

# Imaging and Image Analysis



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# Outline

## - Synthesis imaging as implemented in CASA ( tasks : **clean**, **clean** )

<https://casa.nrao.edu/casadocs/casa-5.1.0/synthesis-imaging>

- Major and minor cycles
- Gridding algorithms
- Deconvolution algorithms
- Types of Images
- Iteration control
- Masks for deconvolution
- Runtime interactivity
- Factors affecting computing cost
- Task interface

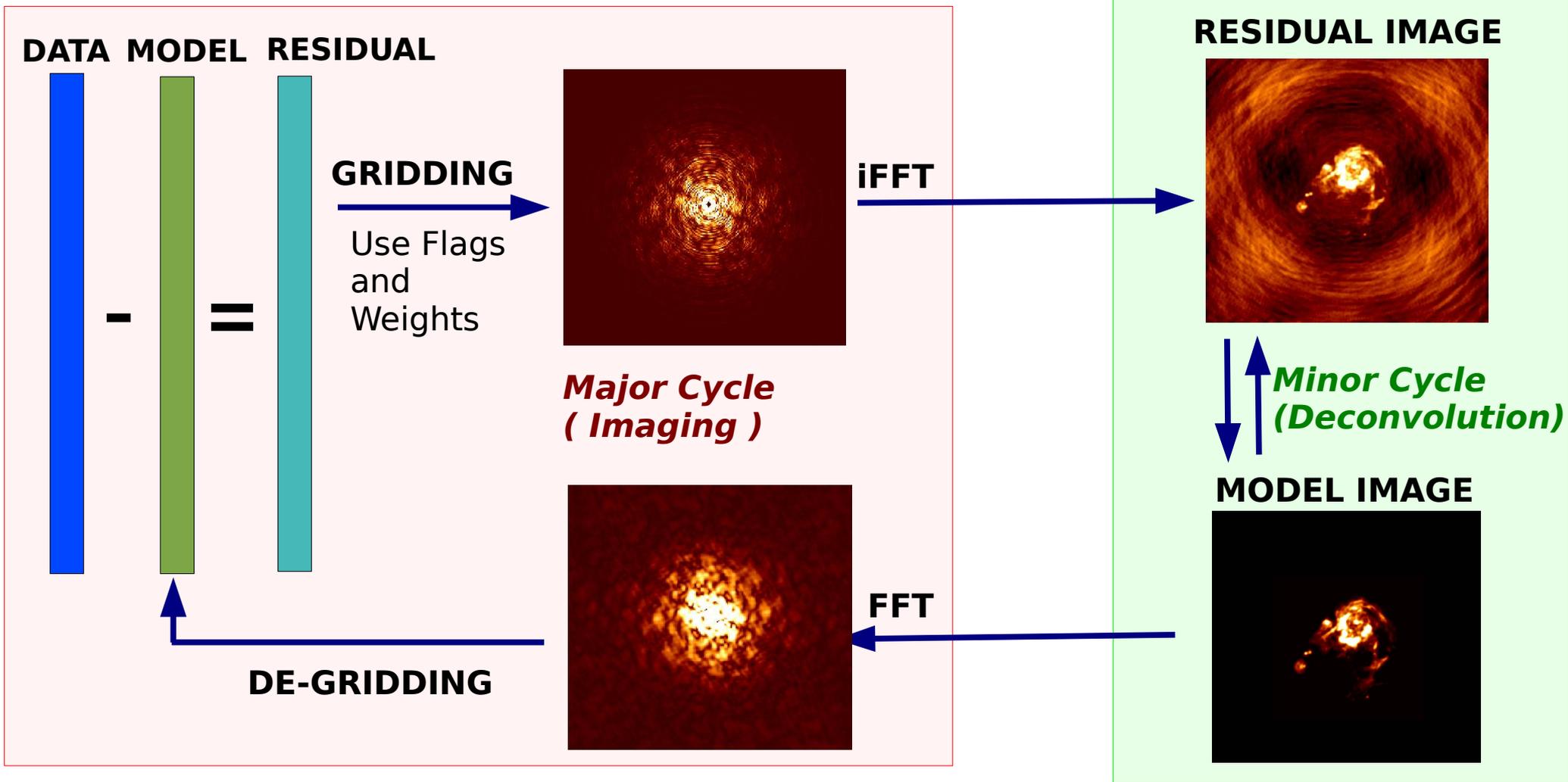
- clean** - Older task
  - Development frozen
  - Task retained for backward compatibility of scripts
  - Tools retained for various utility methods
- tclean** - Newer task
  - Currently developed/used
  - Uses refactored imaging module
  - All major features of **clean** plus several more
  - Used by pipelines

## - Image Viewing and Analysis ( task : **viewer** , 28 **image analysis** tasks)

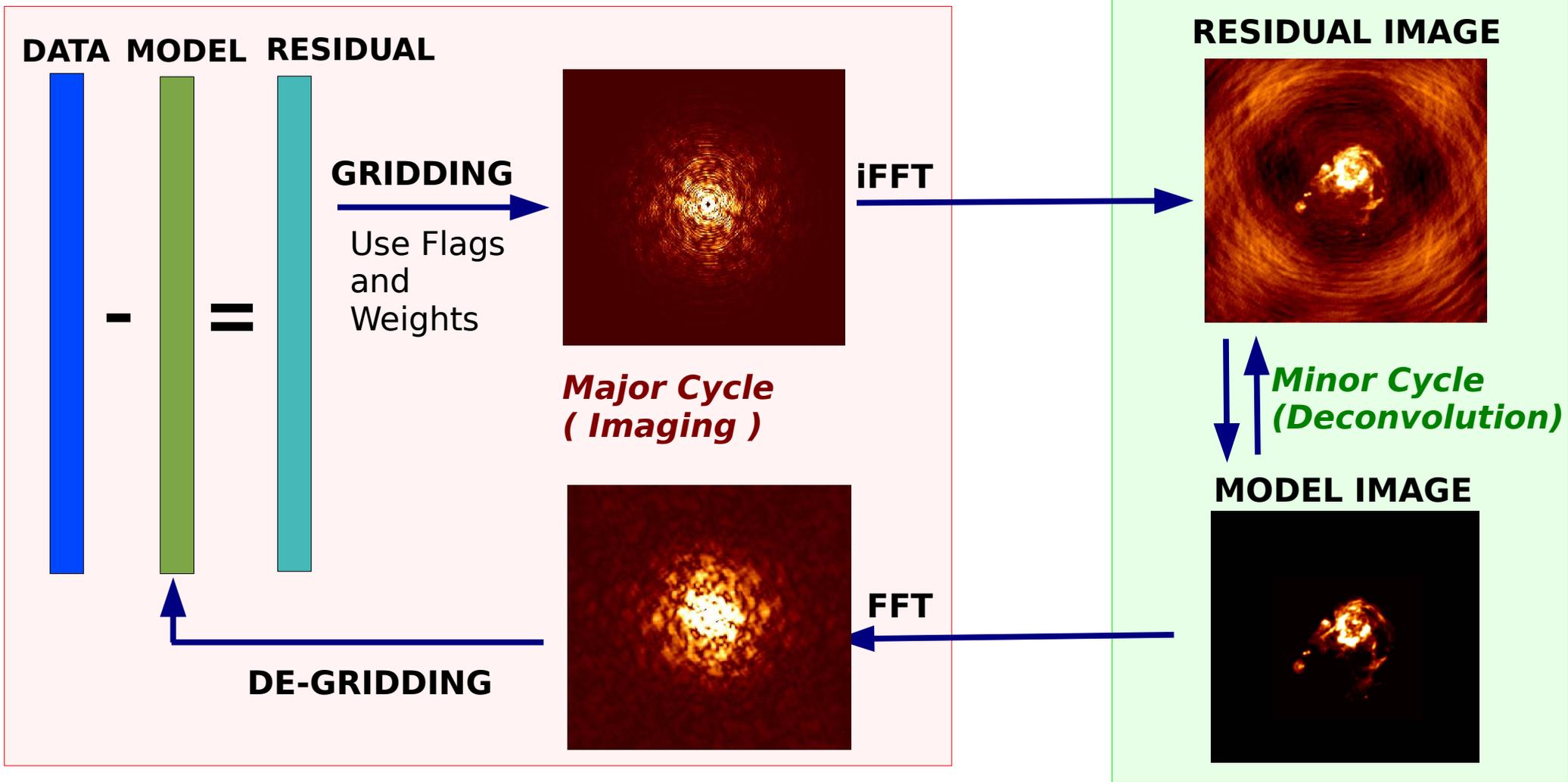
<https://casa.nrao.edu/casadocs/casa-5.1.0/image-cube-visualization>

<https://casa.nrao.edu/casadocs/casa-5.1.0/image-analysis>

# Iterative $\chi^2$ minimization – Major and Minor Cycles



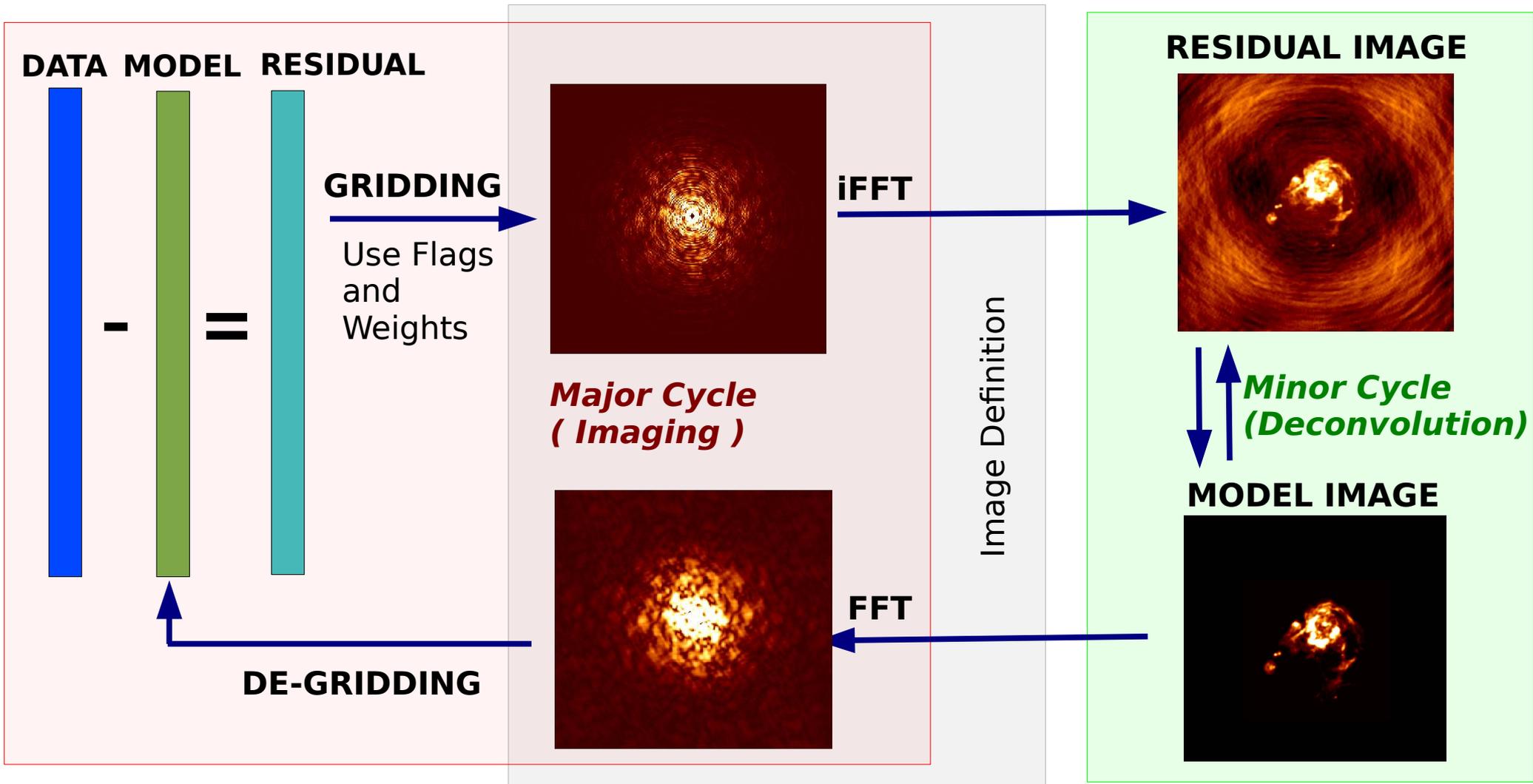
# Iterative $\chi^2$ minimization – Major and Minor Cycles



**Standard gridding,  
W-Projection,  
(WB)-A-Projection,  
Joint Mosaics**

**Clean ( Hogbom,  
Clark, MultiScale,  
MultiTerm, etc... )**

# Iterative $\chi^2$ minimization – Major and Minor Cycles



**Standard gridding,  
W-Projection,  
(WB)-A-Projection,  
Joint Mosaics**

**Cube, MFS,  
MT-MFS, Faceting,  
Stokes, Multi-Field**

**Clean ( Hogbom,  
Clark, MultiScale,  
MultiTerm, etc... )**

# Imaging Algorithms ( Weighting + Gridding )

Image = Weighted average of the data

$$= \frac{\text{sum of ( weight x data )}}{\text{sum of ( weight )}}$$

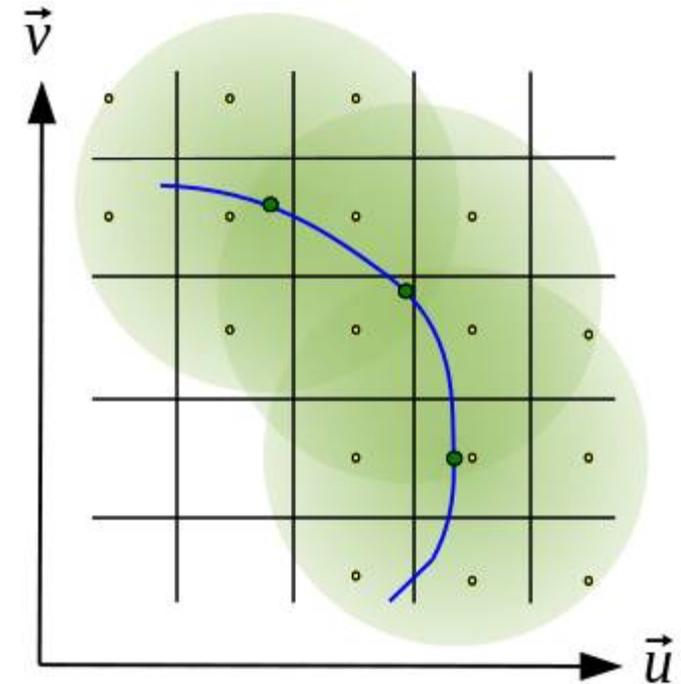
**Gridding** : Accumulate weighted visibilities on a regular grid : convolutional resampling  
( accumulate weights too, for PSF & PB )

Choose different gridding convolution functions  
=> handle wide-field effects (next slide)

**iFT** : Fourier Transform to image domain

**Normalization** : Divide by sum-of-weights  
( and PB, for some gridding algorithms)

- flat noise, flat-sky, pb-square



## Weights

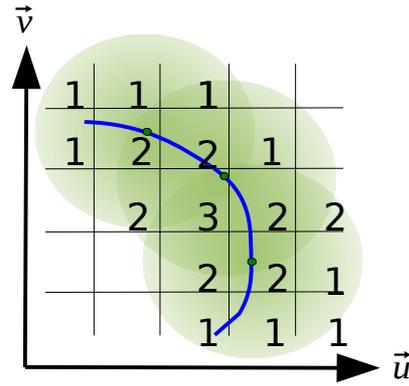
- Data Weights
- Imaging Weights
- Natural
- Uniform
- Briggs

# Imaging Algorithms ( Weighting + Gridding )

## Weighting Schemes :

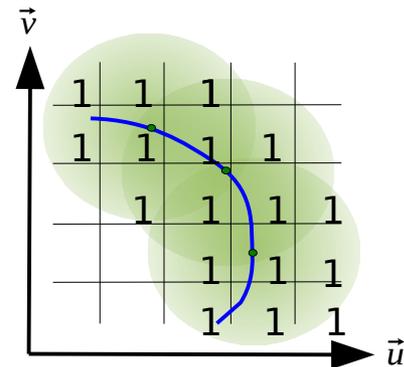
### Natural

- Highest sensitivity
- PSF can be wide



### Uniform

- Narrow PSF
- Suppressed sensitivity

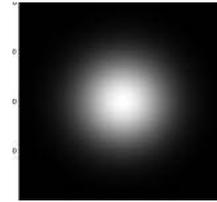


**Briggs/Robust** : Smoothly vary between natural and uniform (robust 0.5 == AIPS robust 0 )

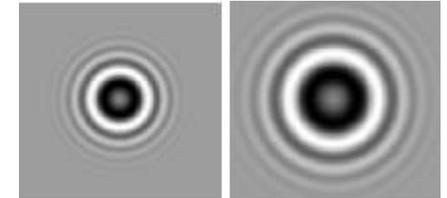
**UV-Taper** : Emphasize large scales

## Gridding : Choice of convolution function

**Standard Imaging** :  
Prolate Spheroidal

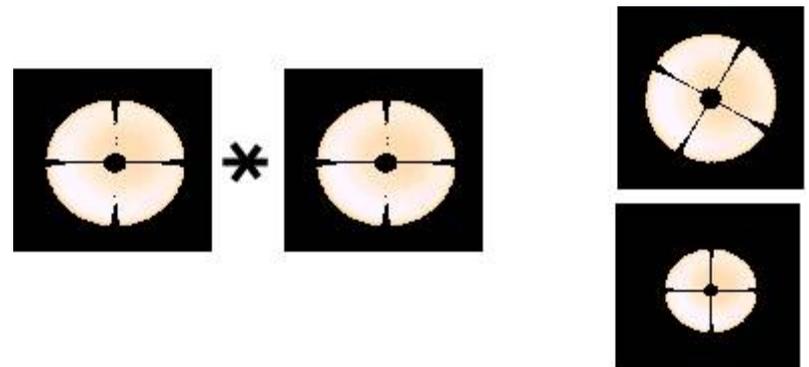


**W-Projection** :  
FT of a Fresnel kernel



**A-Projection** :

Convolutions of Aperture Illumination Funcs + phase gradients for joint **mosaics**



“A”, “W” kernels can change with baseline, time and/or frequency.

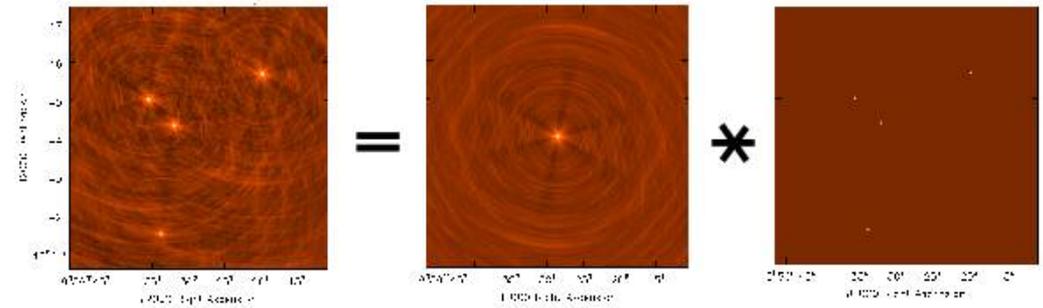
See **Wideband/Widefield** lecture for details

# Deconvolution Algorithms

**Point Source CLEAN** : Sky model is a set of delta functions. Remove effect of PSF

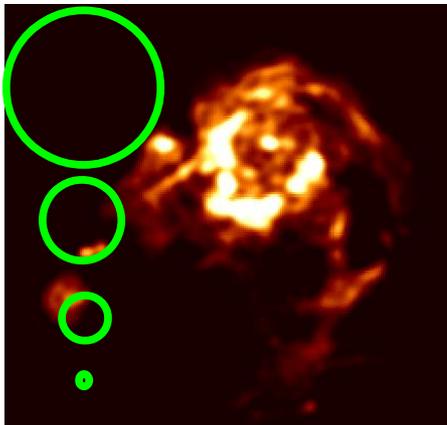
- **Hogbom** (use PSF of same size as image, clean until image edge)
- **Clark / ClarkStokes** (use patch PSF, intermediate model subtraction on the gridded uv-plane )

Convolution Equation ==> Deconvolution



**Multi-Term CLEAN** : Joint deconvolution of a sky model described as a series of basis functions

- **MS-Clean** (sky model is a linear combination of 'blobs' of different sizes)



- **MT-MFS** ( wideband sky model is a Taylor series expansion of Intensity vs Frequency )

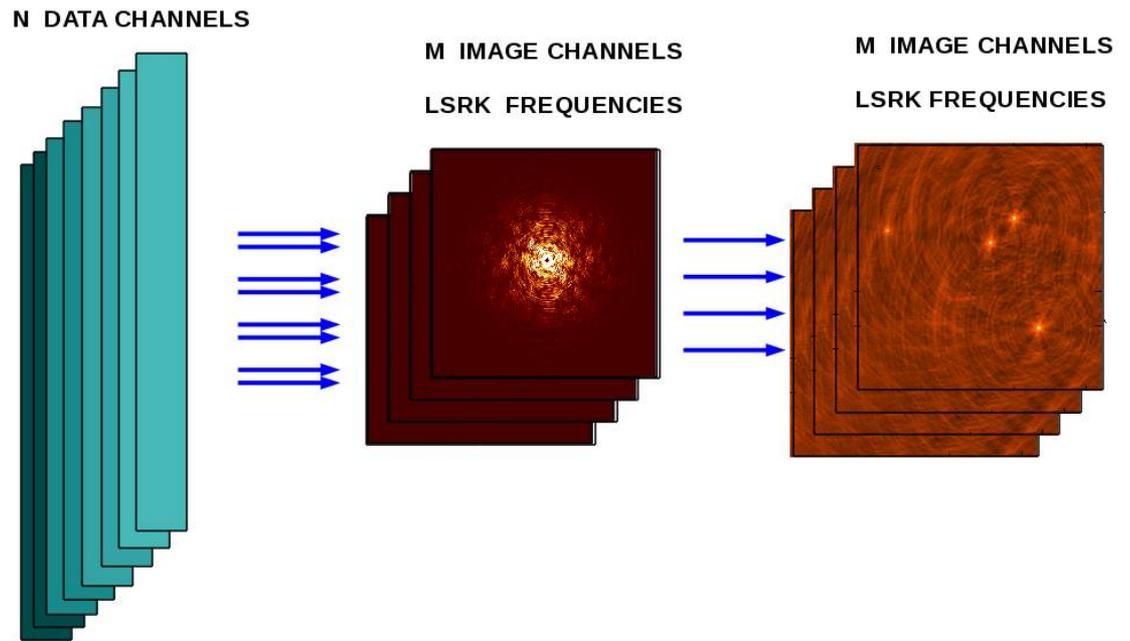
- Solve for intensity and spectrum together
- Includes multi-scale

( Details : [CASADocs](#) and the [Wideband/widefield lecture](#) )

# Image Definition : Spectral Cubes

`specmode = 'cube'`

- N data channels are mapped to M image channels (with binning/interpolation)
- Image coordinates defined by the user : start, width, nchan, outframe ( channel, frequency, velocity )
- Image coordinate system is internally stored in LSRK frame, with a conversion layer to allow relabeling to outframe for display/analysis



- All gridding/imaging is done in the LSRK frame with on-the-fly conversions to LSRK (i.e. no cvel needed)

`specmode='cubedata'` in tclean

- No internal conversion to LSRK.
- Data channels map to image channels (with only binning/interpolation)

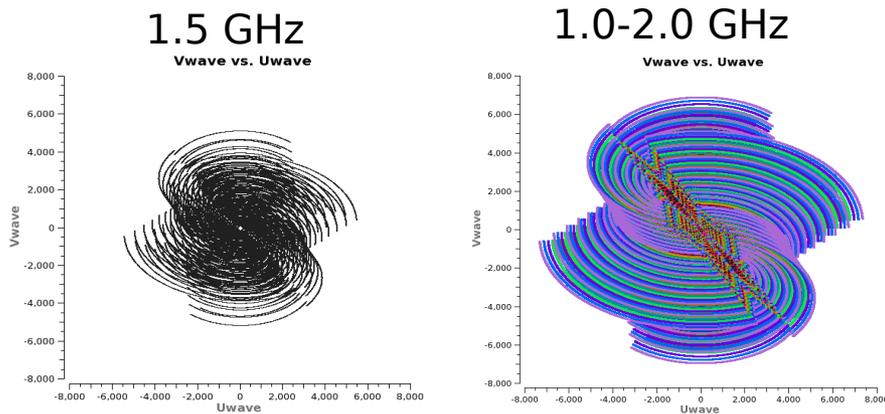
(Soon to come : `specmode = 'cubesrc'` in tclean  
to track moving sources via ephemeris tables)

# Image Definition : Continuum Imaging

specmode = 'mfs', 'cont'

- Data from all channels are gridded onto a single uv-grid, using the appropriate u,v,w coordinates

## - Multi-Frequency-Synthesis

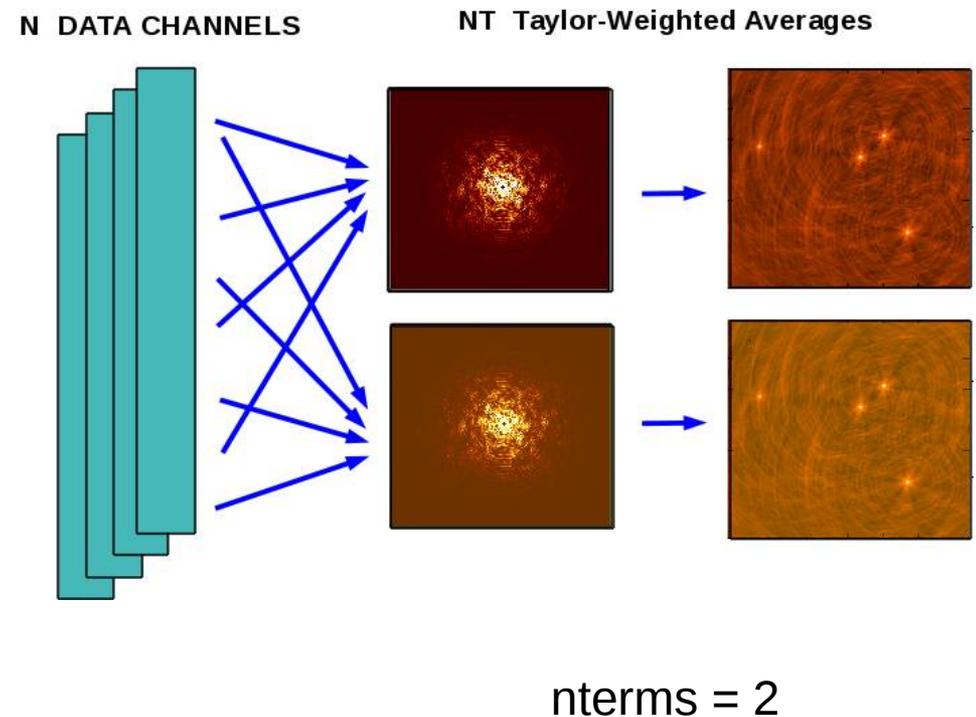
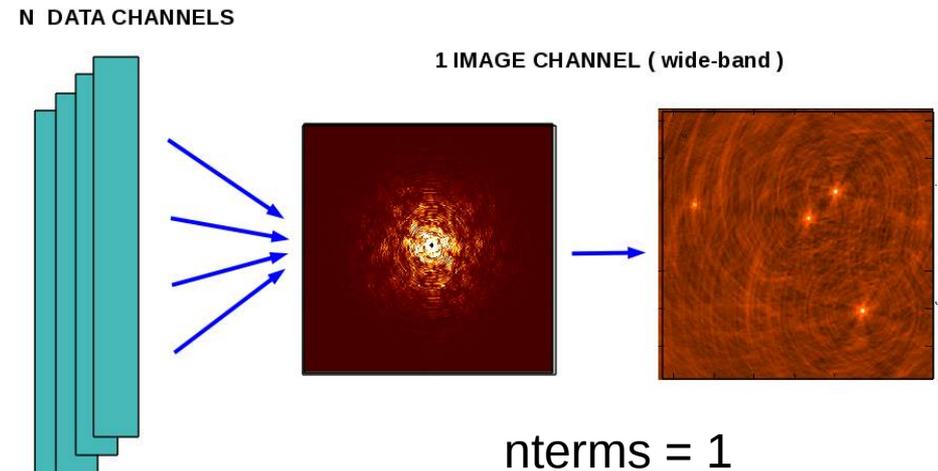


- nterms = 1 (flat spectrum assumption)

- nterms > 1 (Taylor polynomial spectrum)

- Major cycle : nterms Taylor-weighted averages of across frequency

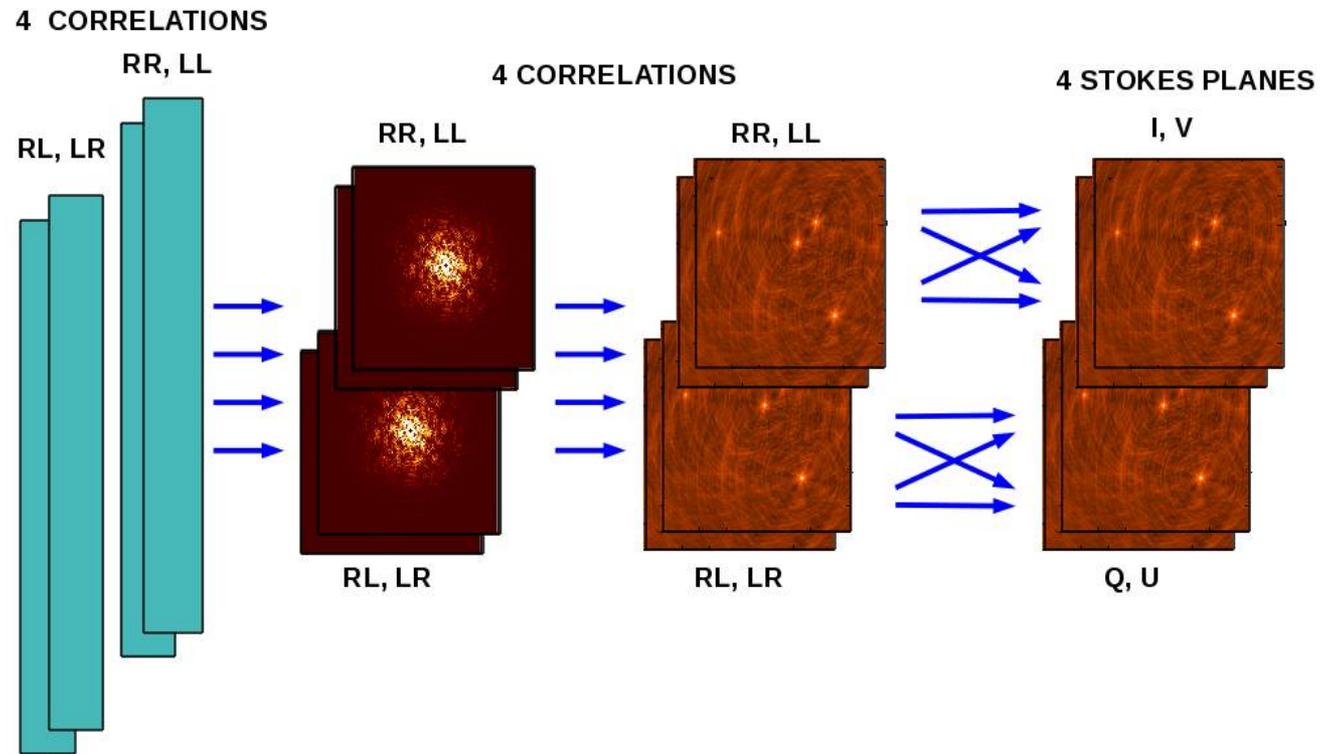
- Minor cycle : Solve for nterms coefficients



# Image Definition : Correlation and Stokes planes

## Polarization image planes

- Data are gridded and iFT  
In correlation basis
- Convert from correlation  
to Stokes basis
- Normalization  
( in Stokes basis )



Users can choose to make images of

R/L => I, Q, U, V, IV, QU, IQUV, R, LL, LR, RL, RRLL, RLLR, 'all'

X/Y => I, Q, U, V, IQ, UV, IQUV, XX, YY, XY, YX, XXYY, XYYX, 'all'

( Soon to come :

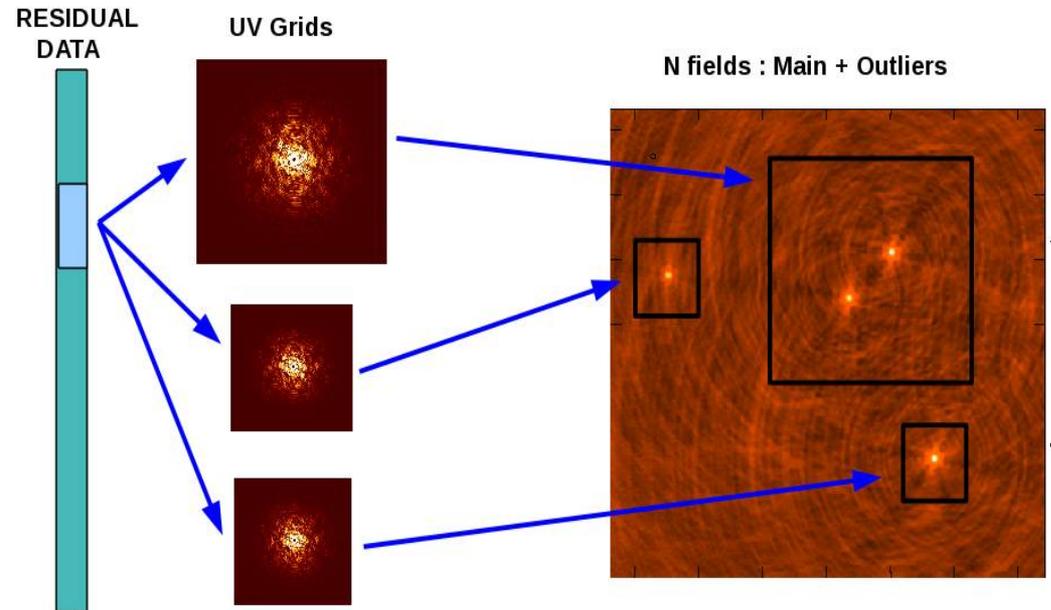
For Stokes I, use data even if some correlation pairs are flagged – “pseudo-I” )

# Image Definition : Multi-field and Facets

## Image partitioning : Same data, multiple images

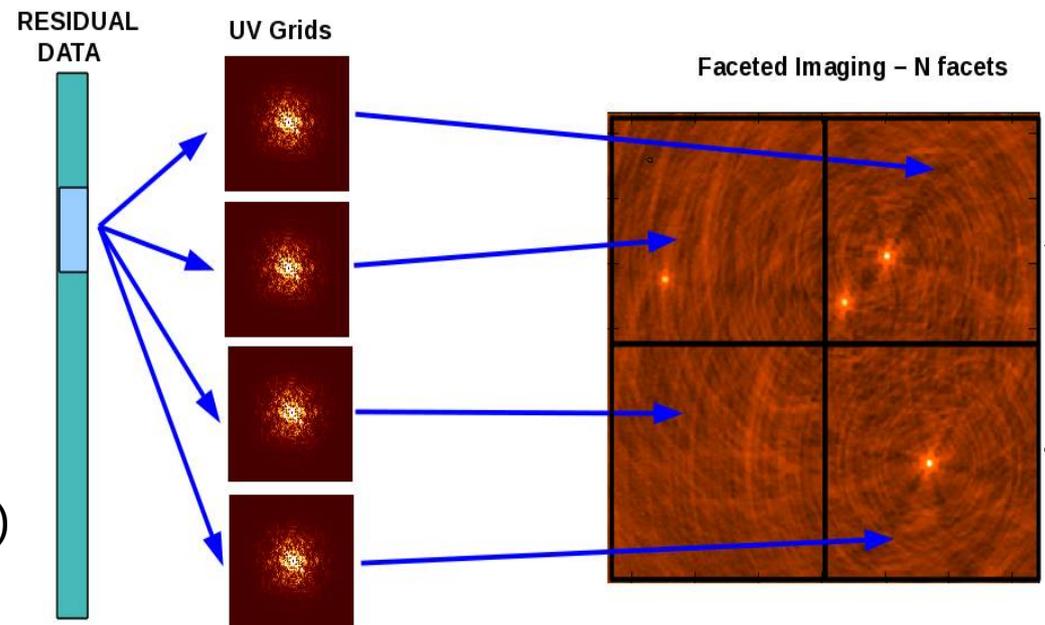
### Multiple fields (outlier fields)

- Usually, one large main field and several smaller outlier fields
- To avoid extremely large images
- Outlier file :  
List of image definition parameters



### Multiple Facets :

- Work with smaller field-of-view images  
To get around the w-term problem  
(non-coplanarity and sky curvature)
- Grid each facet separately onto a subimage, but do a single joint deconvolution (use PSF from first facet)



# Image Definition : Stitched and Joint Mosaics

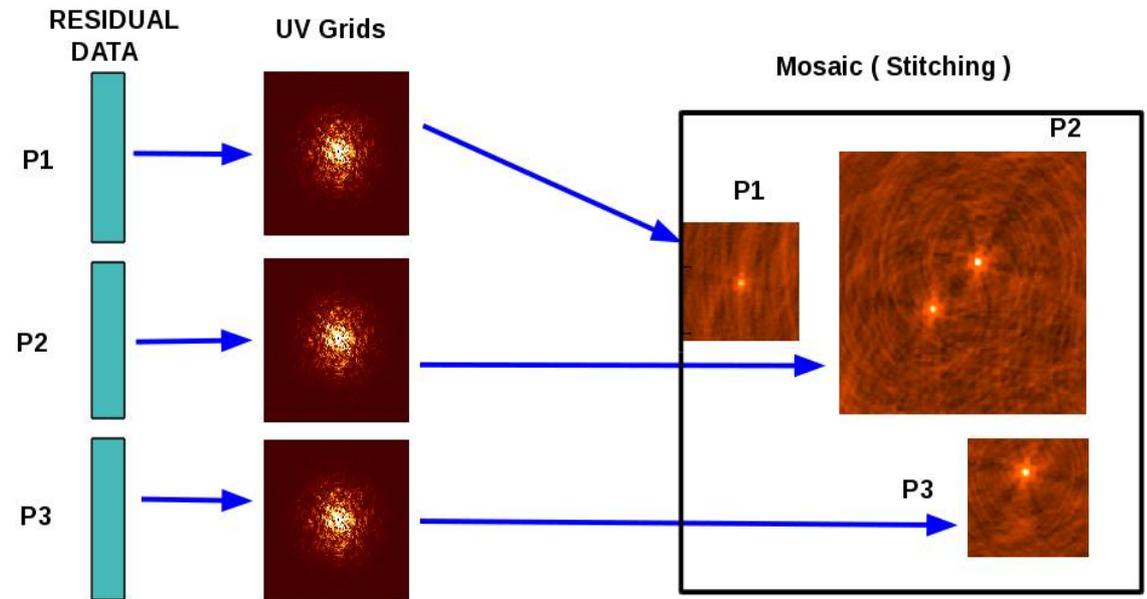
**Mosaic :**  
Combine data from multiple pointings

## Stitched Mosaic

- Image each pointing separately
- Combine them later

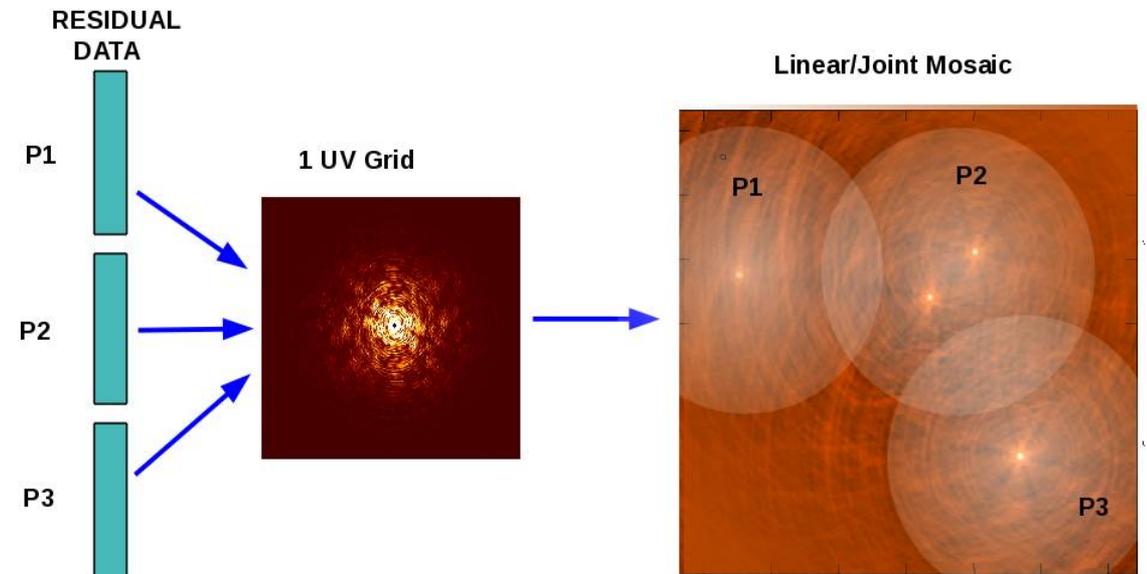
( Im : linearmosaic tool )

( Soon to come :  
'image mosaic', grid separately but  
combine before minor cycle )



## Joint mosaic

- Grid data from all pointings onto single uv grid, with appropriate phase gradients per pointing
- Joint deconvolution  
(assumes spatially invariant PSF )



# Iteration Control

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**Loop gain** (gain) : Controls how much of a source is modeled in each step.  
Choose according to algorithm being used (point source / extended)

**Exit criteria** (niter, threshold, cycleniter, cyclefactor, etc...) :

(1) **Minor cycle stopping criteria** (per chan/pol plane)

- Zero mask (i.e. all pixels = False )
- Iteration limit ( cycleniter )
- Peak residual in mask < cyclethreshold (computed from relative PSF sidelobe level )
- No iterations done
- Detects divergence (10% increase of peak residual across every ~2K iterations )

(2) **Global stopping criteria** (over all channels/pols, checked after each major cycle)

- Peak residual in mask < threshold
- Total number of iterations  $\geq$  niter
- Blank mask in all planes
- No change in peak residual from previous major cycle
- Divergence ( 5-times increase in peak residual from previous or from minimum )

( Soon to come : N-sigma based thresholds and iteration counts (per chan/pol plane) )

# Making and using deconvolution masks

**Mask for deconvolution :** CASA image filled with 1's and 0's  
( Different from internal T/F masks in CASA images )

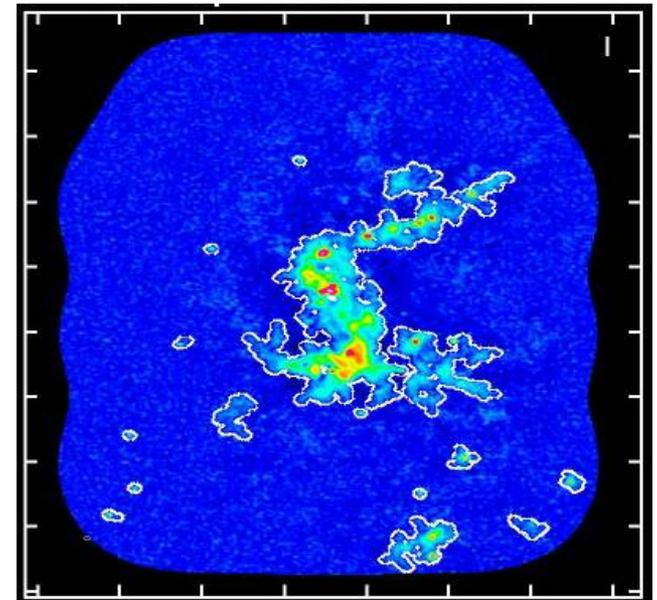
- Supply an image name as input via 'mask' parameter
- Supply a region string as input
- Automatic mask generation (tclean)
- Generate mask by drawing it interactively

**Task makemask :**

- copy from another image's T/F mask,
- LEL expression to compute mask from values
- evaluate a CASA region file to form a 1/0 mask
- extend/expand existing masks across planes

**Automasking :**

- auto-thresh (threshold based masks, using automatically computed rms)
- auto-multithresh (adaptively grow/prune masks as cleaning proceeds )
- pbmask : outer limit for mask based on PB gain

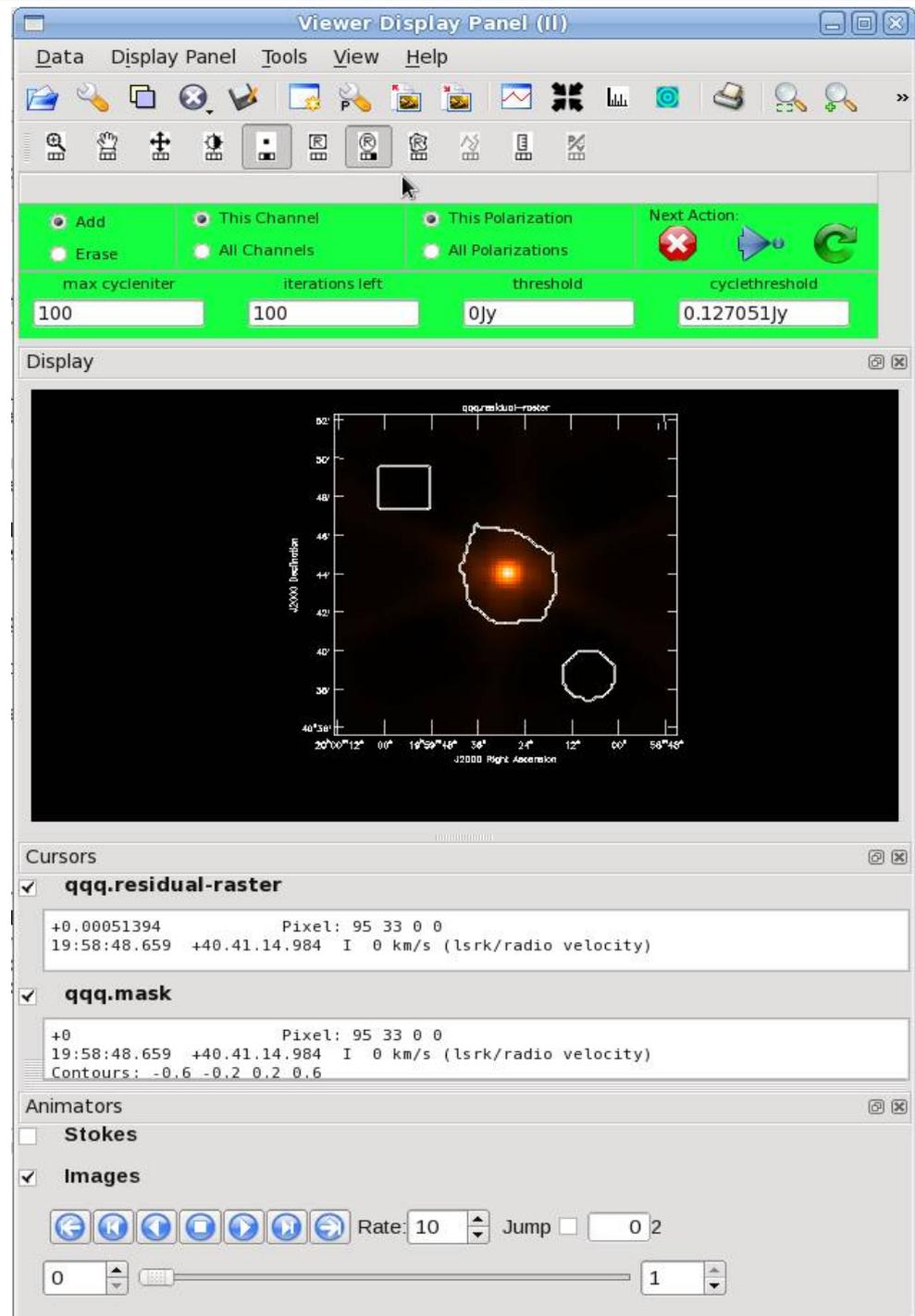


*Image from C.Brogan*

# Interactive masking (and iteration control)

## Interactive GUI

- View/edit masks in between major and minor cycles
- In tclean, the results of automasking will be visible and available for editing
- Edit masks using fixed shapes or freely defined region drawing  
( choose from menu via mouse button mapping )
- Copy masks across channels/pols or keep them separate
- Options to Stop, Continue silently, Continue/pause after next major cycle
- Iteration control options....



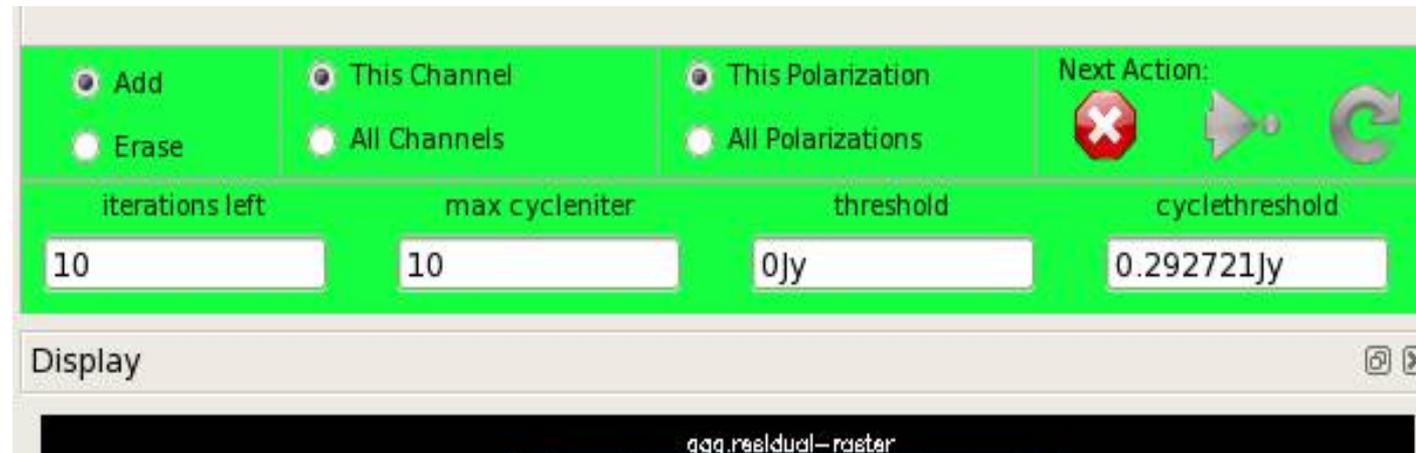
# Interactive masking (and iteration control)

## Interactive GUI

clean



tclean



- Added ability to control major cycle triggers directly (cycleniter, cyclethreshold)

In the future : Use the new CARTA viewer for interactive masking, monitor/control

- Use CARTA's region creation/manipulation tools
- Runtime display of convergence history (peak residual, total flux...)
- Ability to interrupt at logical stopping points, and edit iteration controls

# Some miscellaneous features

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- **Restart Options** ( `calcpsf=F`, `calcres=F` ) : Ability to restart from minor cycle without recomputing residual image and psf (`tclean`)
- Use the Measurement Set in **read-only mode** (multiple processes imaging same MS)
- **Saving a CLEAN model** (`savemodel`) : Possible during the last major cycle  
(or) in a separate step : `calcres=F`, `calcpsf=F`, `niter=0`
  - `modelcolumn` ( store model visibilities on disk in the MS )
  - `virtual model*` ( store gridder params to compute model visibilities on-the-fly when needed)

---- Can sometimes use the saved model for continuum subtraction (via task `uvsub`)  
( Note : 'imcontsub' and 'uvcontsub' fit arbitrary smooth polynomials per pixel/baseline )
- **Combining Single Dish and Interferometer data**
  - Combine SD image with restored output interferometer image (task `feather`)
  - Use SD image as starting model for Interferometric imaging (`clean`, soon to come in `tclean`)
- **Tool Interface** (`tclean`) : Python layer (`PySynthesisImager`) is useful for prototyping  
[https://casa.nrao.edu/casadocs/casa-5.1.0/global-task-list/task\\_tclean/examples](https://casa.nrao.edu/casadocs/casa-5.1.0/global-task-list/task_tclean/examples)

# Factors affecting runtime (and memory use)

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**How long will my imaging run take ? Well, it depends....**

- **Image size** : All operations scale by  $N^2$  (for an  $N \times N$  image),  
Choose a size optimized for fftw (factored by 2,3,5,7).
  - **Data size** : Major cycle scales linearly with data volume
  - **Gridder** : Size of gridding convolution function ranges from 3 pixels on a side to  
~few x 100 on a side => Orders of magnitude difference in runtime
  - **Deconvolver** : MS-Clean, MTMFS are more expensive than Hogbom/Clark
  - **Iterations** : Number and frequency of major cycles, appropriateness of minor cycle  
algorithm to sky structure, number of minor cycle steps (and loop gain)
  - **Sky structure** : High/low SNR, Calibration artifacts, easy/hard convergence, etc ?
  - **Machine specifications** : RAM / Swap, Disk speed (local, nfs, ssd), nCPUs, Ncores,..
- ( Soon to come : A resource predictor for tclean to estimate peak memory use  
**Status : Under development**
- Soon to come : Parallelization of entire run (cube) and for major cycles (continuum)  
**Status : Exists and in a late stage of testing/validation )**

# Task Interface ( tclean )

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**vis** = 'msname' or [ 'msname1', 'msname2', ... ]

- Selection options (field, spw, ... )

**imagename** = 'xxx'

- Image definition options ( imsize, cellsize, startmodel, ... )

**specmode** = 'mfs' or 'cube'

- Channelization options (start, step, nchan,... in different units, reffreq )

**gridded** = 'standard', 'widefield', 'mosaic', 'imagemosaic', 'awproject'

- Algorithm-specific parameters (wprojplanes, cfcache, pblimit, etc....)

**weighting** = 'natural', 'uniform', 'briggs'

- Options for some choices (robust,...)

**deconvolver** = 'hogbom', 'clark', 'multiscale', 'mtmfs',

- Algorithm-specific options (scales, nterms, ... )

**usemask** = 'user', 'pb', 'auto-thresh', 'auto-multithresh'

- Mask creation options (pbmask, threshold, maskname, etc...)

**niter** = N

- Iteration control options (gain, threshold, cycleniter, cyclefactor, interactive=T/F...)

**parallel** = True/False ( when started up with 'mpicasa' )

( + outlier files, restart option (calcres=T/F, calcpsf=T/F), savemodel, restoration, etc...

# Output Images

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<b>imagename.psf</b>	Point Spread Function
<b>imagename.pb</b>	Primary Beam
<b>imagename.residual</b>	Residual Image (or initial Dirty Image)
<b>imagename.model</b>	Model Image after deconvolution
<b>imagename.image</b>	Restored output image
<b>imagename.image.pbcor</b>	Primary Beam corrected image
<b>imagename.mask</b>	Deconvolution mask
<b>imagename.sumwt</b>	A single pixel image containing sum of weights per plane
<b>imagename.weight</b>	Image of un-normalized sum of PB-square (for mosaics and A-Projection)
<b>imagename.XX.{tt0,tt1,tt2}</b>	Multi-term images representing Taylor coefficients (of polynomials that model the sky spectrum)
<b>Imagename.workdirectory</b>	Working directory for partial products during parallel runs

# Image Viewing and Analysis

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## Viewing Images

Task **viewer** ( or **casaviewer** from the command line )

- Image cubes ( 2D displays with animation controls for other dimension(s) )
- Rasters, contours, vectors, markers
- Stacks of images, each of which can be a 2D, 3D or 4D 'cube'
- Region drawing + statistics + histograms
- Spectral Profile tools : Line overlays, Spectral fitting, P/V diagrams, etc
- 2D image fitting
- Colormap editing, intensity scaling, etc on the 2D raster (per image, or global)

( scripting the viewer via task **imview** )

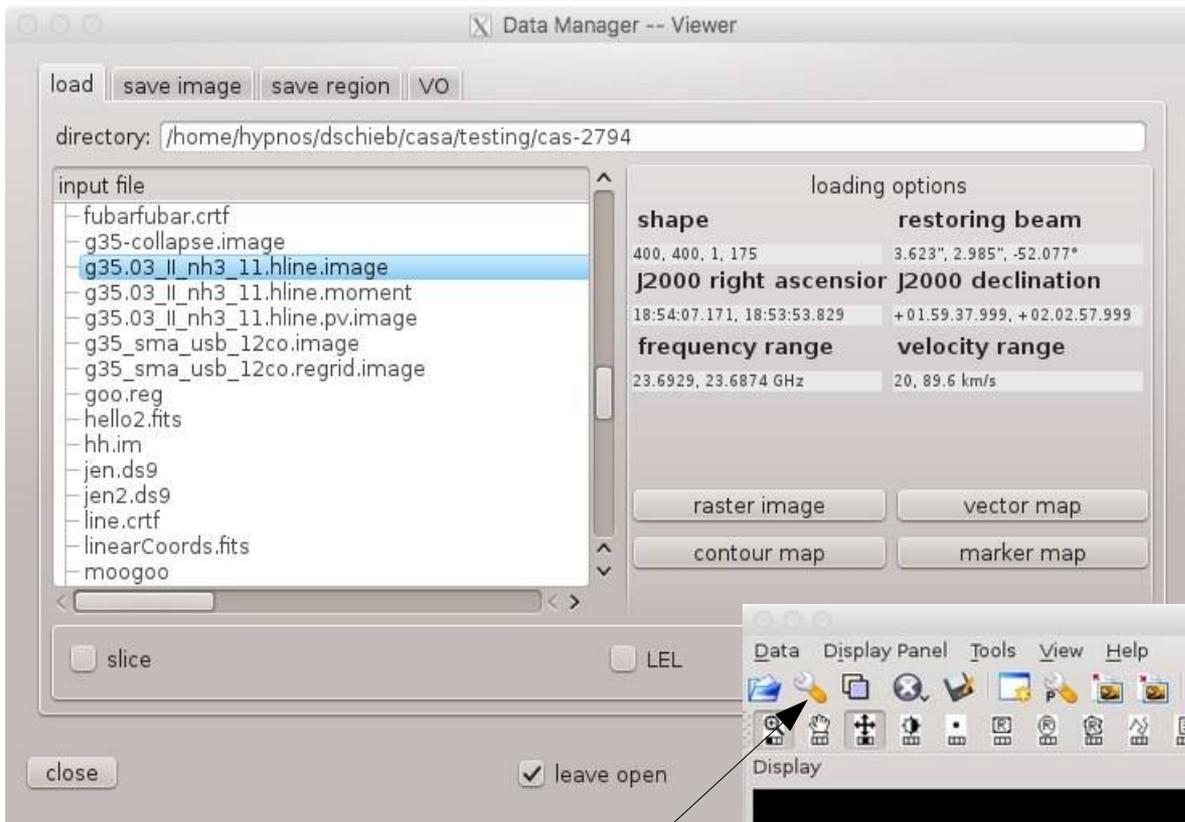
CASAdocs : <https://casa.nrao.edu/casadocs/casa-5.1.0/image-cube-visualization>

## Image Analysis ( **28 separate tasks** )

- Import/export, viewing headers, reformatting (regridding, spectral collapse, etc)
- Mathematical analysis – image plane + spectral
- Working with Regions, masks, image sub-selection syntax

CASAdocs : <https://casa.nrao.edu/casadocs/casa-5.1.0/image-analysis>

# Image Viewing and Analysis – Viewer task + GUI

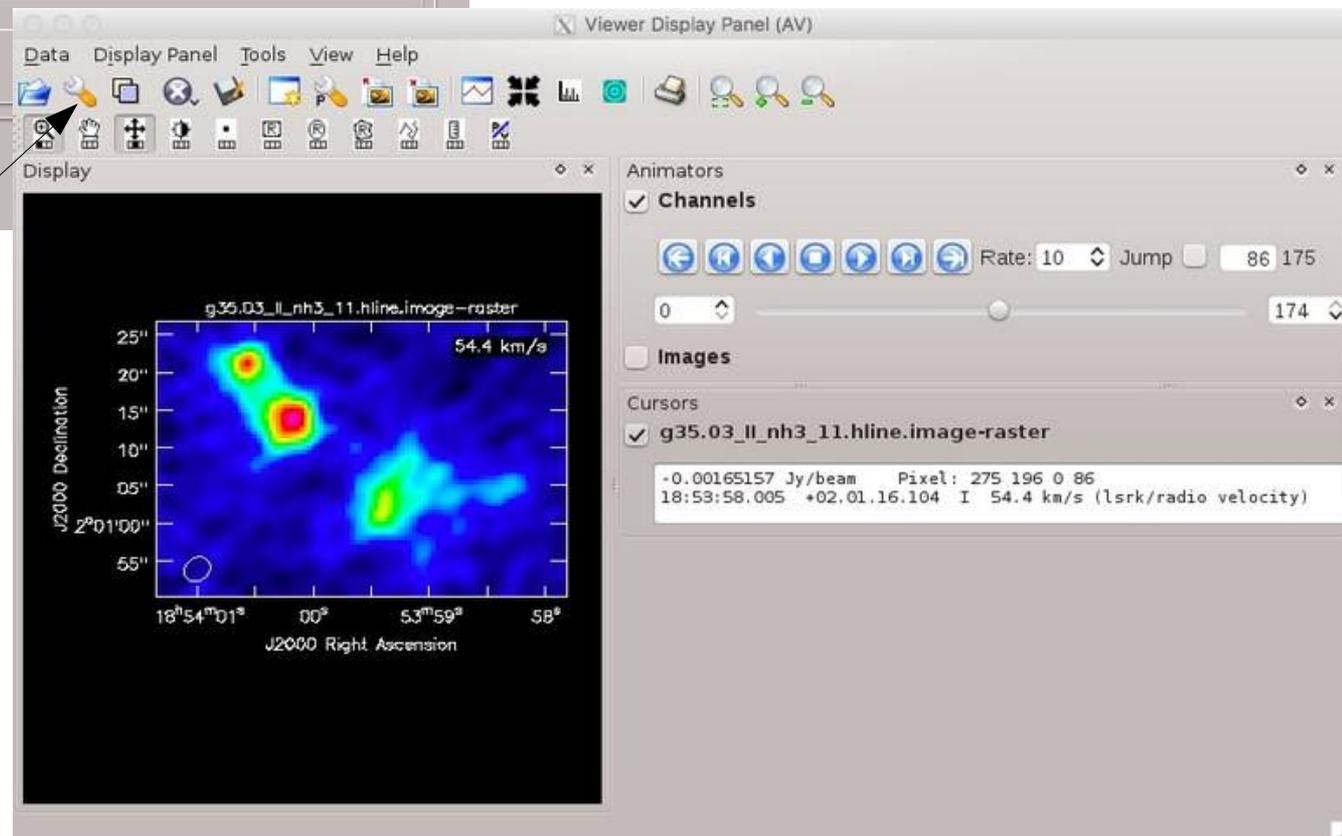


## Data Manager

- Select images and display type

## Viewer Display Panel

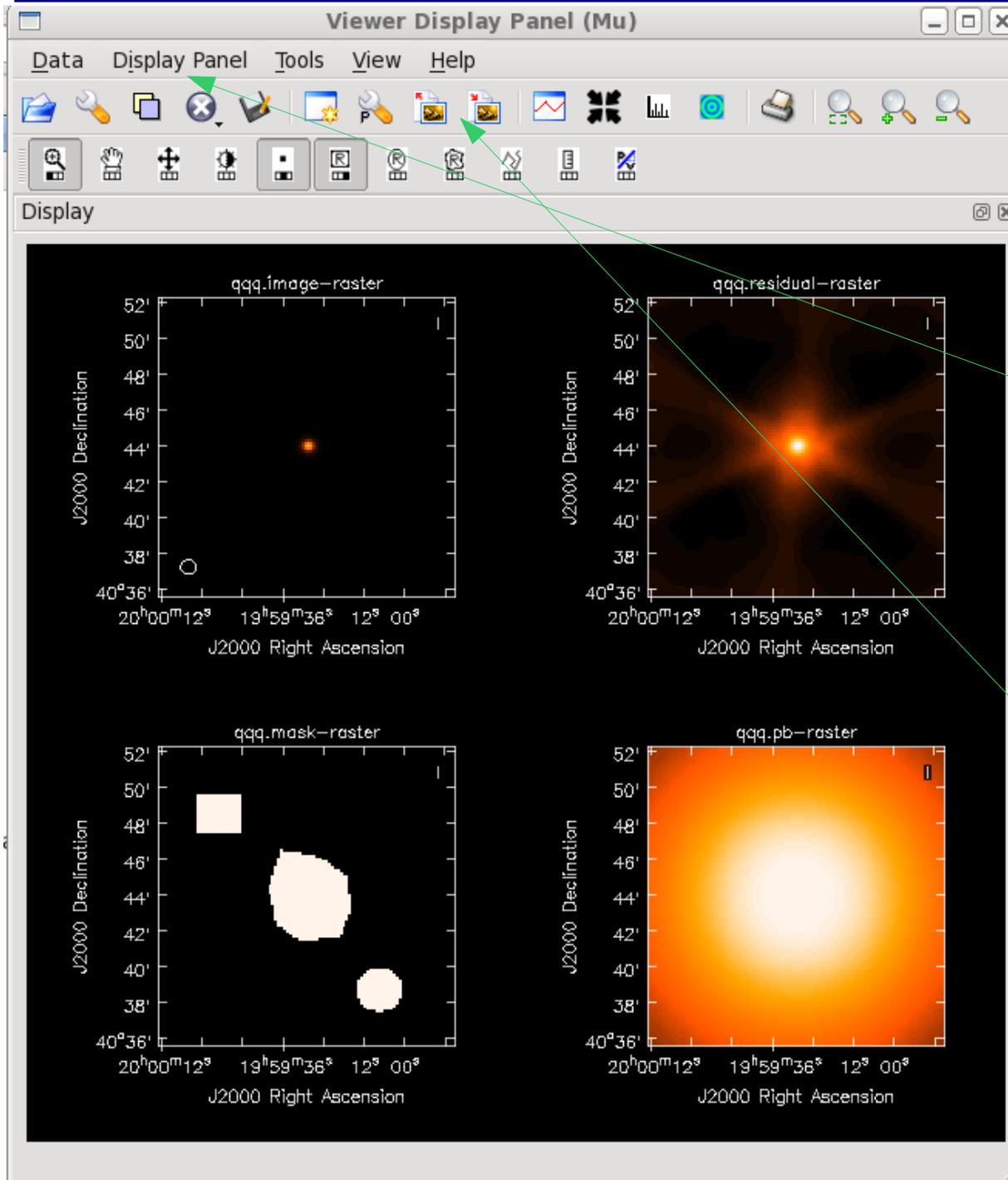
- Cursor tracking display
- Axes animation controls
- Drawing tools by mapping mouse buttons to functions



## Display settings

- Adjust panel
- Color scale, data range
- Axis labels
- Display/animation axes
- Coordinate systems
- ...etc...

# Image Viewing and Analysis – Stacks and Multi-panel displays



## Multi-Panel displays

- Load several images (one at a time)
- Configure Nx, Ny in the Display Panel options
- Zooming / stats / region drawing can operate on all images
- Adjust display options per image or globally

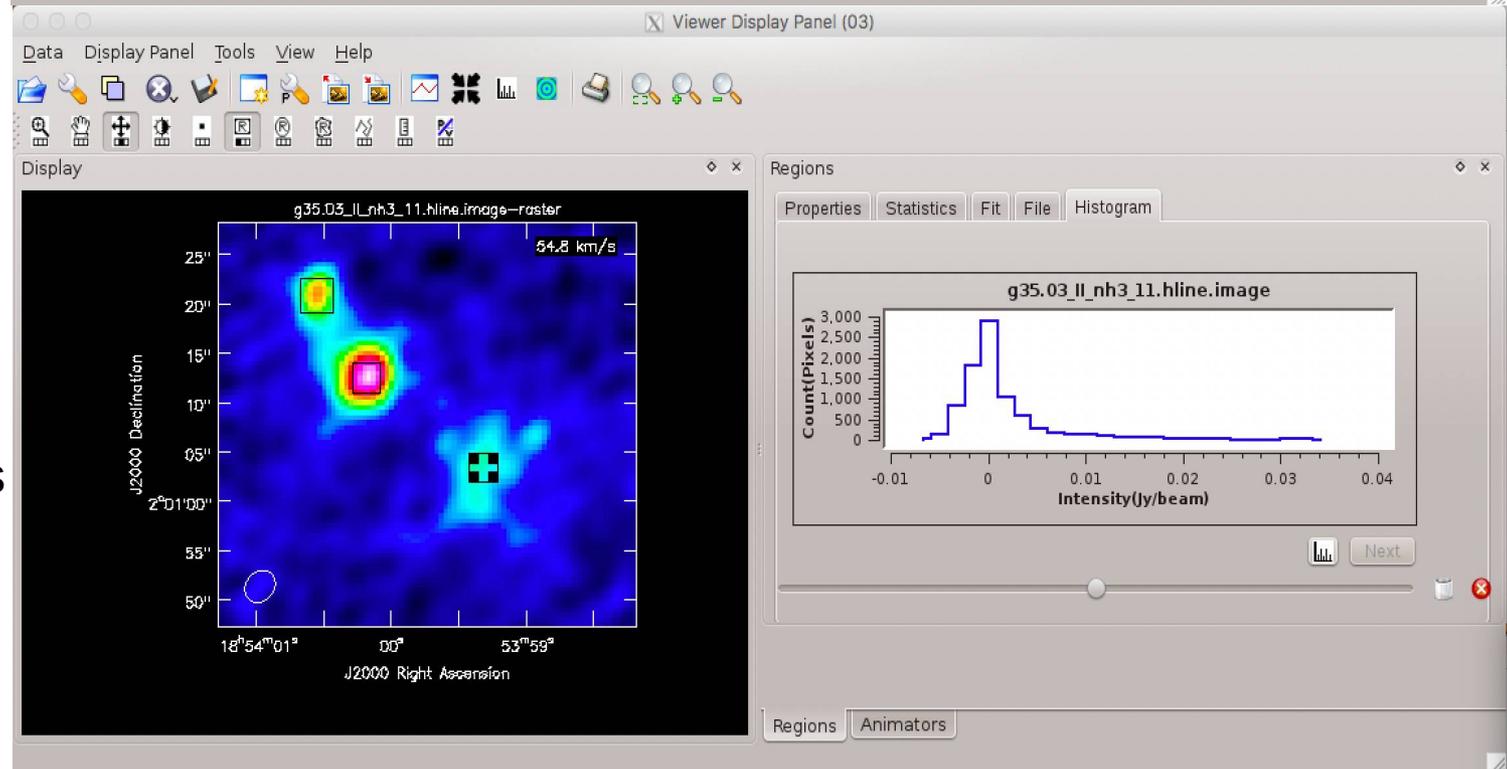
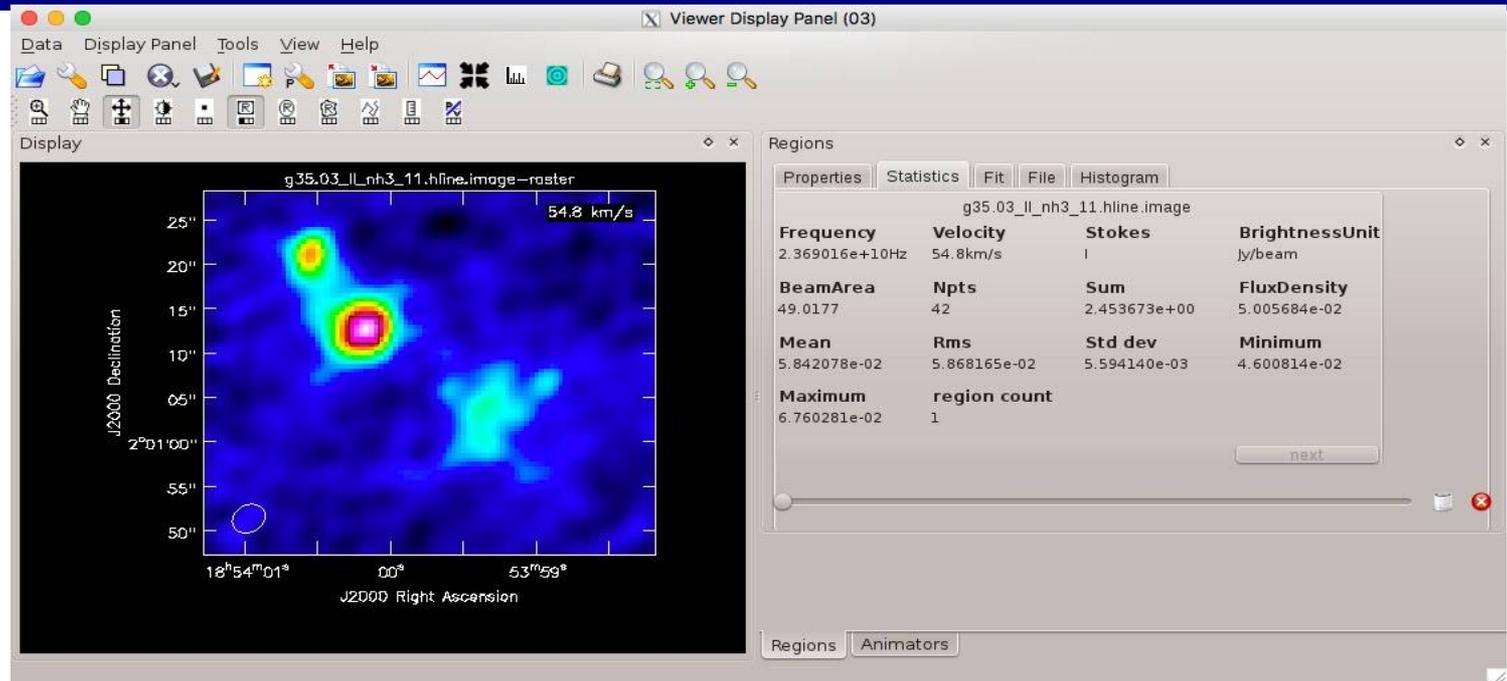
## Preserving viewer State

- Save current state of the GUI (xxx.rstr file on disk)
- Restore later upon fresh restart of the viewer

# Image Viewing and Analysis – Region statistics

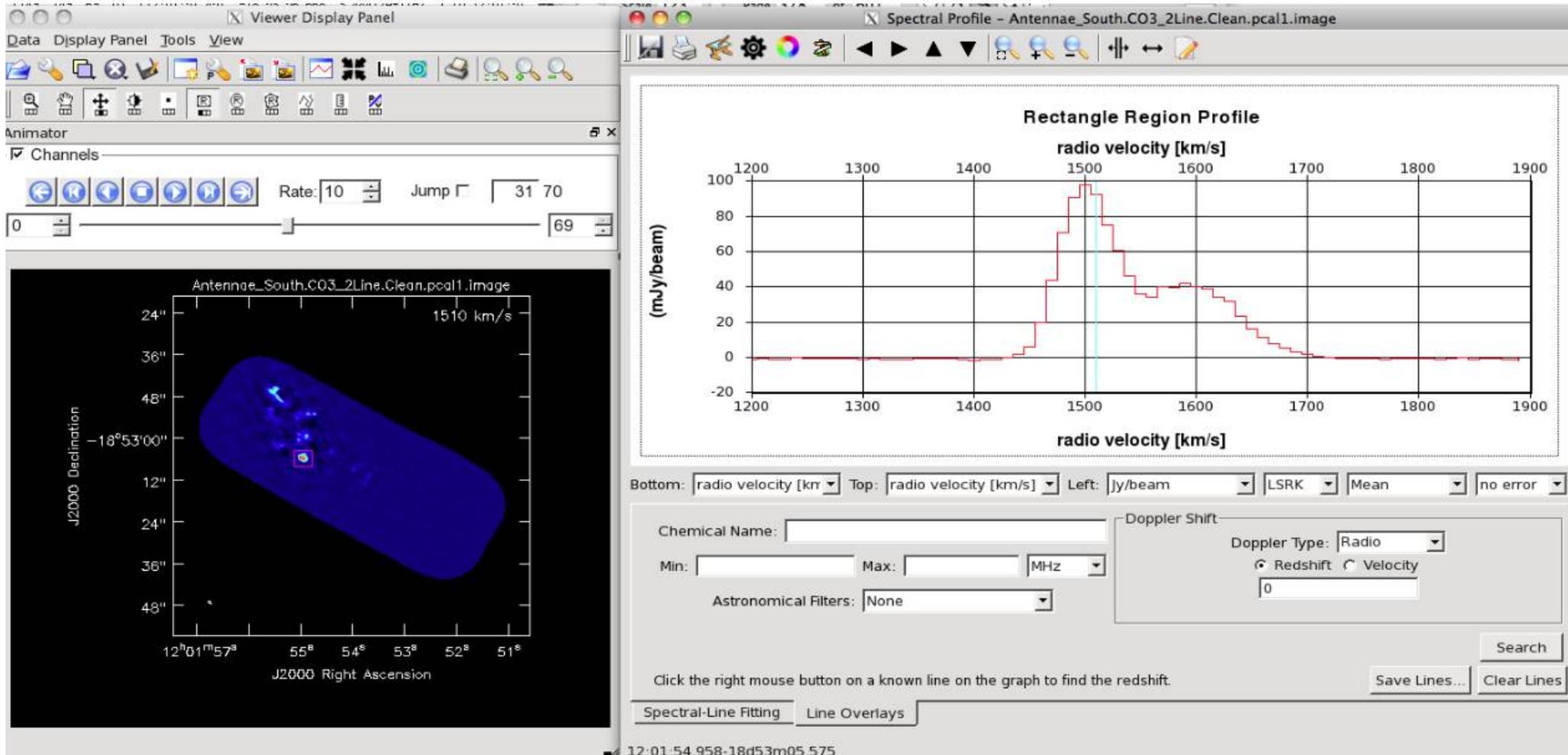
## Regions and Statistics

- Draw regions of Different shapes
  - Save/import region files
  - Display statistics
  - Display histograms
  - Animation axis for Stacks of images
  - Region stats also printed to terminal
- ( Task **imstat** also does Image statistics )



# Image Viewing and Analysis – Viewer's Spectral Profile tool

View spectra of flux within selected regions

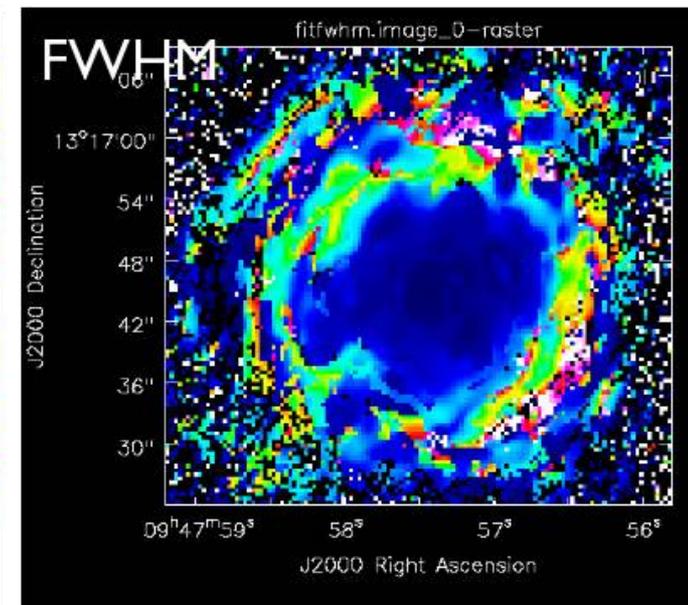
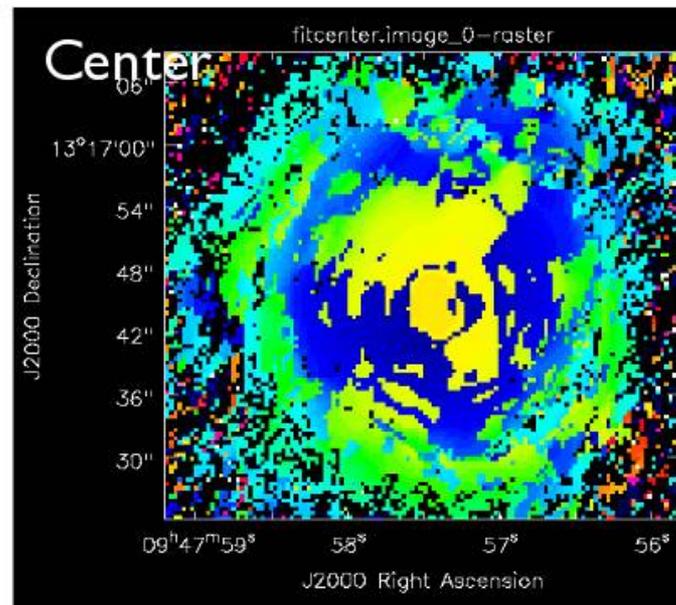
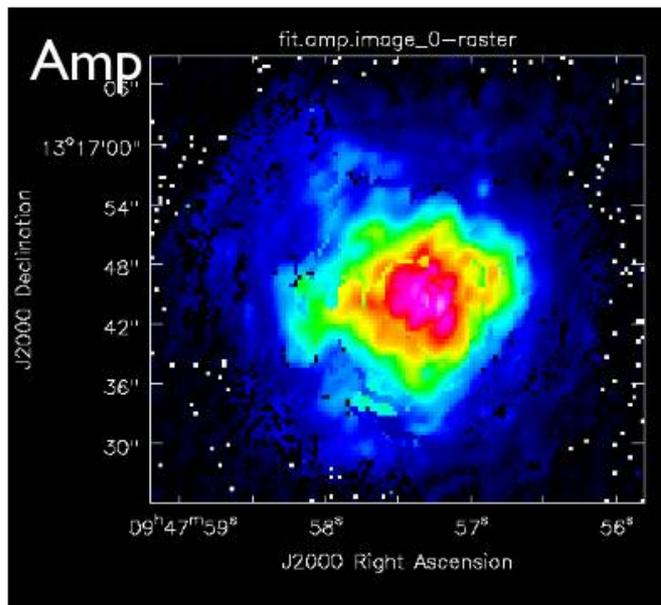


- Spectral Line fitting tool (also done by task **specfit**)
- On-the-fly spectral frame conversions
- Line overlays (via queries of Splatalogue spectral line database)

# Image Viewing and Analysis – specfit task

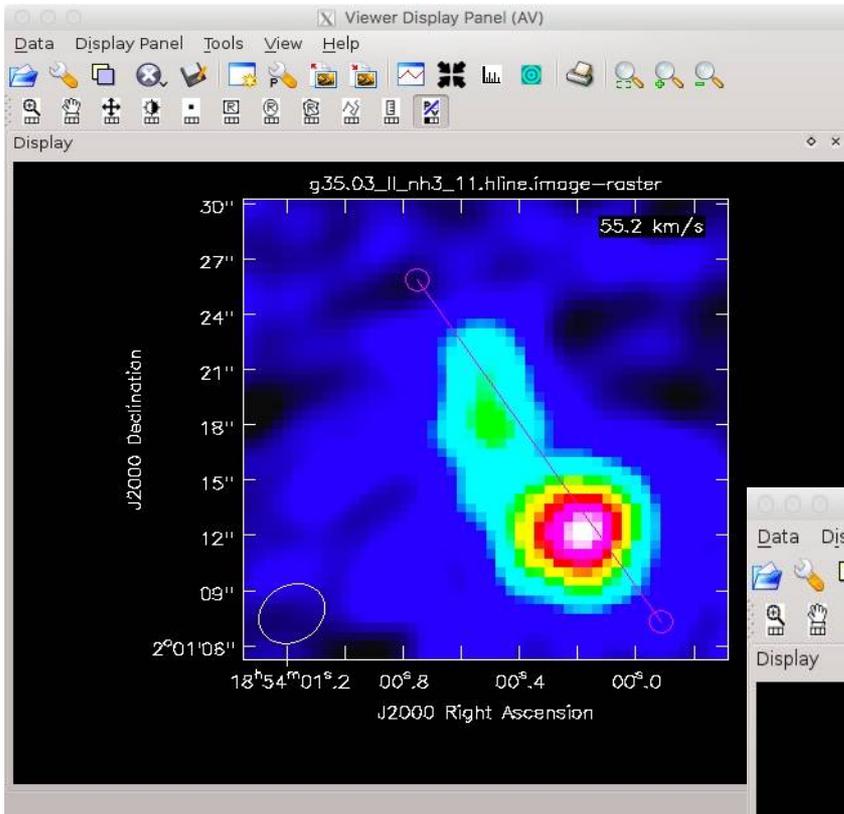
## Specfit task

- Fits 2D Gaussians and/or polynomial models to an image/region, typically along the Spectral axis.
- Can fit multiple Gaussian multiplets with constraints
- Can be per-pixel ==> Output images of Amp, FWHM, Center velocity

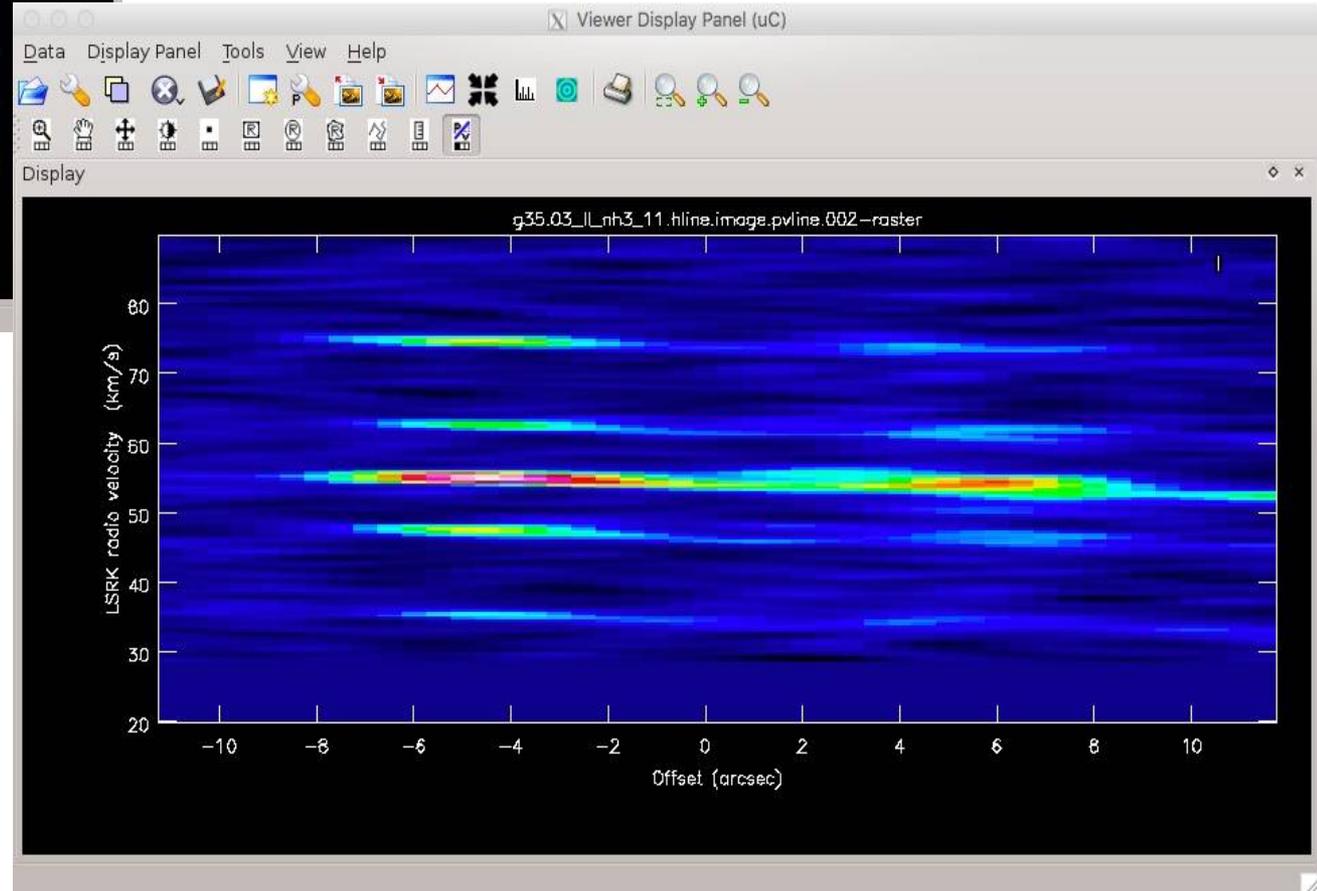


*Images from J.Ott*

# Image Viewing and Analysis – Position/Velocity diagrams



Select a cut across the image,  
and make a P/V diagram



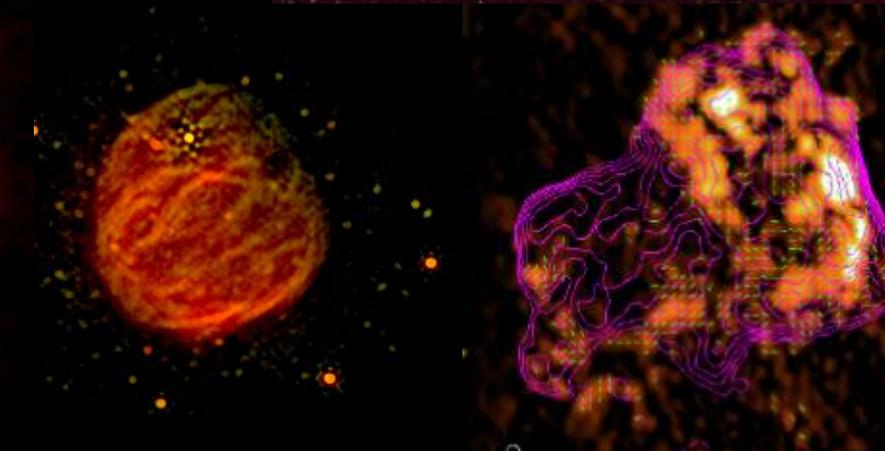
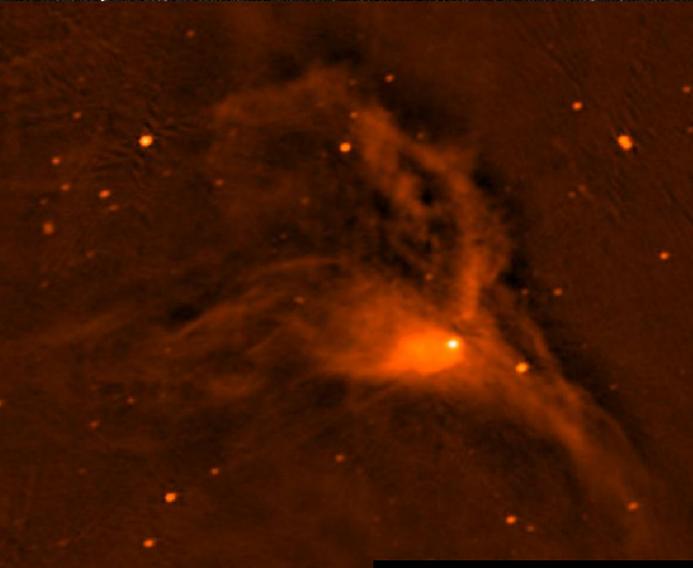
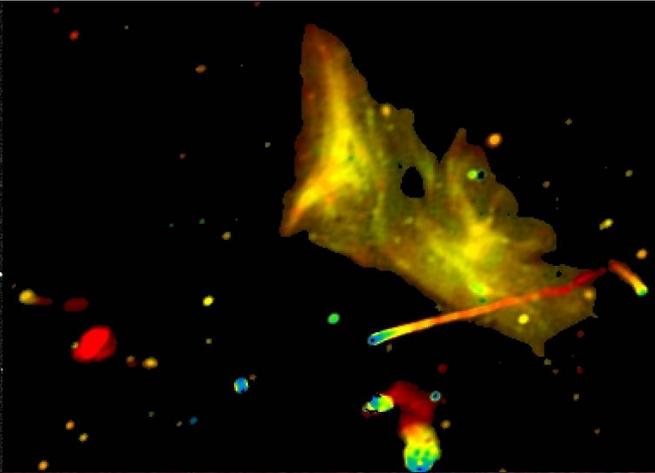
( Task **impv** also constructs  
a Position-Velocity diagram )

# Image Analysis – 28 different tasks

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- Import/export : **importfits, exportfits, imval**
- Viewing headers : **listfits, imhead, imhistory**
- **Reformatting** (resizing, binning, regridding to different csys, collapsing axes)
  - **imsubimage, imtrans, imregrid, imreframe, imrebin, imcollapse**
- **Spectral Analysis** : **imcontsub, immoments, impv, specsmooth, specfit, specflux, plotprofilemap, rffit, spxfit, slsearch (splattotable)**
- **Image domain analysis** : **imfit, imsmooth, immath, imstat, imdev**
- **Additional topics** (please see [CASAdocs](#)):
  - Tool interface : [ia tool](#)
  - Working with [Regions](#) ( syntax and CASA Region File Format )
  - Working with [T/F masks](#) within images (e.g. [ia.maskhandler\(\)](#) tool)
  - Working with [Lattice Expression Language](#) (for math, and sub-selection)
  - Read pixel values into Python arrays/dictionaries for custom analysis  
E.x. `data = imval('xxx.image')`

# Questions ?



*Images from  
Golap, Marvil,  
Maercker, Rau,  
Ott, Owen,  
Bhatnagar,  
Brogan*