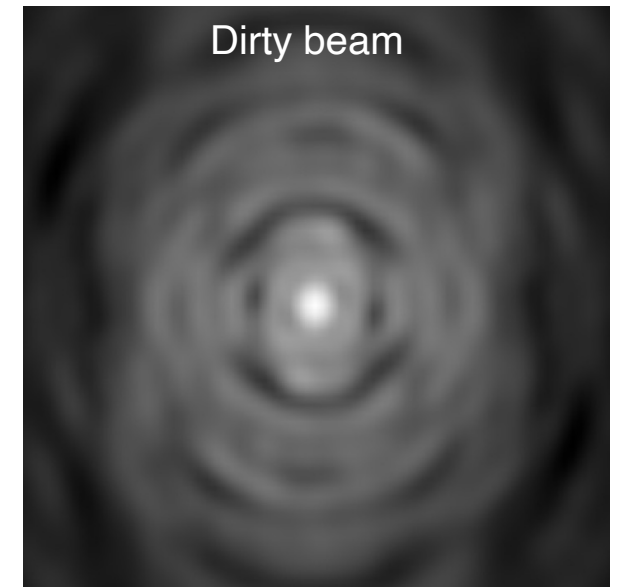
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Credits: Daniel Tafuya, EIRIS 2019



Deconvolution

CLEAN method principal steps (Clark's algorithm):

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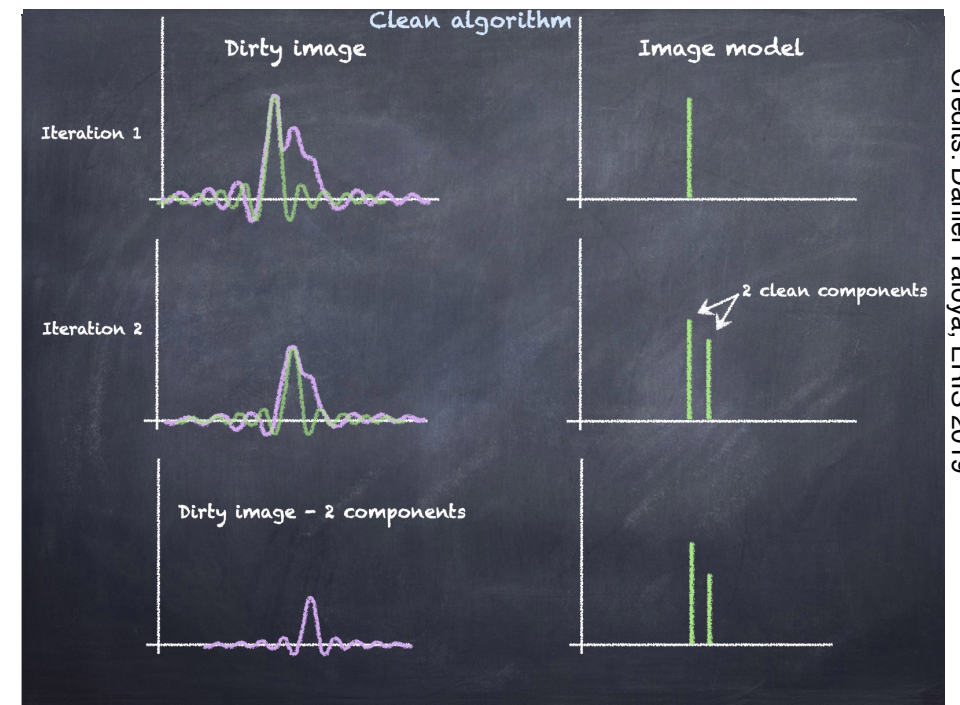
2) Identify strongest peak as a delta component

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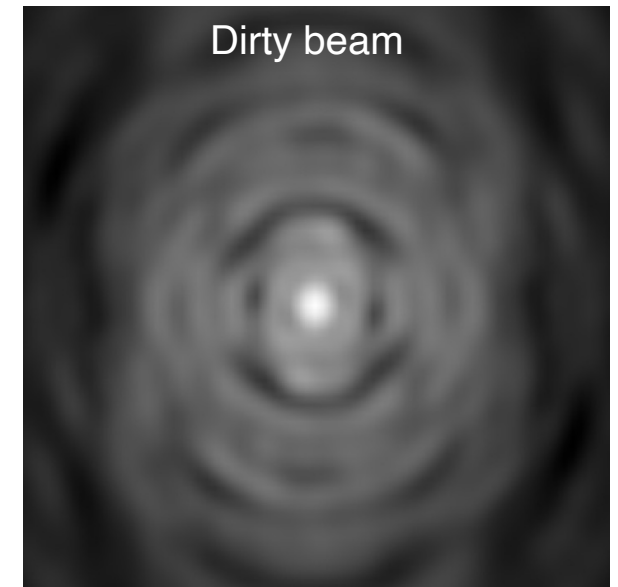
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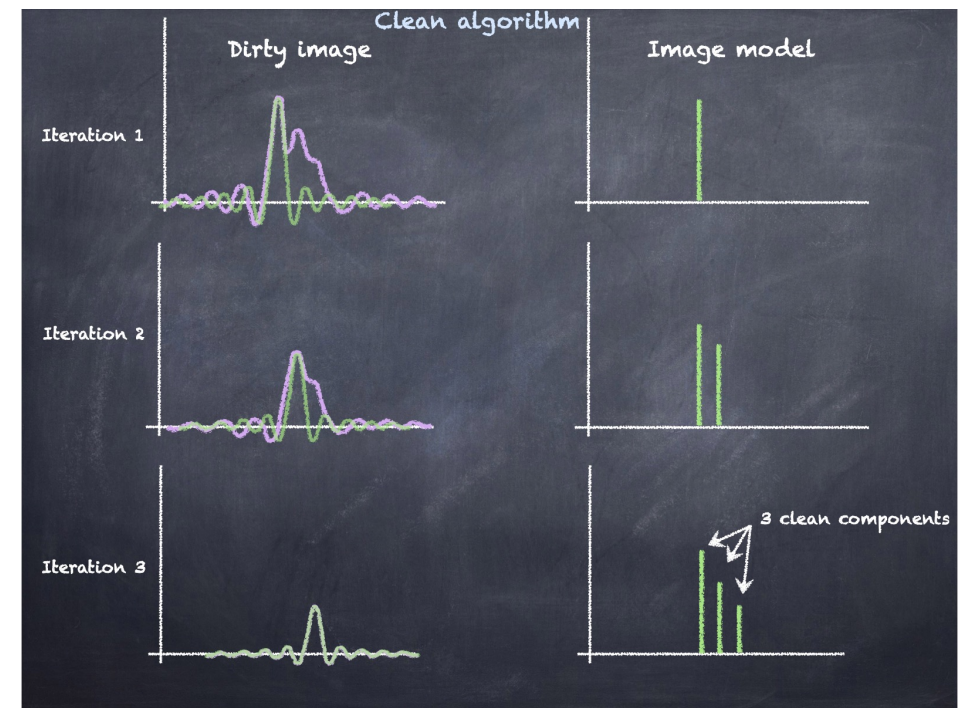
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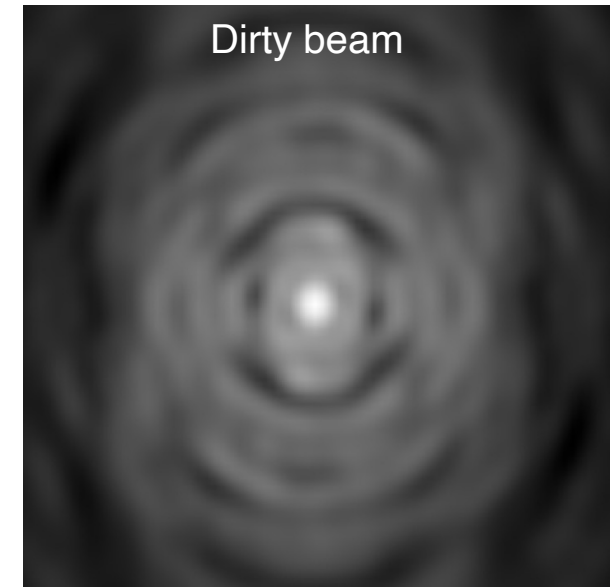
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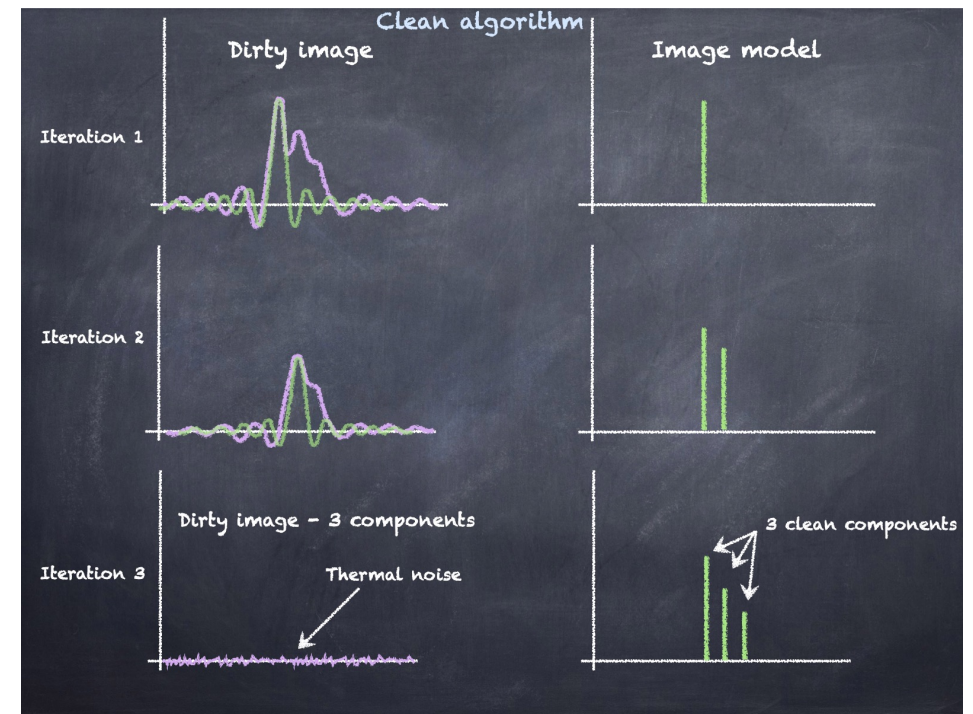
Credits: Daniel Tafuya, EIRIS 2019



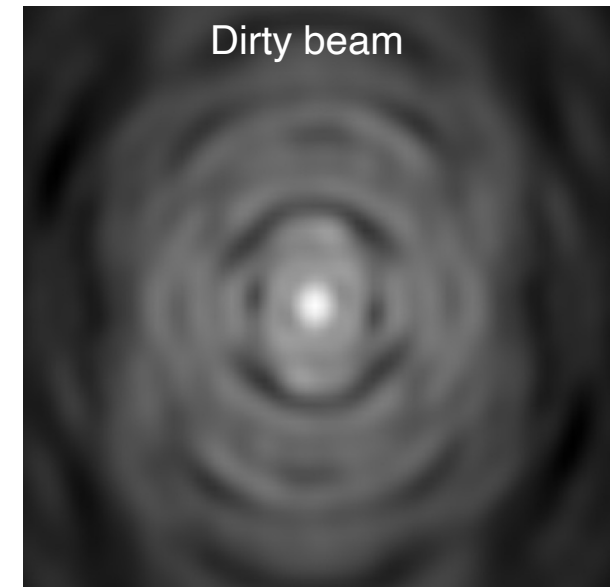
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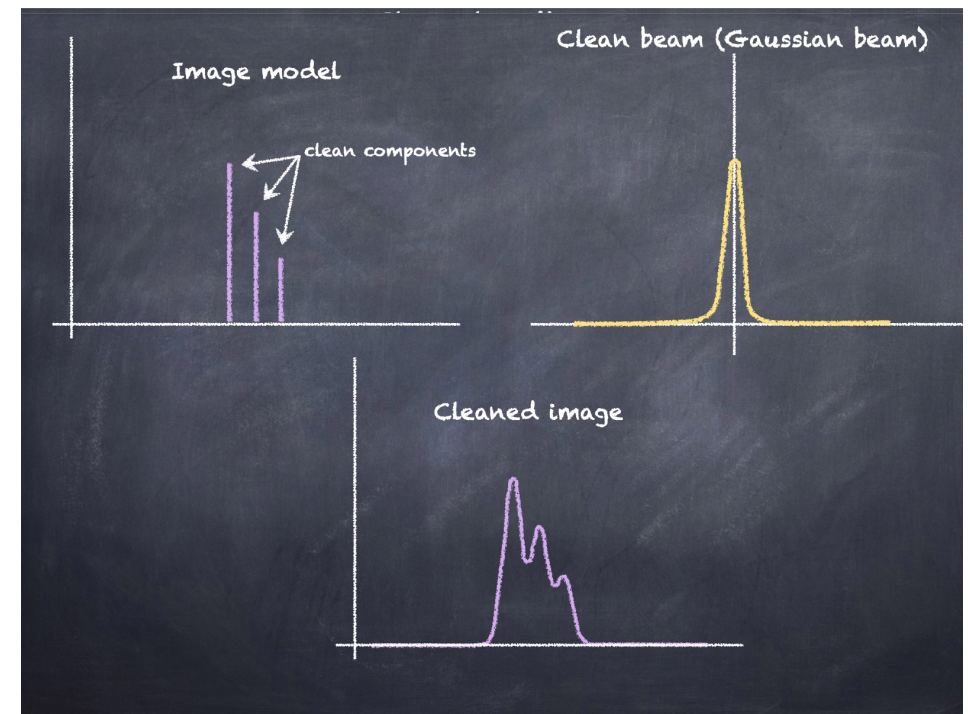
Credits: Daniel Tafuya, ERIS 2019



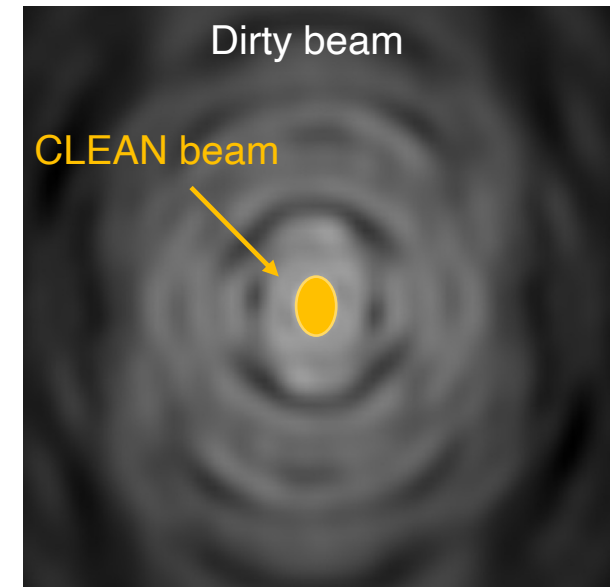
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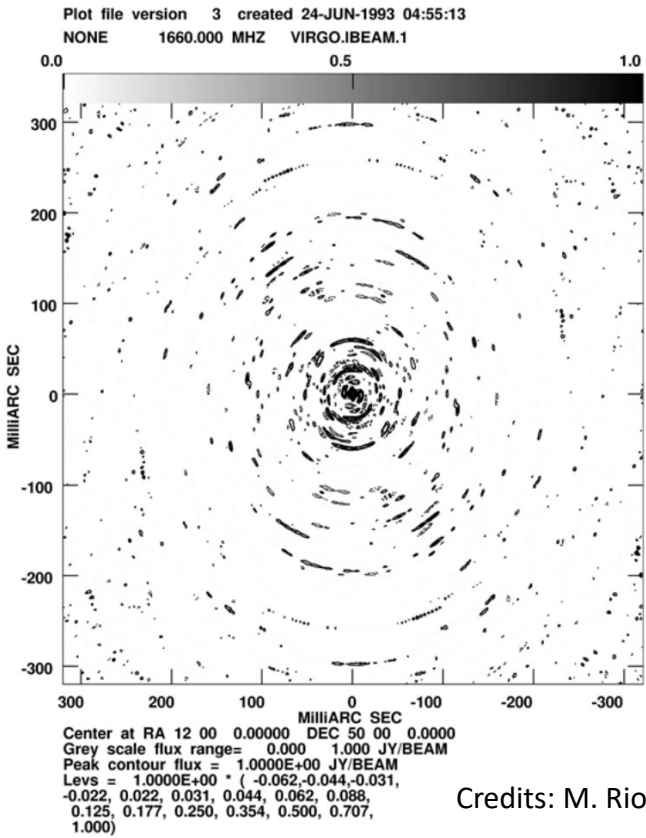


Credits: Daniel Tafuya, ERIS 2019

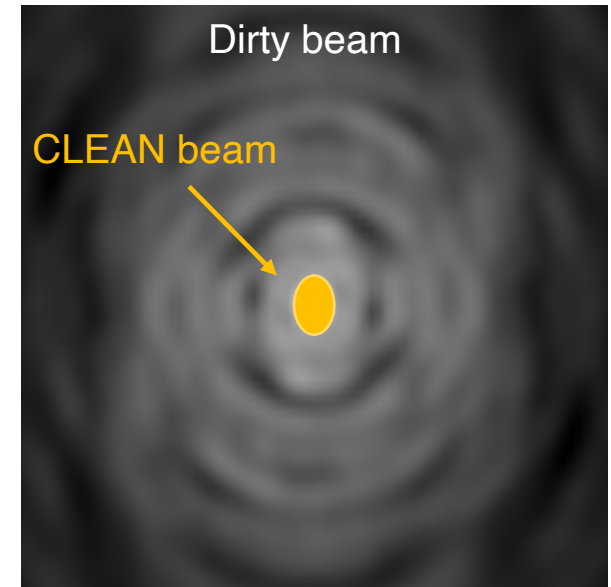
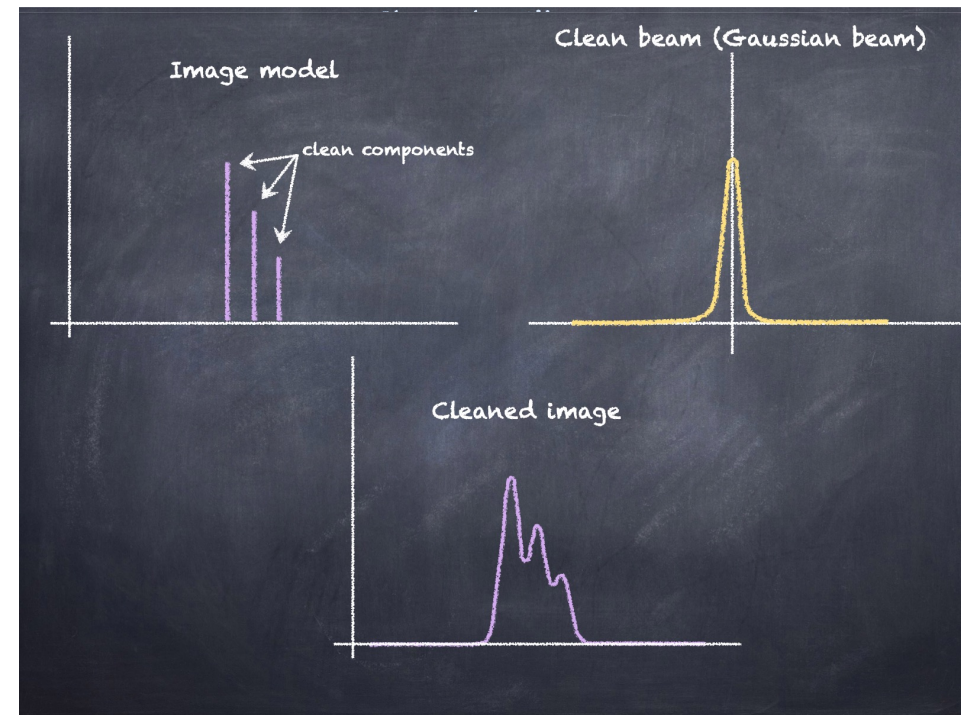
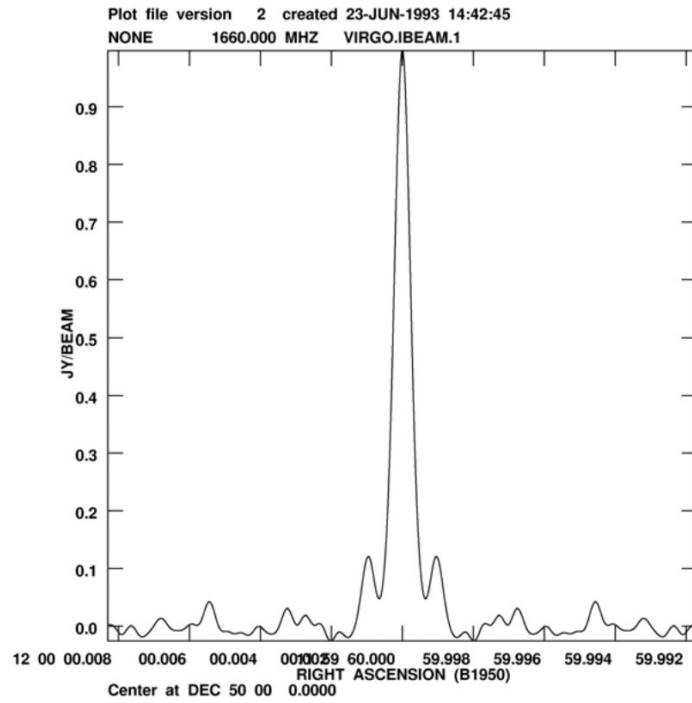


Deconvolution

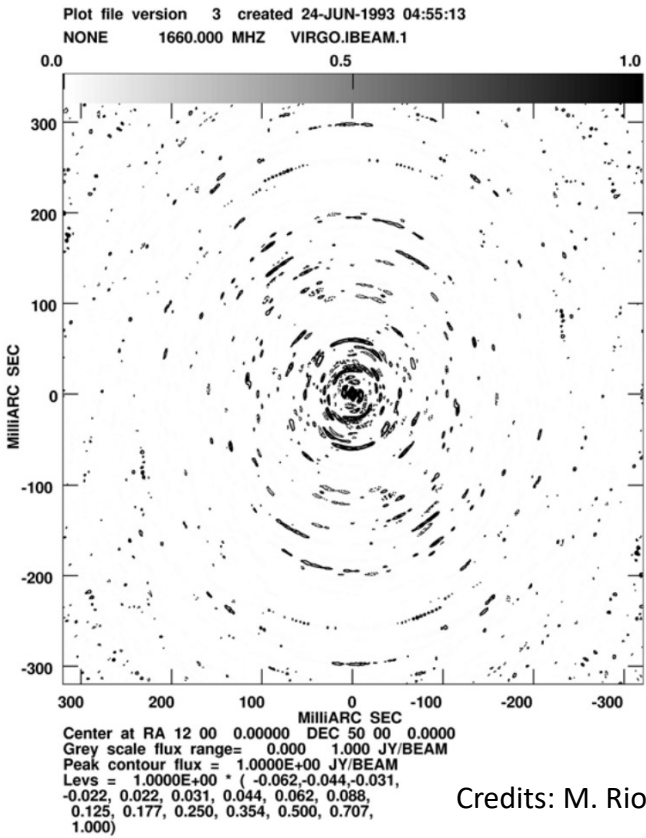
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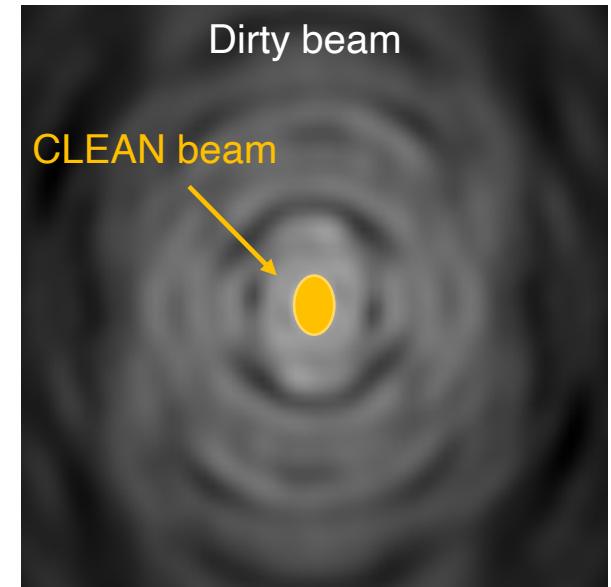
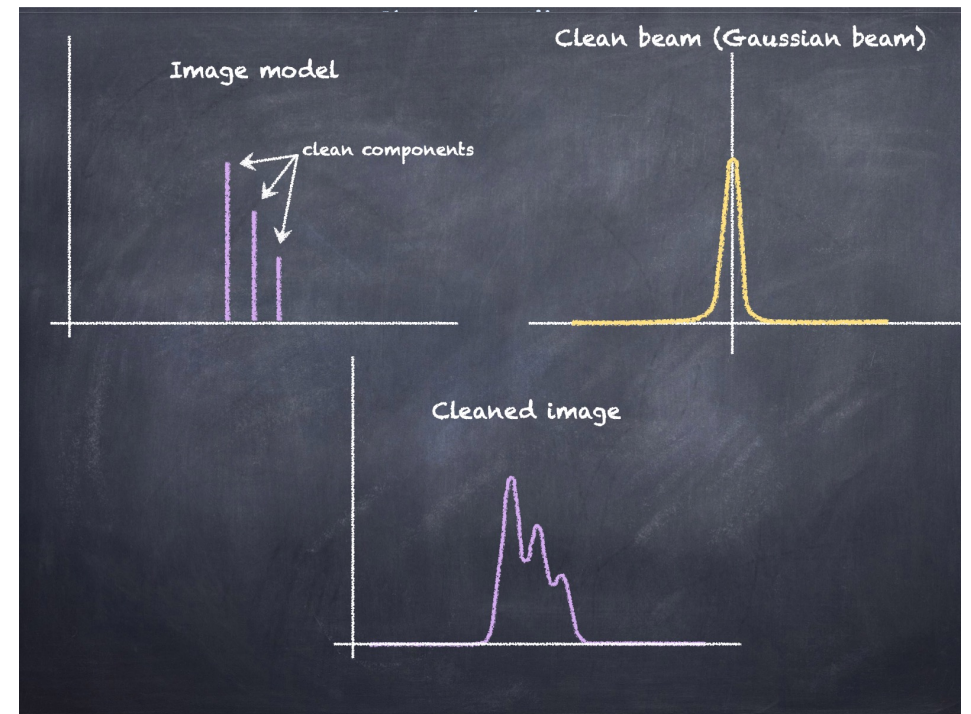
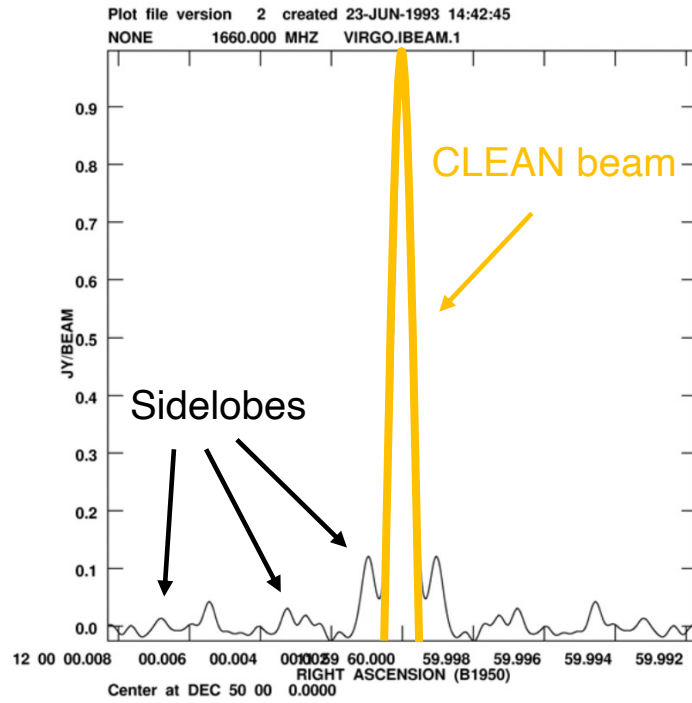
Credits: M. Rioja



Deconvolution



Credits: M. Rioja

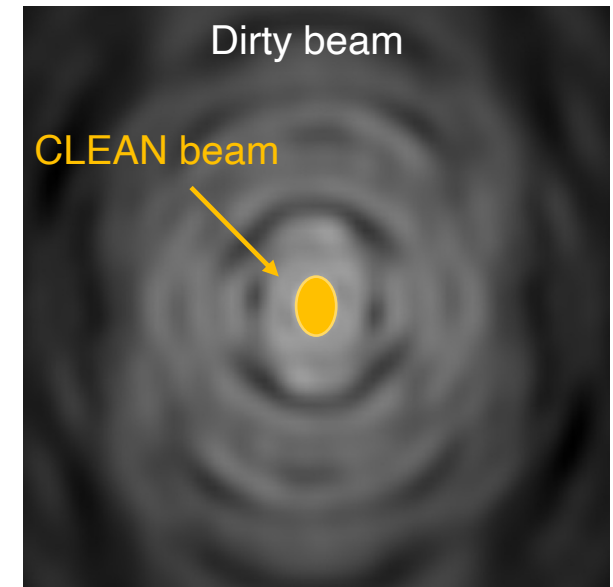
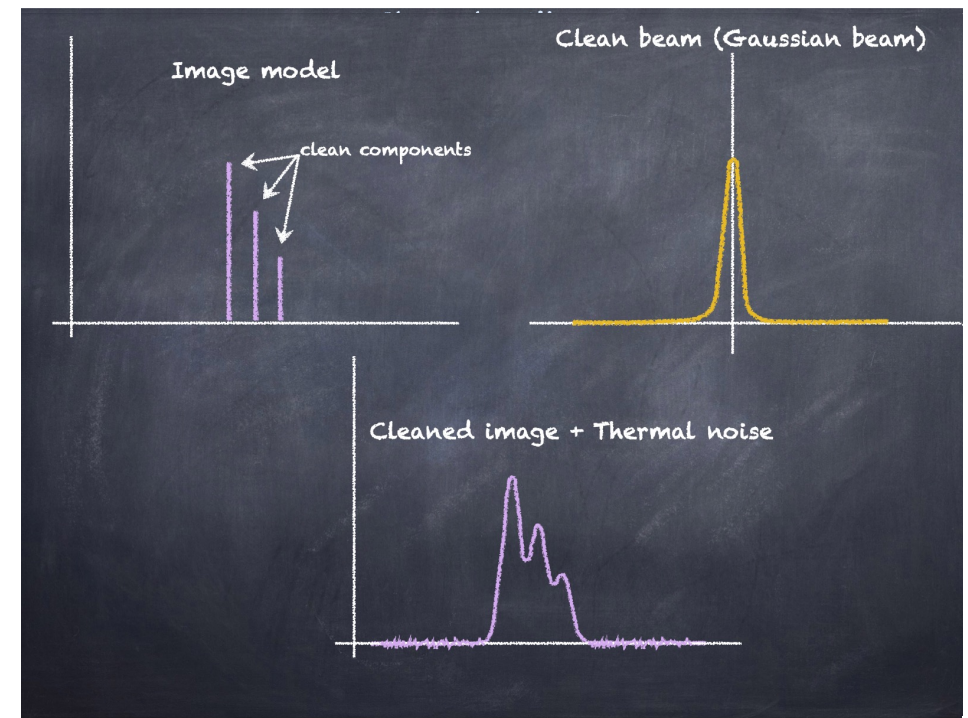


Deconvolution

CLEAN method principal steps (Clark's algorithm):

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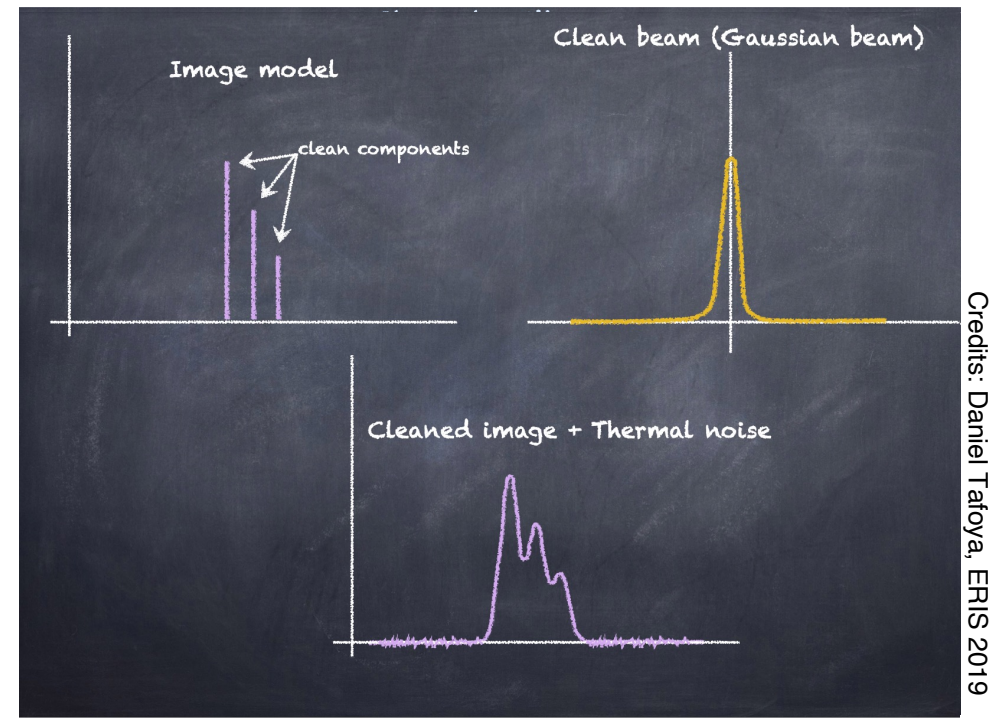


Deconvolution

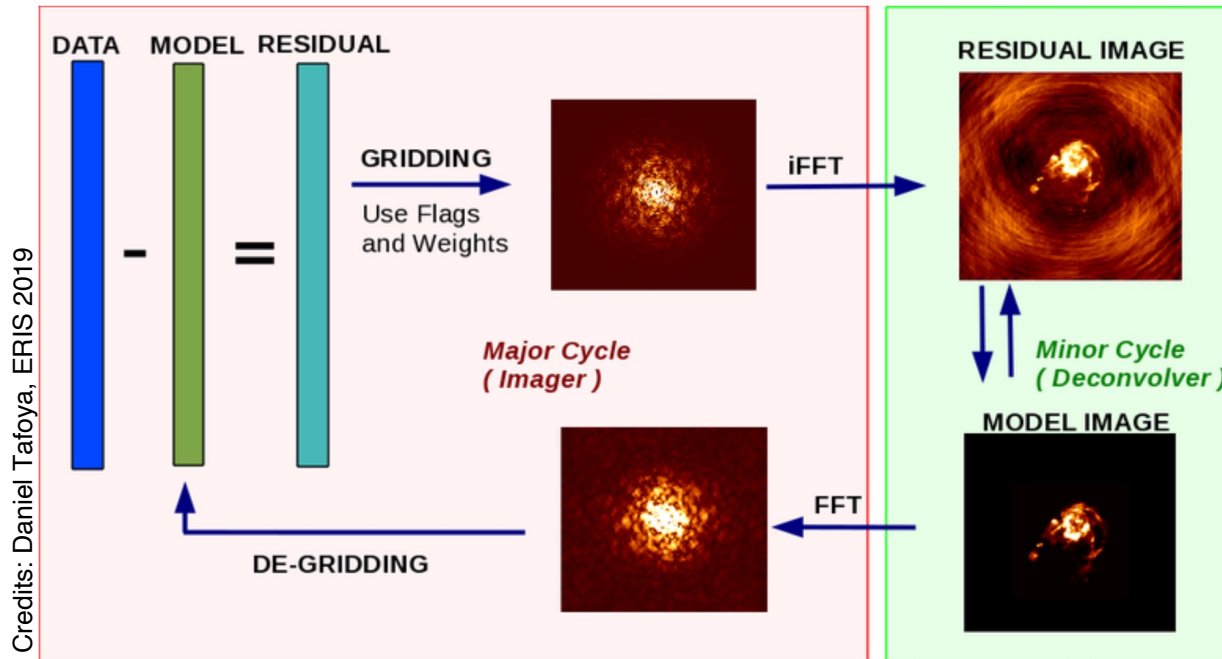
CLEAN method principal steps (Clark's algorithm):

Minor cycle

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Credits: Daniel Tafuya, EIRIS 2019



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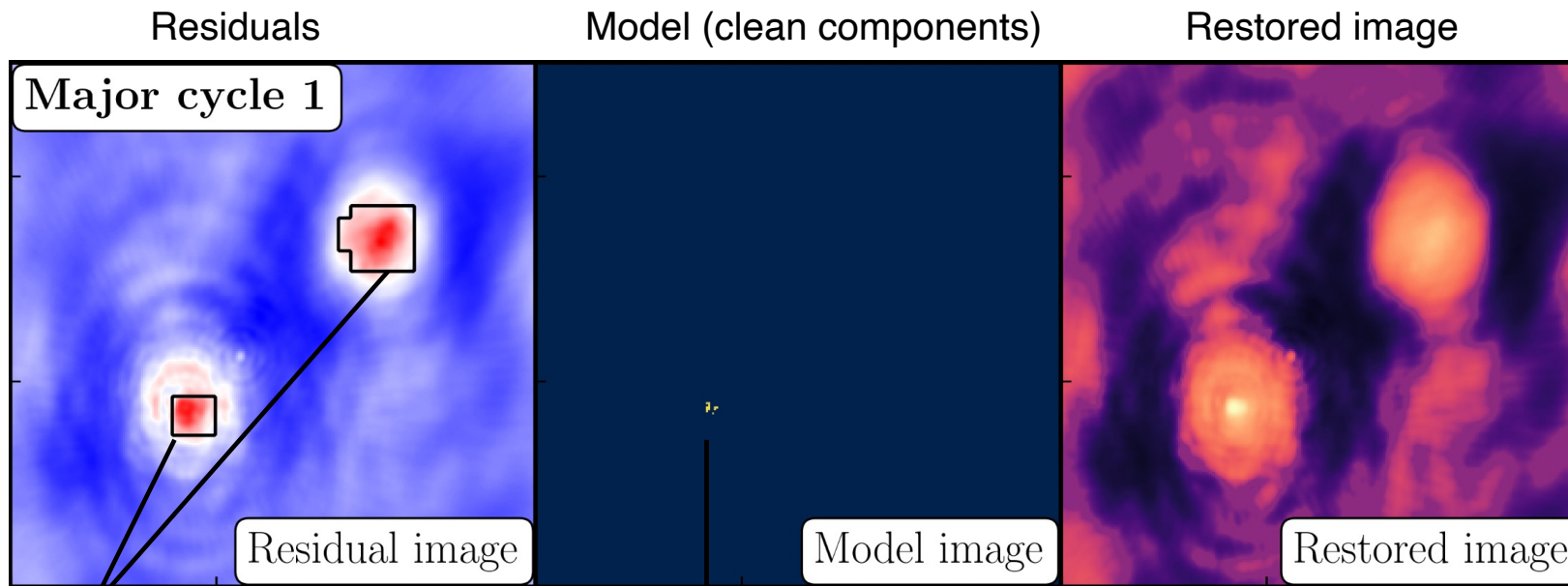
The major cycle implements FT between the data and image domains

The minor cycle operates purely in the image domain

(The 2-cycles approach makes the deconvolution faster)

Deconvolution in practice

CLEAN in action



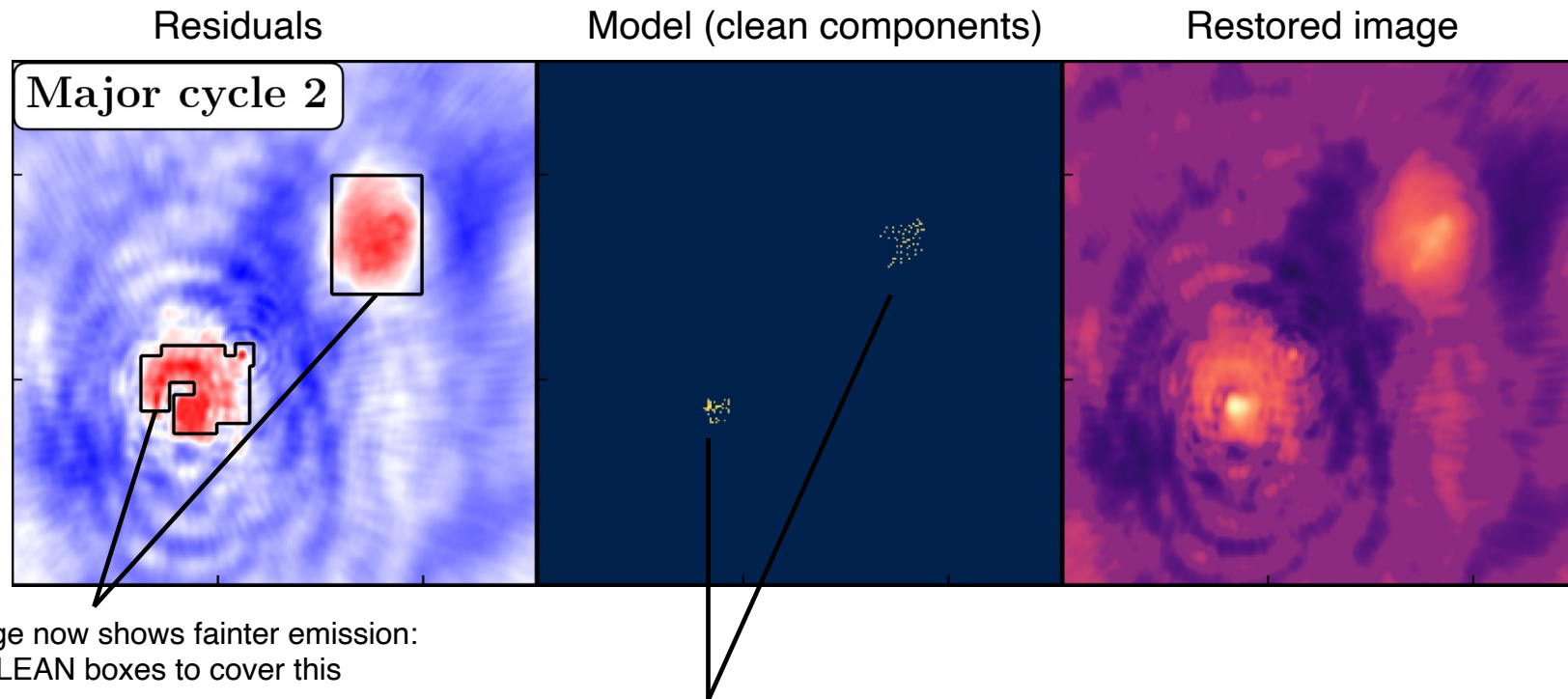
To make this process converge faster we use the so-called CLEAN boxes (mask)

Also useful to not let CLEAN go to sidelobes (see next slides)

CLEAN components obtained during several minor cycles

Deconvolution in practice

CLEAN in action

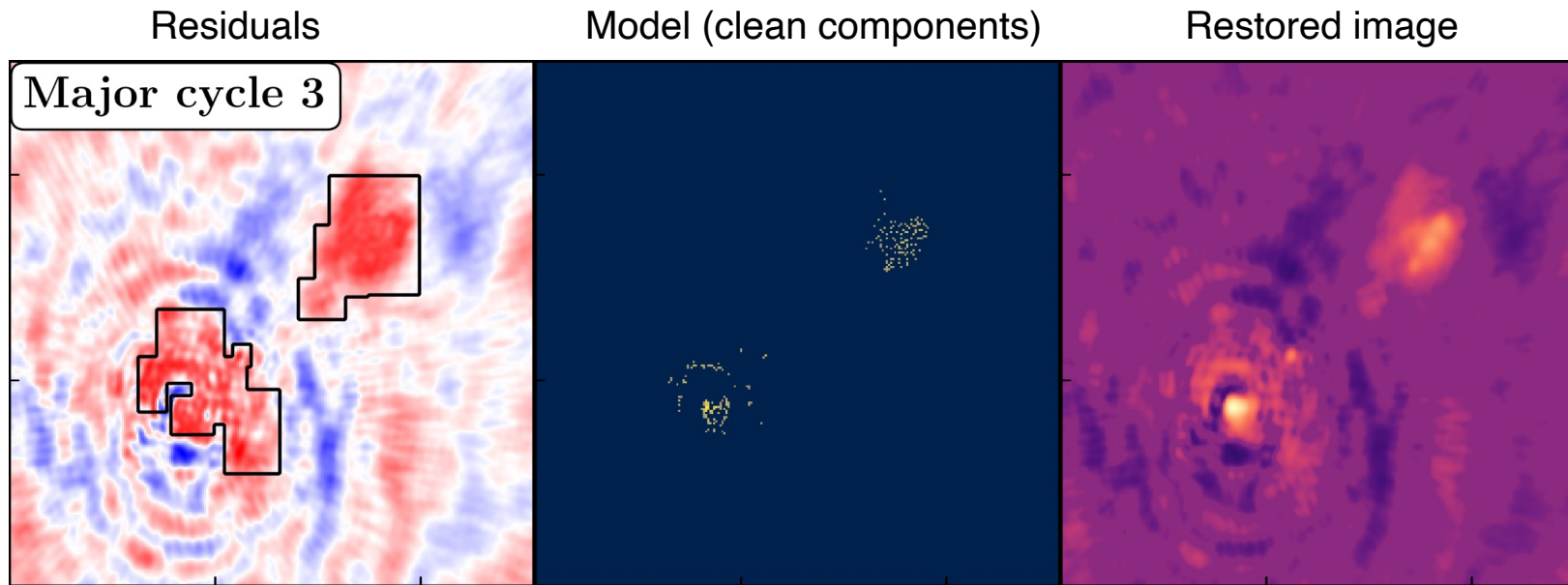


The residual image now shows fainter emission:
we enlarge the CLEAN boxes to cover this

New CLEAN components added to the previous ones

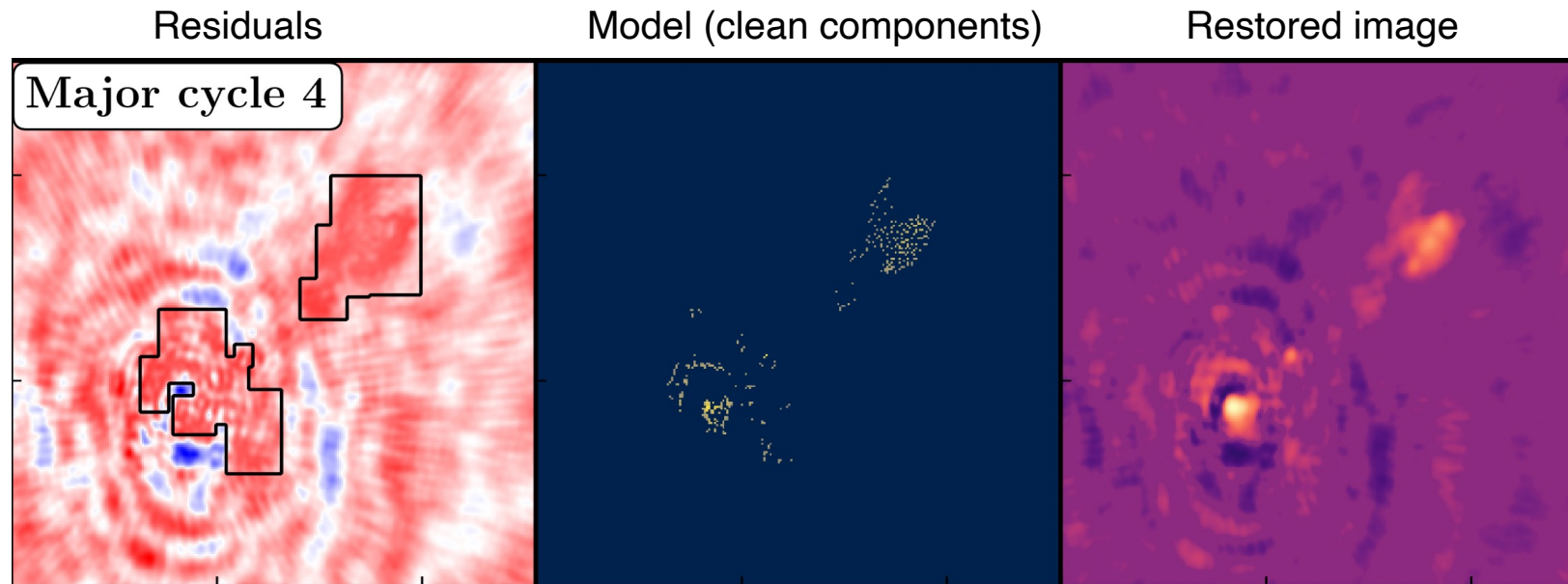
Deconvolution in practice

CLEAN in action



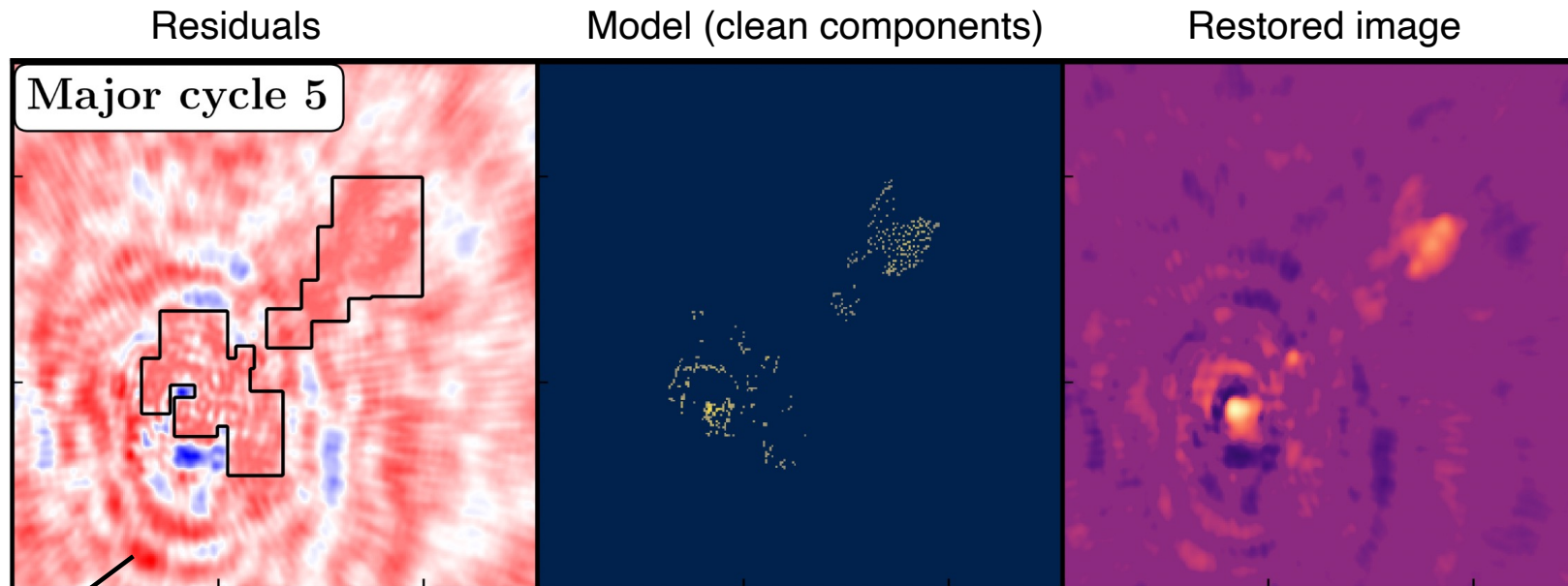
Deconvolution in practice

CLEAN in action



Deconvolution in practice

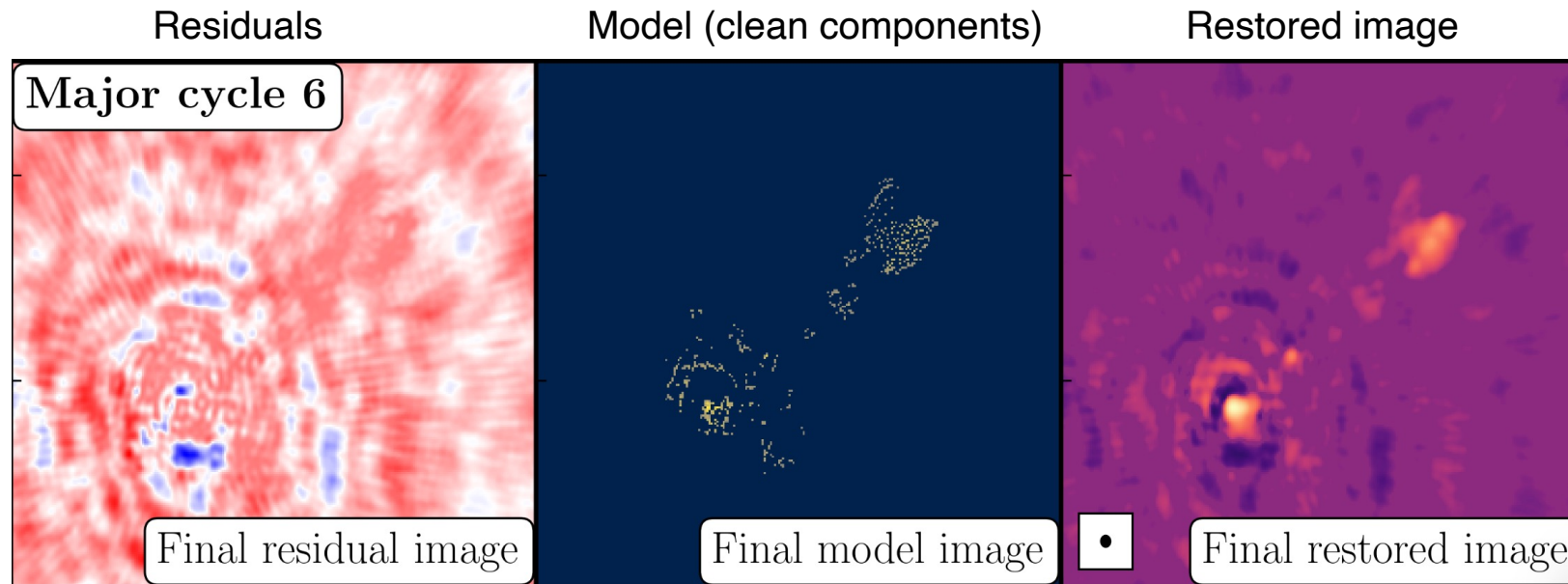
CLEAN in action



This emission is brighter BUT it's due to sidelobes!
It's always a good idea take a look at the dirty beam before starting cleaning
+ CLEAN boxes prevent the CLEANing of sidelobes

Deconvolution in practice

CLEAN in action



Residual image should look like «only noise»

CLEANing stopping criteria

- **Visually**, when your residuals contain only noise – this means that you cleaned all the flux density of the source
- **Convergence**: Check the logger for max-min (possibly symmetrical), total flux density should increase while cleaning (if not, stop), noise level should decrease (if it does not change anymore, stop → overcleaning)
- **Negative peak identified** (negatives can indicate that CLEAN is now working on sidelobes/noise, but it can also indicate that CLEAN is trying to fix earlier mistakes)
- **Smallest peak identified below a threshold** – which can be noise-based (e.g. 3 x theoretical noise estimated with exposure calculator – thermal noise)
- **Number of iterations** (not the best criterion, as you may end up doing too much or too little cleaning)

CLEANing issues and recognizing errors



CLEANING-related

- Interpolation of **unsampled (u,v) spacings** (in particular short spacings) : reconstruction of largest spatial scales is always an extrapolation (CLEAN boxes help)
- Assumption of **point-sources for extended structure** is not great
- **Under- and over-cleaning** are often an issue (over-cleaning: rms in logger does not change anymore)
- **Computationally expensive**, as it requires iterative, non-linear fitting process (CLEAN boxes help this too)

Calibration and data-handling related

- **Bandwidth and time smearing**
- **Amplitude/phase errors** from previous calibration and/or unflagged data (**symmetric/antisymmetric artefacts**)

Source-related

- **Variability** of the source
- **Spectral variations of the source** – multi frequency synthesis (gridding different frequencies on the same (u,v) grid is now standard)

CLEANing issues and recognizing errors



CLEANING-related

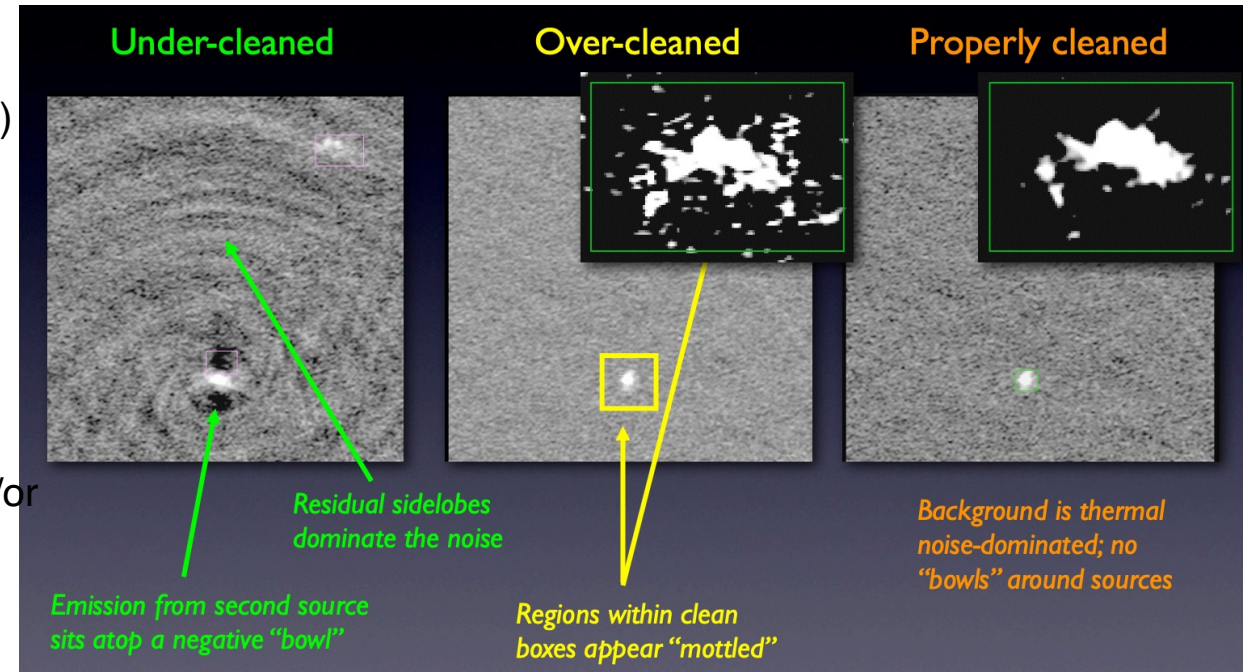
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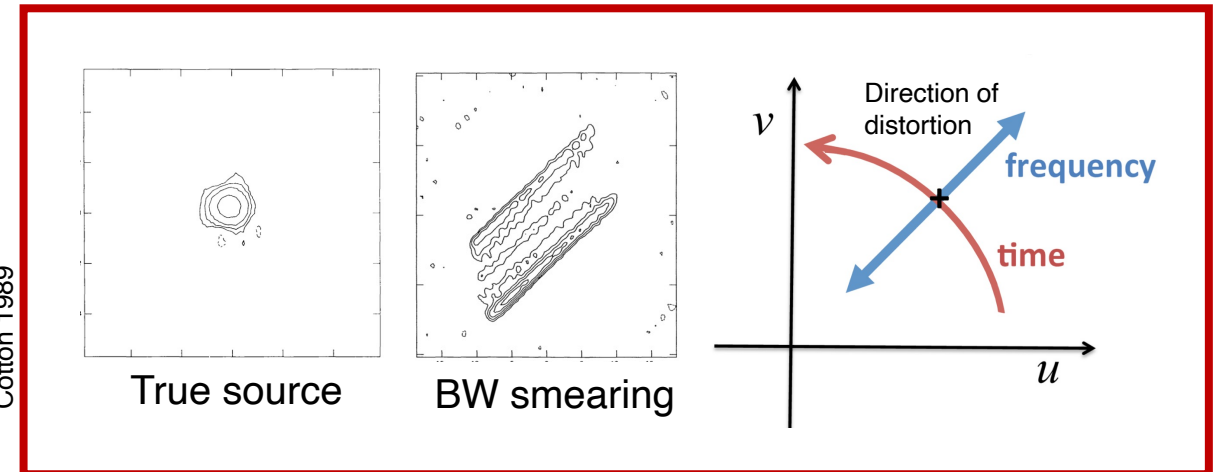
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Cotton 1989



CLEANing issues and recognizing errors



CLEANING-related

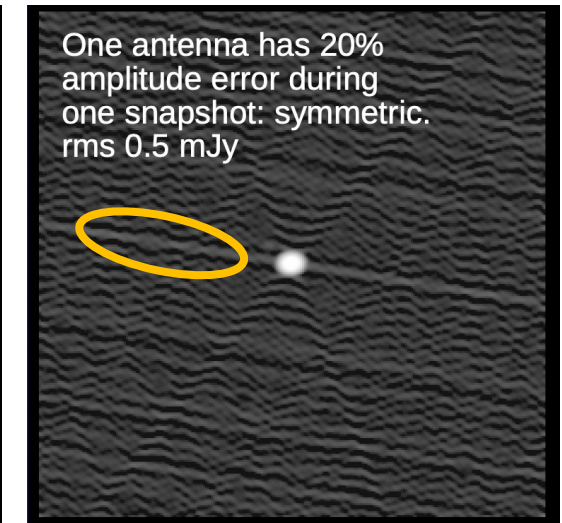
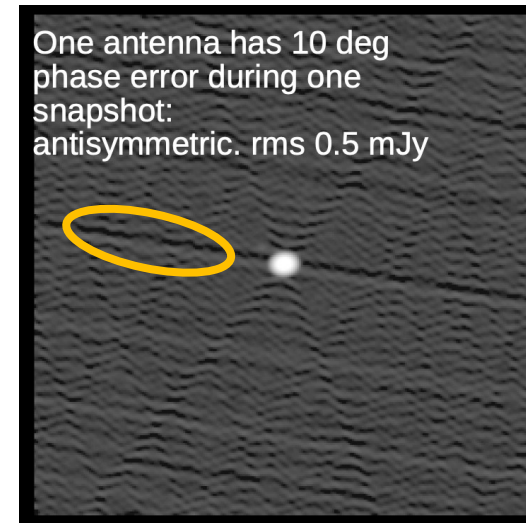
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CLEANing issues and recognizing errors



CLEANING-related

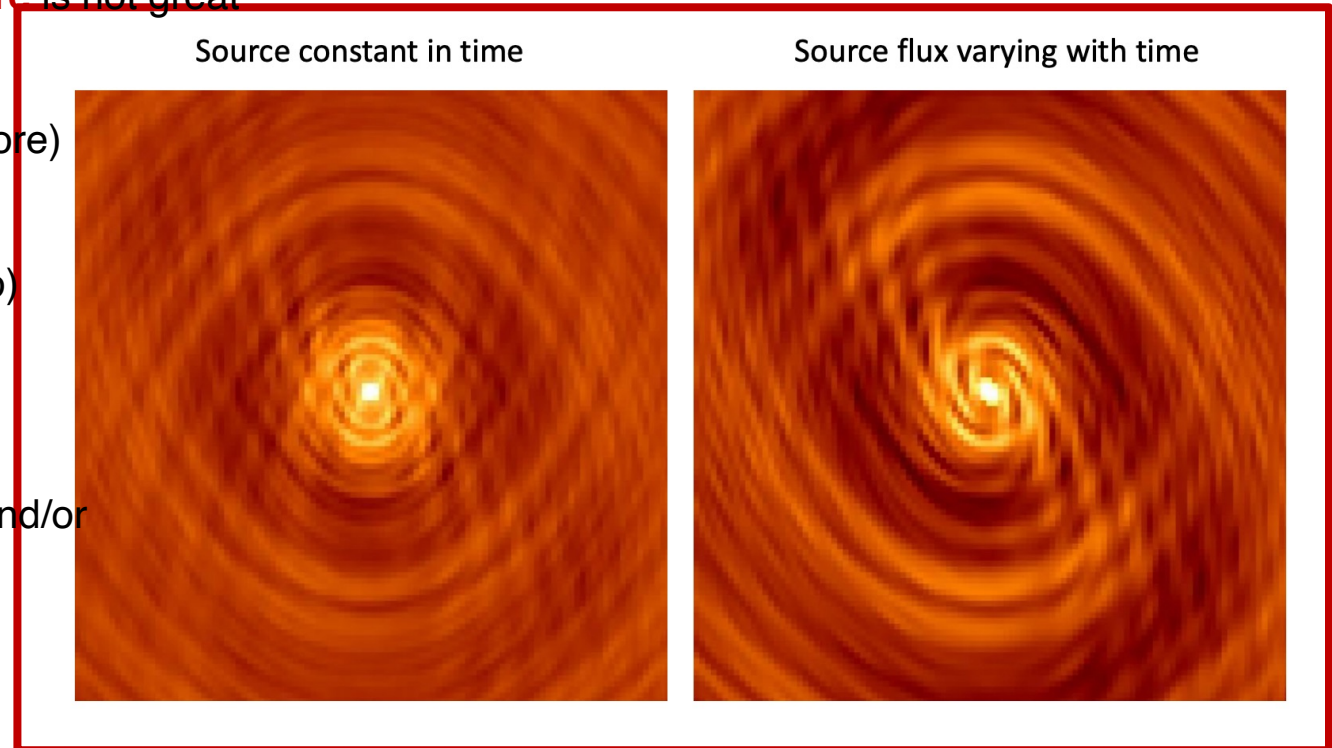
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CLEANing issues and recognizing errors



CLEANing-related

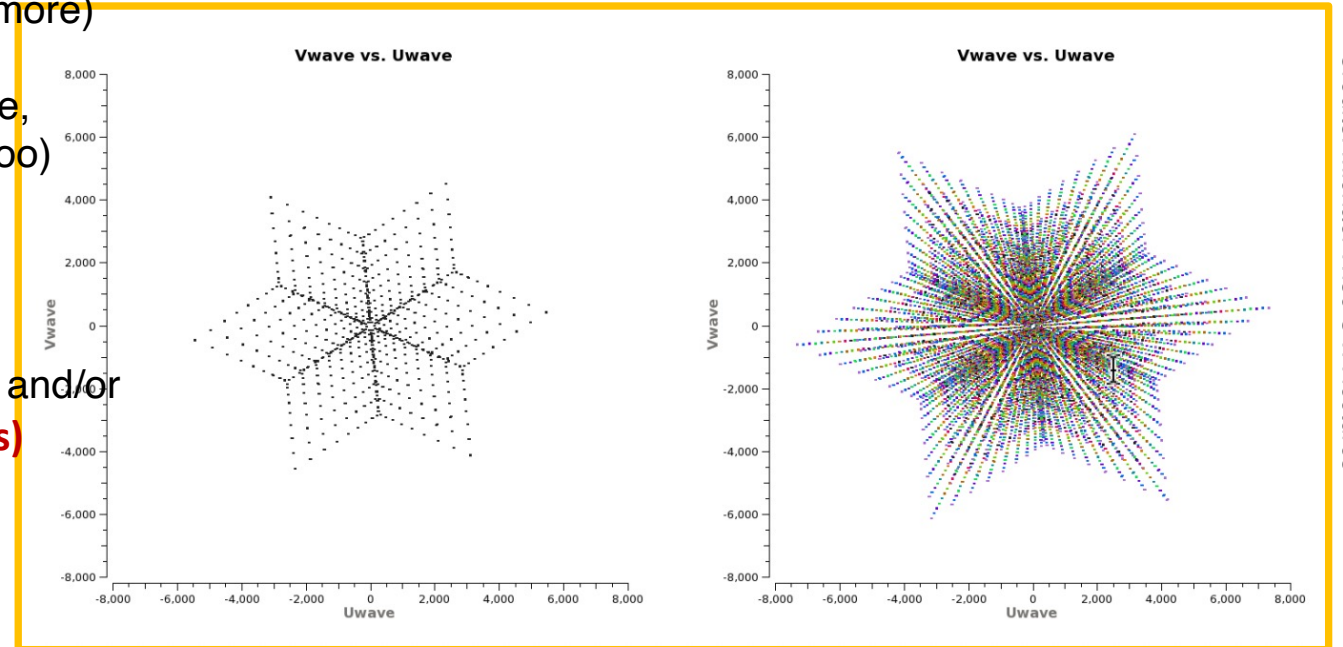
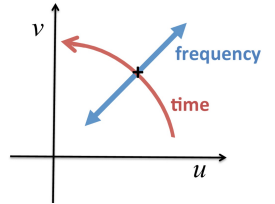
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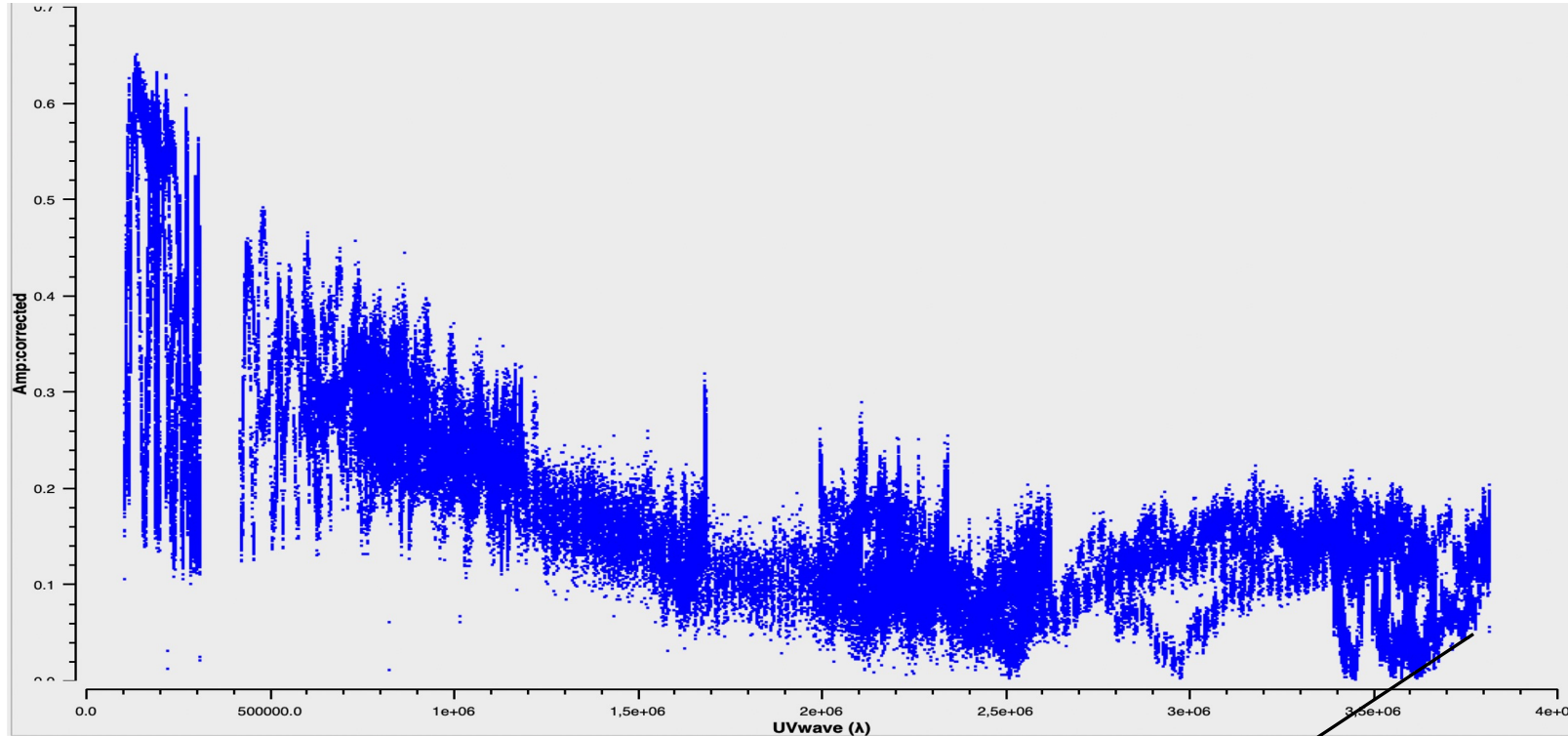
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Credits: Michael Wise UVA lectures

Determining imaging parameters

Imaging parameters: pixel size



Nyquist sampling theorem in astronomical terms

minimum number of pixels required to obtain a fit to a Gaussian must be at least 3 pixels

$$\text{cell} \approx \frac{180}{\pi N_s} \times \frac{1}{D_{\max} [\lambda]} \text{ [deg]}$$

N_s typically = 5-7
must not be smaller than 3!

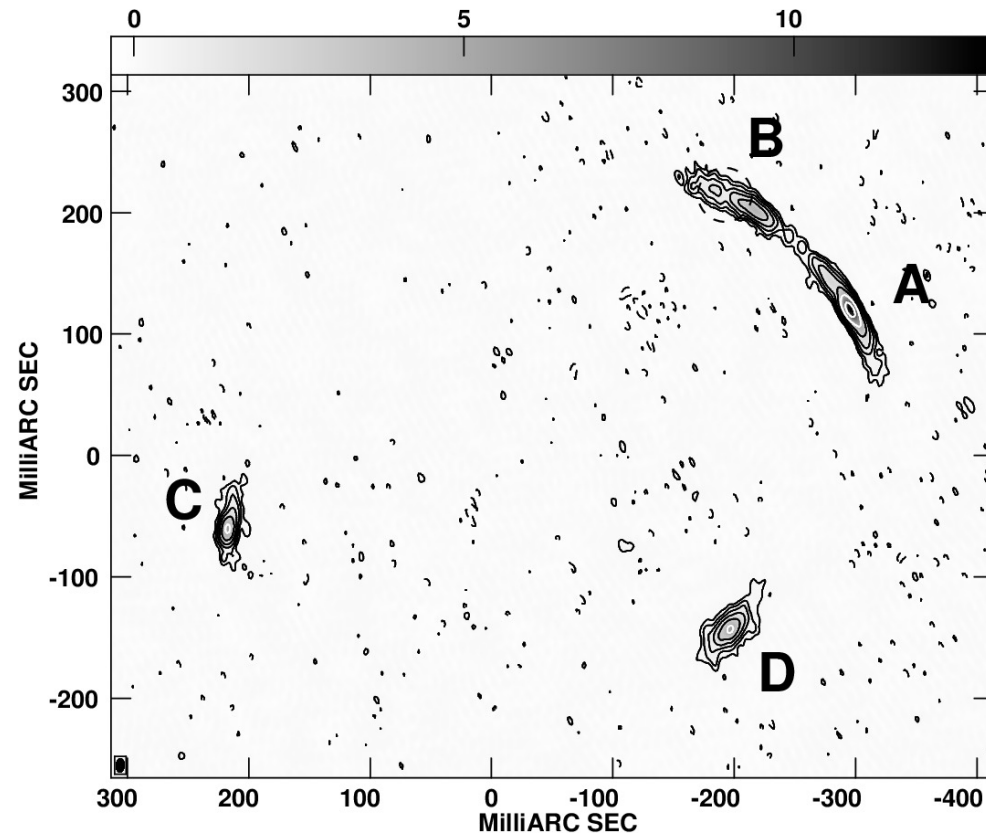
Imaging parameters: field of view / image size

The source size is typically much smaller than the entire Field-of-View (FoV), which corresponds approximately to the single-dish beam $\approx \lambda/D$ (homogeneous array)

But it is also limited by time and bandwidth smearing:
a typical FoV for VLBI is of the order of a few arcseconds

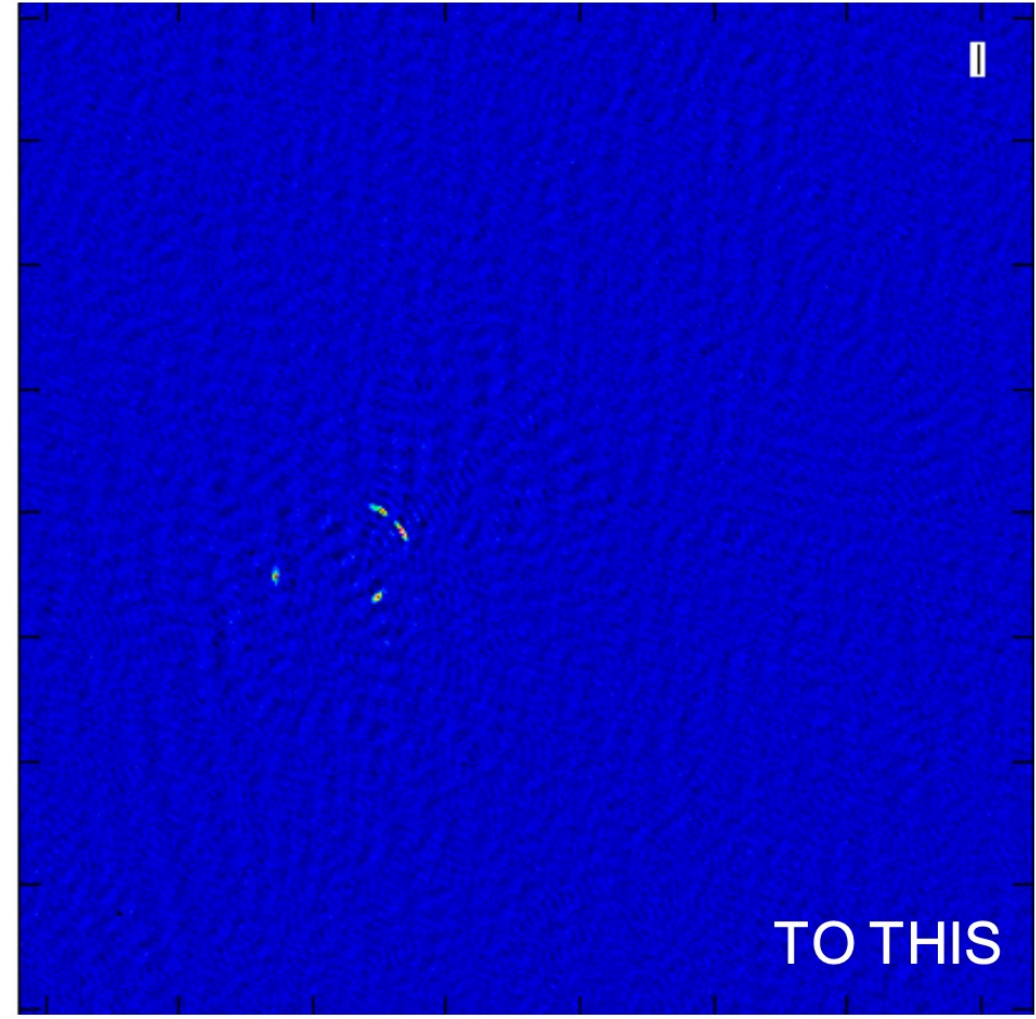
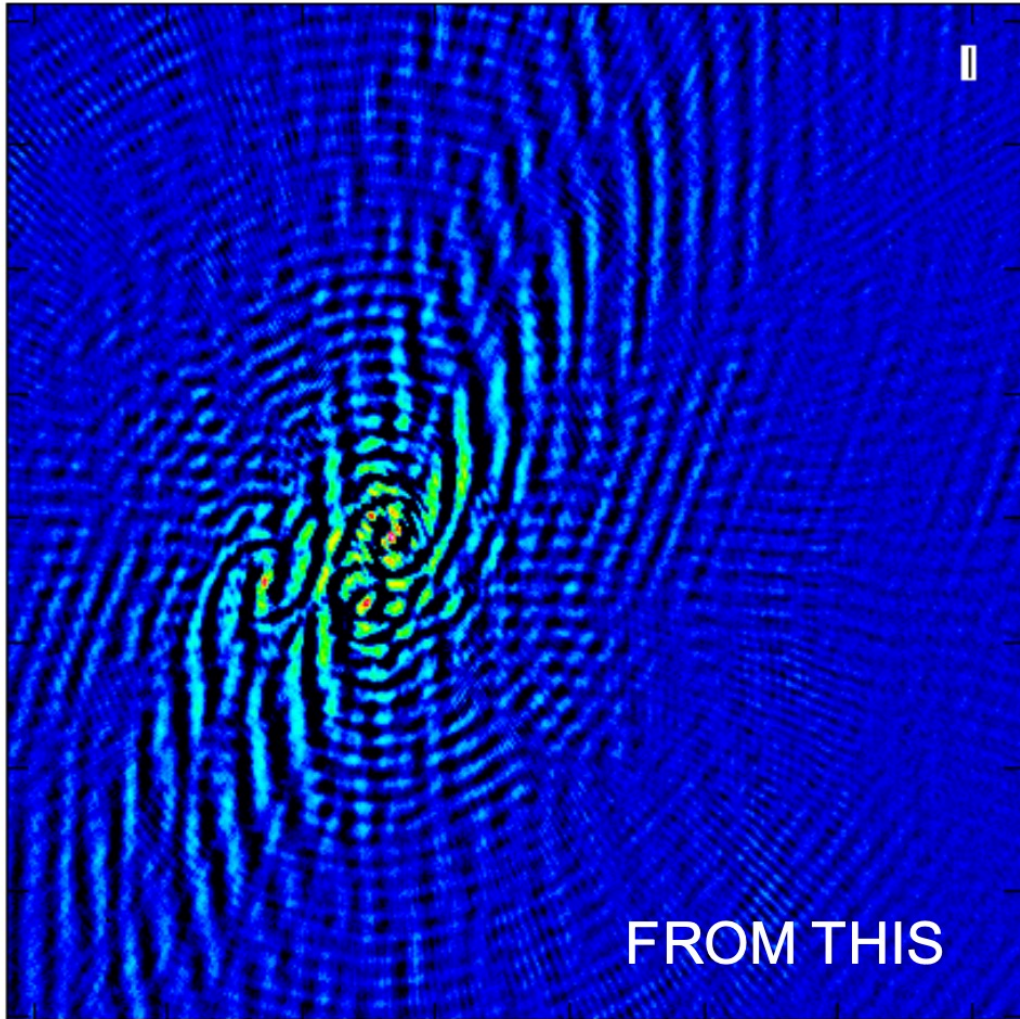
Also: it's always good to check
what is already known about your target!

For the test data for this CASA-VLBI workshop
the target is JVAS B0128+437
(L-band VLBA observations)



Biggs et al. 2004, MNRAS, 350, 949
and <https://arxiv.org/pdf/astro-ph/0412677.pdf> (Biggs+ 2004 EVN proceeding)

Let's create a CLEAN image



Imaging parameters in CASA: tclean

imagename

field

cell

imsize

deconvolver

niter

weighting

If you don't know what a
parameter means, its units...
just type
help tclean

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Size of image in pixels – typically power of two 2^n (128x128, 256x256, 512x512 etc.)

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deconvolver

CLEAN algorithm -- Clark or Hogbom algorithms are fine for starting

niter

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niter If = 0 computes the dirty image; if > 0 runs major and minor cycles (sub-parameter '*cycleniter*')

weighting

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niter If = 0 computes the dirty image; if > 0 runs major and minor cycles (sub-parameter '*cycleniter*')

weighting $V_{\text{obs}}(l,m) = V_{\text{true}}(u,v) S(u,v)$ $S(u,v)$ is 1 if there's a measurement and 0 elsewhere
so-called **natural weights**

Imaging parameters in CASA: a slide about weights

weighting

$V_k \rightarrow$ AMP(a_k) PHASE(ϕ_k) NOISE(σ_k) WEIGHT (w_k)

Better rms, worse beam

Natural

$$w_k = 1 / \sigma_k^2$$

«more weights on short baselines», best sensitivity (important for more extended structures) but poor beam shape with overemphasized sidelobes

Robust

(Briggs 1995)

$$w_k = 1 / (S^2 + \sigma_k^2)$$

$$S^2 = \frac{(5 \times 10^{-R})^2}{\bar{w}}$$

R = robustness (or robust factor) and it goes from -2 to 2
Average variance weighting factor over the grid cell in the image

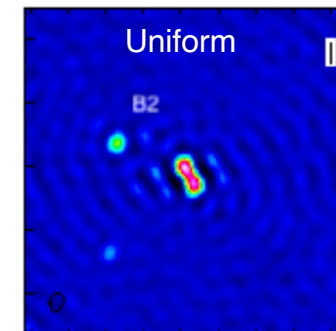
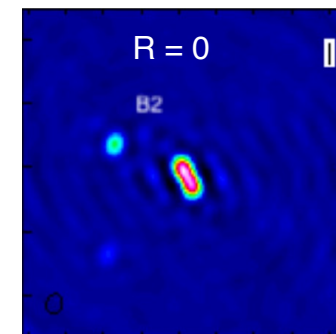
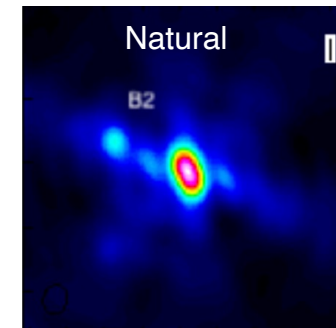
Uniform

Sampling density function

$$w_k = 1 / \varrho(u_k, v_k)$$

«more weights on long baselines», better resolution (tighter main lobe) and lower sidelobes

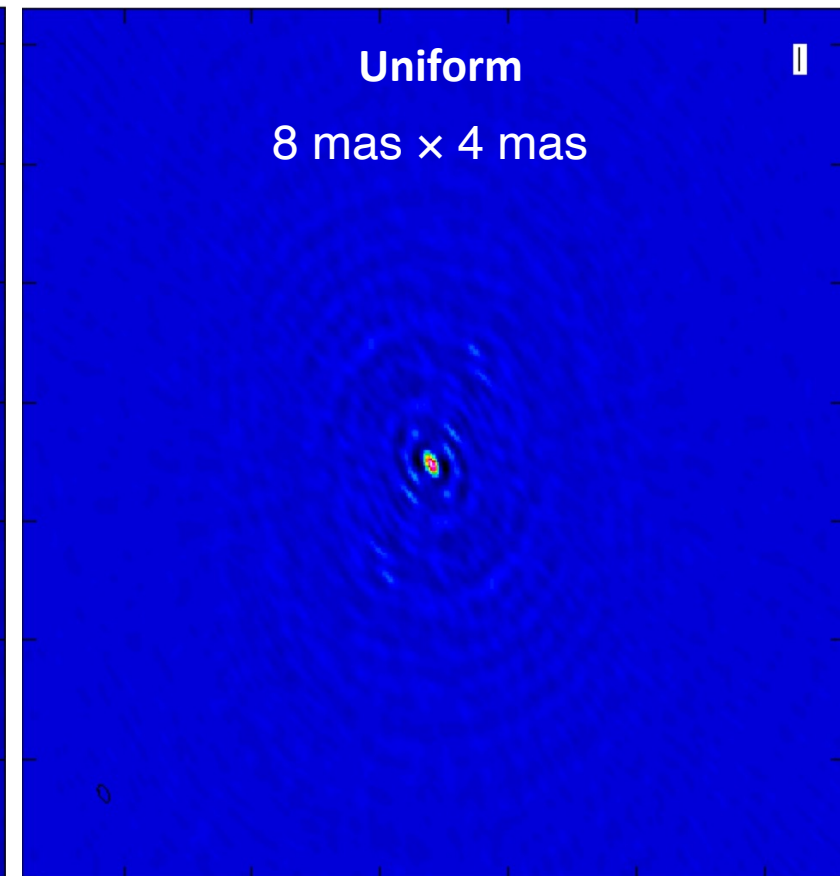
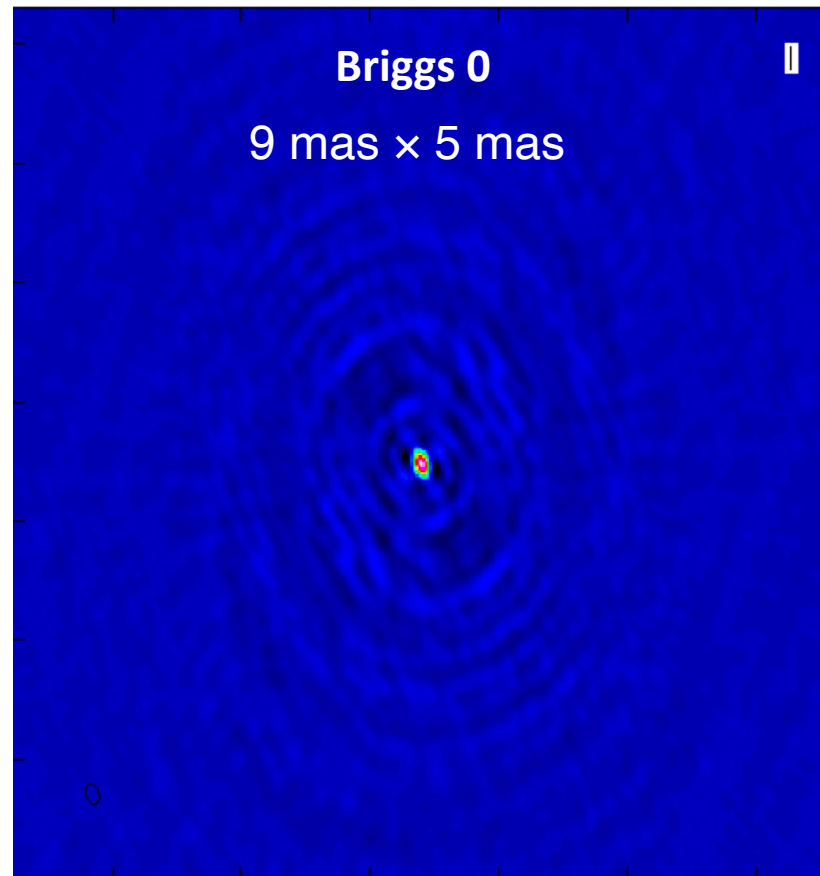
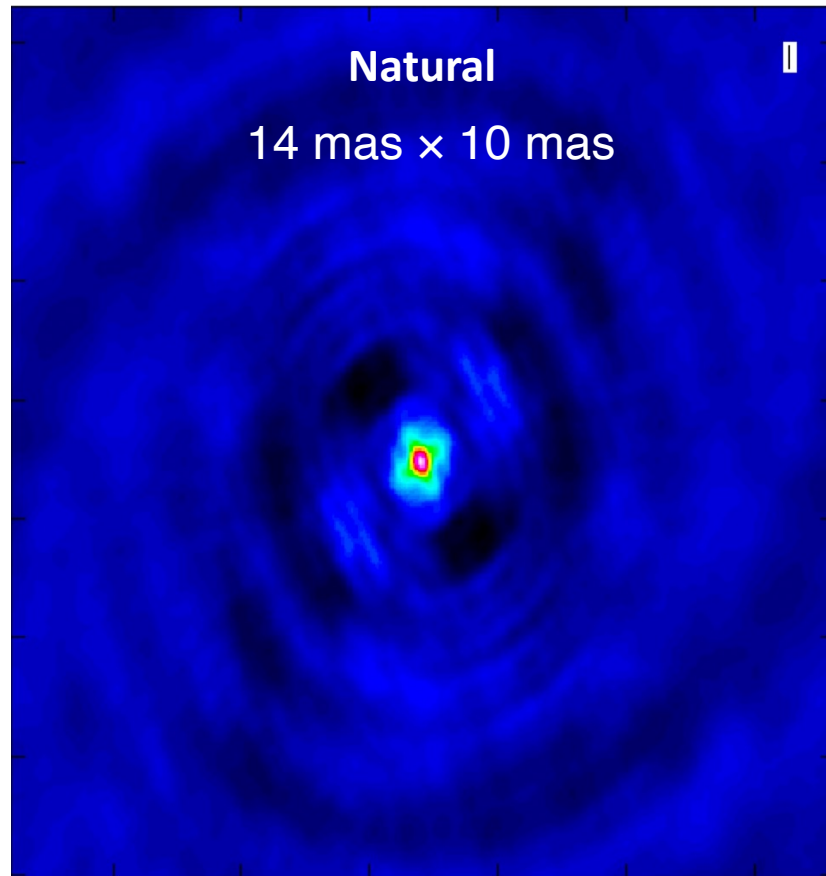
Better beam, worse rms



Imaging parameters in CASA: a slide about weights

weighting

Different weights lead to different PSF (Dirty Beam) hence we should adjust the cellsize according to the chosen weights



Imaging in CASA: interactive cleaning

Once you are happy with the choice of parameters ... go `tclean`

The screenshot shows the CASA `tclean` interactive cleaning interface. The top toolbar contains icons for panning, zooming, and other navigation functions. Below the toolbar are several control panels:

- Panning and zoom controls:** Includes buttons for pan, zoom, and other navigation actions.
- Generate regions/masks:** Includes buttons for adding, erasing, and generating regions and masks.
- Final residual brightness threshold to CLEAN to:** A panel with input fields for `max cycleniter` (set to 100), `iterations left` (set to 1000), `threshold` (set to 0Jy), and `cyclethreshold` (set to 0.750603Jy).
- Animators:** A panel for controlling the animation of the cleaning process, including a `Rate` slider (set to 10) and a `Jump` dropdown (set to 0 2).
- Cursors:** A panel for displaying the current cursor position and information about the currently opened images.

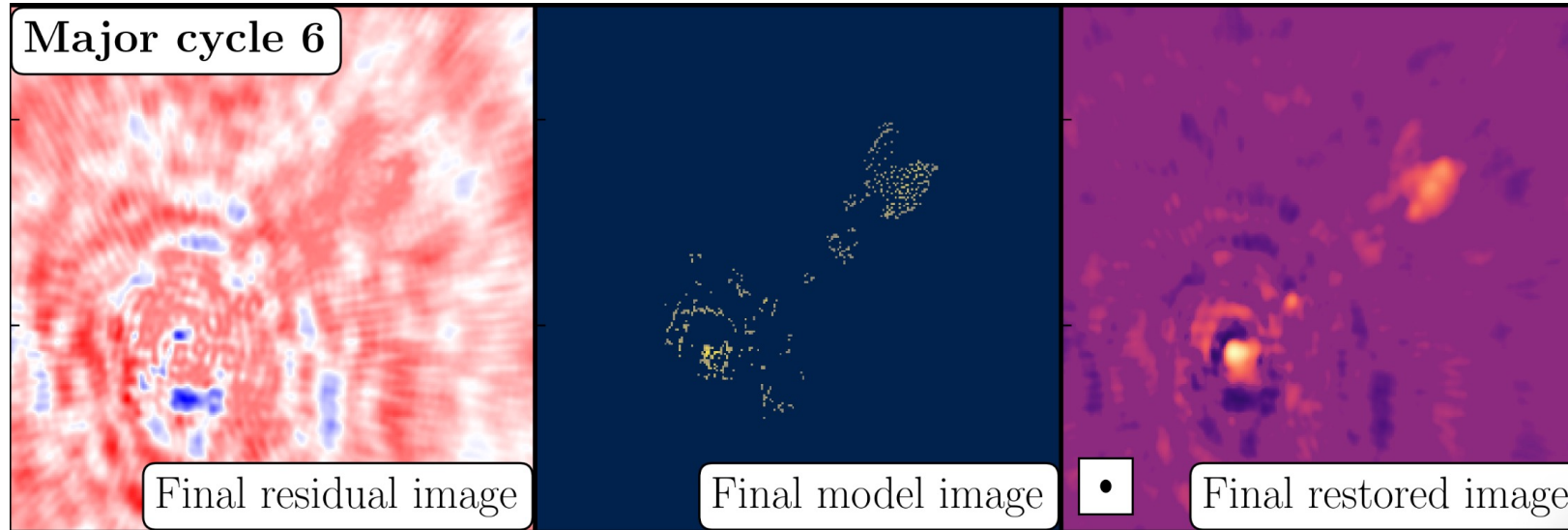
The main display area shows a residual image titled `1848+283_p_only.residual-raster`. The image is a color-coded map of the residual flux, with a central bright spot and surrounding structure. The axes are labeled `J2000 Declination` and `J2000 Right Ascension`.

Annotations and red arrows point to various features:

- Iterations left (from setting niter parameter):** Points to the `iterations left` input field.
- Max number of CLEANs per major cycle:** Points to the `max cycleniter` input field.
- Continue CLEAN indefinitely until threshold reached or iterations left = 0:** Points to the `Next Action:` button.
- Continue CLEAN for only one major cycle i.e. max cycleniter or cyclethreshold reached. This returns interactive screen to modify masks. Complete major cycle if residual image reaches this peak brightness:** Points to the `cyclethreshold` input field.
- Animator for when multiple images to be deconvolved at one i.e. spectral line cubes or polarizations:** Points to the `Animators` panel.
- Currently opened images and info on where the mouse pointer is:** Points to the `Cursors` panel.
- Residual image (dirty image minus cleaned flux):** Points to the main display area.

Imaging in CASA: interactive cleaning

Continue the cleaning process until your image looks like noise



ALWAYS take a look at the logger!

E.g., Flux density in the model should increase

Major Cycle 1	model=0->0.177024,
Major Cycle 2	model=0.177024->0.381401,
Major Cycle 3	model=0.381401->0.504565

If it keeps decreasing: **STOP CLEANING!**

model=0.686961->0.683864

Imaging in CASA: output of tclean

.image → Final cleaned map

.tt0, .tt1 ... → Suffix to indicate Taylor terms for multi-term wideband imaging if «mtmfs» is used as deconvolver

.alpha and .alpha.error → spectral index and its error map

.mask → mask used (clean boxes)

.model → clean components

.psf (for tt0, tt1,...) → dirty beam

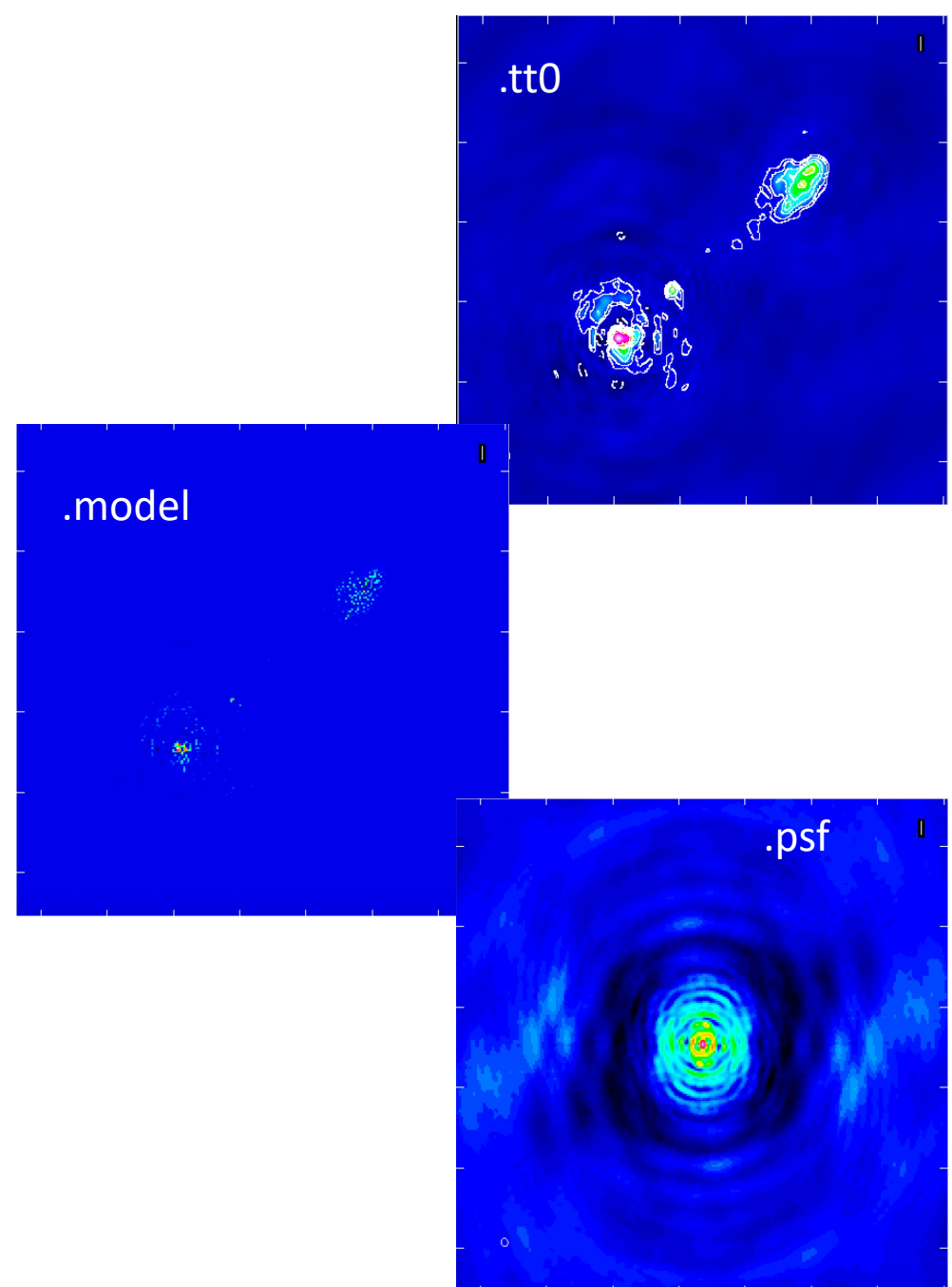
.pb → primary beam

.residual → residual image (data – model)

.sumwt → sum of the weights

Other details can be found here

<https://casa.nrao.edu/docs/taskref/tclean-task.html>



Measuring image properties

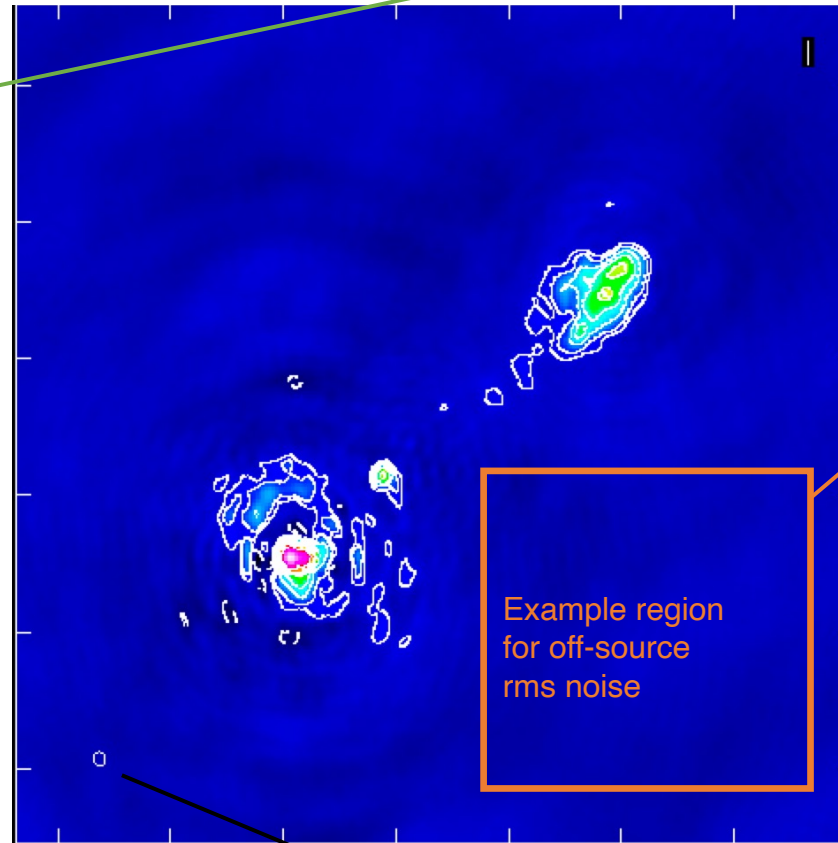
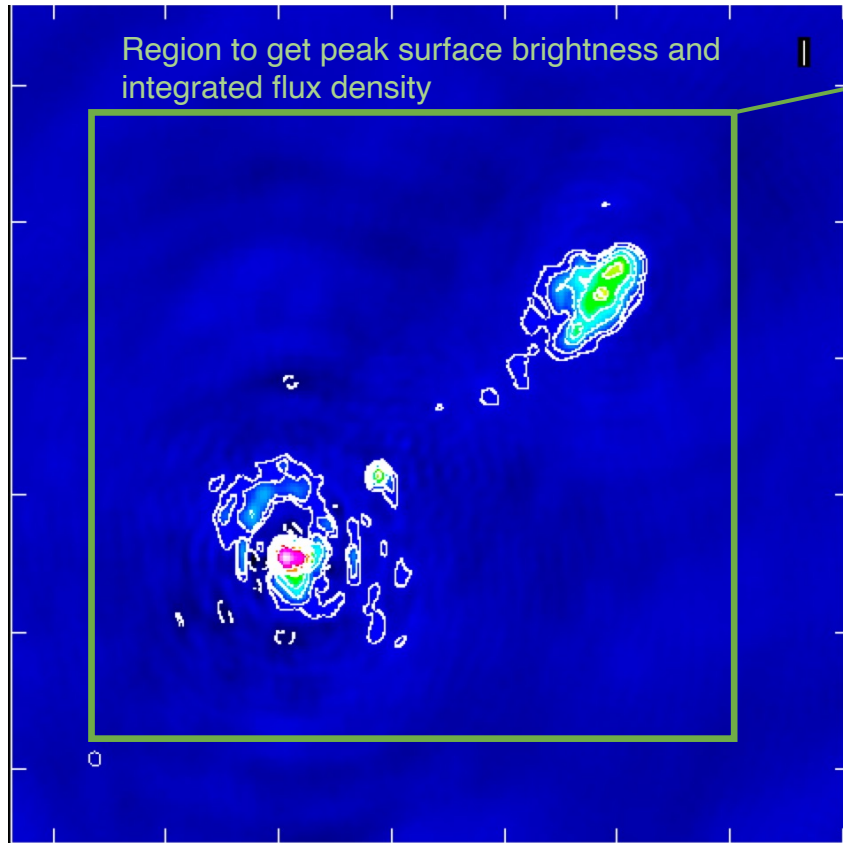
See also https://www.jb.man.ac.uk/DARA/ERIS22/imaging.html#image_properties

Imaging in CASA: measuring image properties

Directly using IMVIEW
Or using the task IMSTAT

The peak surface brightness is in Jy/beam

While the integrated flux density is in Jy



The rms of the image is in Jy / beam

What is the dynamic range (peak / off-source rms) of your image ???

Restoring beam (CLEAN beam)

Imaging in CASA: measuring image quality

- **Off-source rms** noise close to theoretical noise
- **Dynamic range** (peak / off-source rms) -- typical (good) values 10^2 - 10^6
- «**Fidelity**» – difference with an input model (need *a priori* info)
- Off-source rms **noise structure quite uniform**, close to a Gaussian random field («no stripes»): check for any phase and amplitude errors (see previous slides)
- any «weird» structure might be a symptom that something went wrong (at the deconvolution stage and/or during calibration)

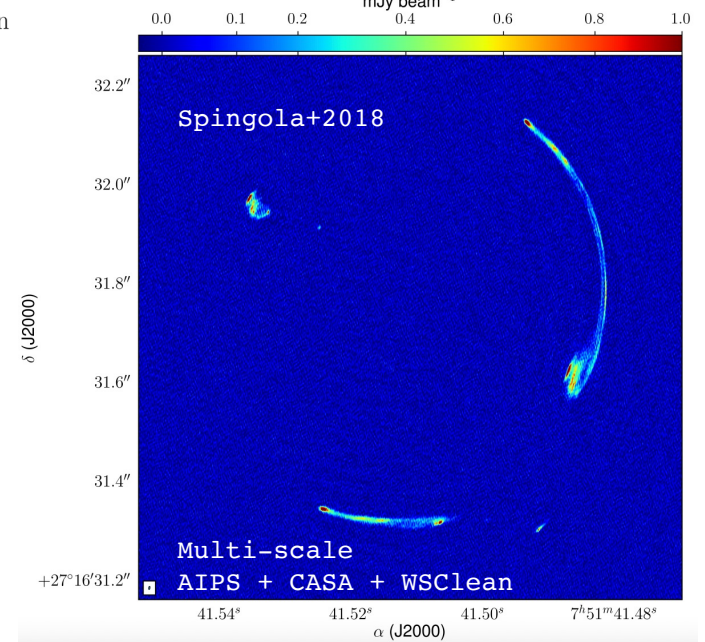
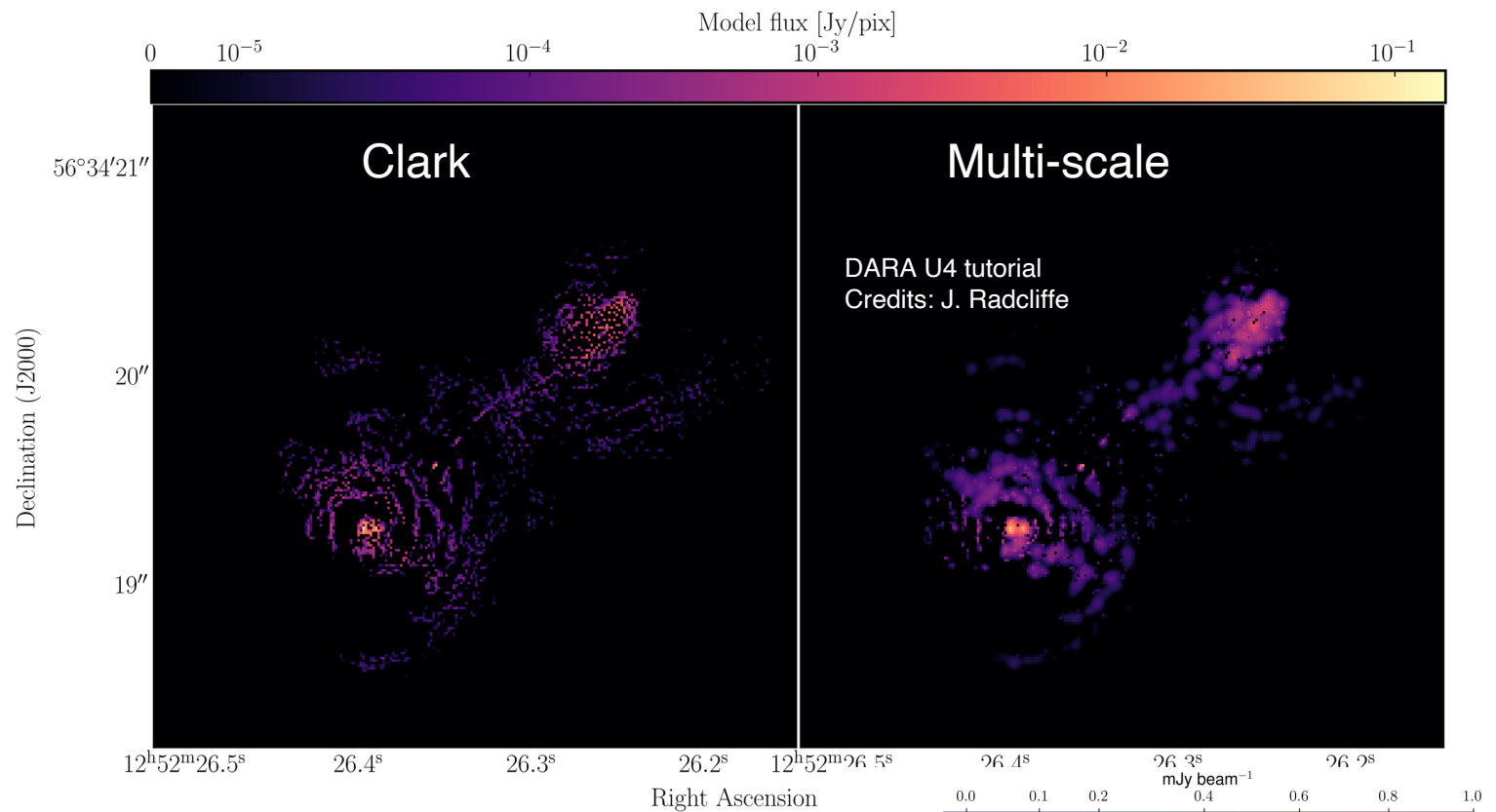
Multi-scale cleaning

Deconvolver: `mtmfs`

Default: clean components are *delta functions* – it does not work very well for extended sources!

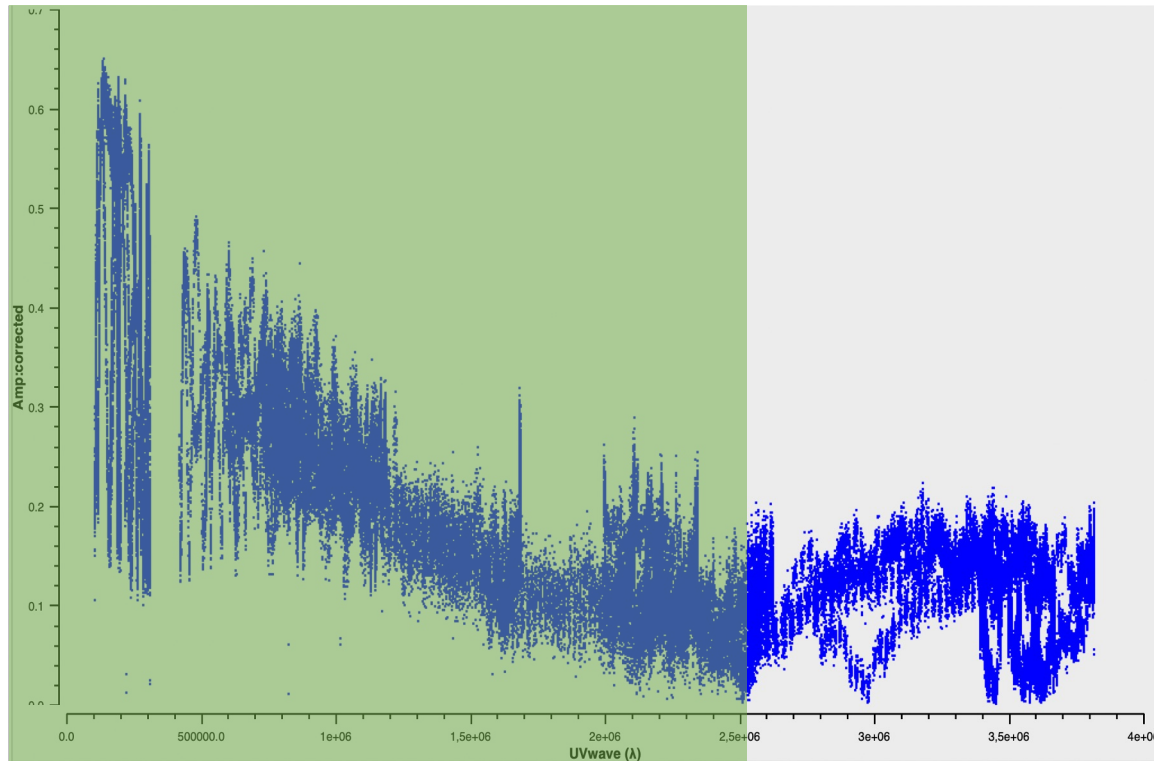
Multi-scale = clean components that can be «extended» (e.g., Gaussian functions) on different scales

See Cornwell 2008; Rau & Cornwell 2011; Ofringa & Smirnov 2016



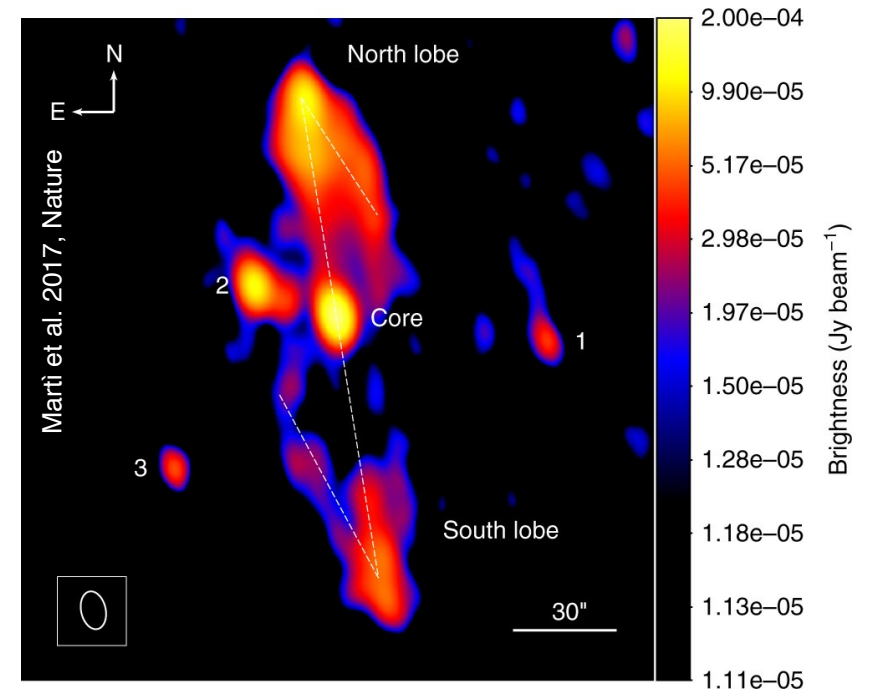
Tapering

It's like smoothing the data in the image plane by convolving with a Gaussian PSF with FWHM corresponding to the chosen $k\lambda$ to some extent increases sensitivity to medium-large angular scale structures



`uvtaper = ['2.5 Mlambda'] (tclean)`

circular taper of HWHM=25 Mega-lambda



Tapering on 25 $k\lambda$ was applied to the interferometric visibilities to better enhance the extended emission

References

Chapters 7 and 8 of «Synthesis imaging in radio astronomy II» (Edited by Taylor Carilli and Perley)

Campbell 2019 http://old.evlbi.org/user_guide/fov/fovSFXC.pdf

Interferometry and Synthesis in radio imaging (Thompson, Moran and Swenson) <https://link.springer.com/book/10.1007/978-3-319-44431-4>

Previous ERIS imaging lectures can be found here <https://www.astron.nl/events/eris-2022/>

Lecture on imaging by Michael Wise https://www.astron.nl/astrowiki/lib/exe/fetch.php?media=ra_uva:ra_uva_lecture8.pdf

Images in the first slide: Spingola+2018, Giovannini+2018(NatAs), Boccardi+2016, Hartley+2019, McKean+2011, Giroletti+2020, Radcliffe+2016, Johnston+2020, Kellermann+2007

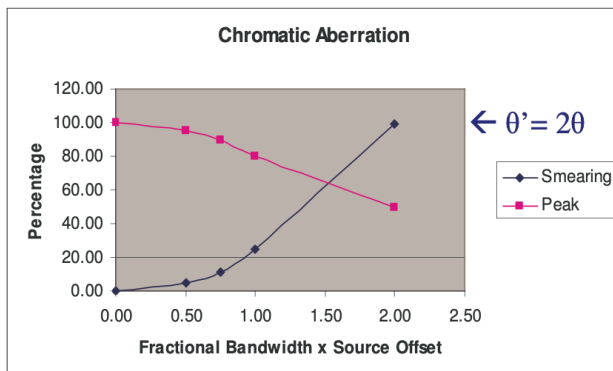
Extra slide: bandwidth and time smearing

Bandwidth smearing

- Bandwidth smearing (chromatic aberration) will produce radial smearing and reduction in source peak

- Parameterized by the product of the fractional bandwidth and the source offset in synthesised beamwidths

$$\delta\nu/\nu_0 \times \theta/\theta_{HPBW}$$



- Can be alleviated by observing and imaging in spectral line mode with many narrow frequency channels gridded separately prior to Fourier inversion – reduces $\delta\nu$

- Detailed form of response depends on individual channel bandpass shapes.

Time smearing

- Time-average smearing (de-correlation) will produce tangential smearing

- In general cannot be easily parameterized. At Declination= $+90^\circ$ a simple case exists where the effects can be parameterized by the equivalent product:

$$\omega_e \delta t_{int} \times \theta/\theta_{HPBW}$$

Where ω_e is the Earth's angular rotation rate and δt_{int} is the integration time interval in the dataset

- For other Declinations the effects are more complicated. However they can be alleviated by ensuring that δt_{int} is small enough such that there at least 4 samples per turn assuming a maximum rate of θ/θ_{HPBW} turns in 6 hours