Synthesis imaging using CASA

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Radio interferometric observations

The products from the array are Visibilities (Fourier domain, u-v), not images (physical domain, x-y).



What to do with visibilities calibrated?

- Stay in the uv-domain and perform your analysis in the Fourier domain (works for simple source geometry)
- Reconstruct a model sky brightness distribution, then perform your analysis in the physical domain

 $V_{\text{obs}}(u, v) = I_{\text{model}}^{\text{dirty}} I_{\text{model}}^{\text{dirty}}(x, y)$ $I_{\text{model}}^{\text{clean}}(x, y) \sim I_{\text{true}}(x, y)$



TW Hya 7mm cont. (Hughes et al. 2007)



Nomenclature

- Dirty beam (synthesized beam): point-source response (point spread fn.), FT of uv-coverage and normalize the peak to unity [Jy/beam]
- Dirty map: FT⁻¹ of visibilities [Jy/beam]
- Clean beam: (usually) a 2D Gaussian approximation of the main lobe of the dirty beam [Jy/beam]
- Clean map: the sum of clean component map convoluted with clean beam and the residual map [Jy/beam]
- Clean component map: an assumable of point sources determined from CLEAN algorithms [Jy/pixel]
- Residual map: a dirty map with clean components (convolved with dirty beam) removed, usually noise-like [Jy/beam]

CLEAN (Högbom 1974)

recommended to read

- An (iterative) algorithm proposed/summarized by J. A. Högbom
 - I: compute dirty beam and dirty map
 - II: subtract over the whole dirty map with a dirty beam pattern (multiplied by a factor, e.g., 0.1) which is centred at the point where the dirty map has its maximum of absolute value of intensity
 - III: Repeat step II, each time replacing the dirty map by the remaining map (i.e., residual map) from the previous iteration
 - IV: once the iteration is satisfied, generate the clean map by adding the final residual map with the clean component map convolved with the clean beam



dirty beam

10

15

1.0 0.8

0.6

0.2 0.0 -0.2

-15

-10

beam)

<u>}</u> 0.4

intensity





What if...

- I iterate the CLEAN process too many times?
- the strength of sidelobes is similar to the main lobe? (i.e., dirty map is really "dirty")



dirty beam

15

1.0 0.8

0.6 0.4

0.2 0.0

JISIT





dirty beam

10

15

1.0 0.8 0.6

> 0.4 0.2

0.0 -0.2

ensity





dirty beam

10

15

1.0 0.8 0.6

> 0.4 0.2

0.0 -0.2

ensity



Real data: a face-on spiral galaxy





CLEAN: CASA-setup

CLEAN with CASA



CASA 4.7.2 (<u>https://casa.nrao.edu/casa_obtaining.shtml</u>)

	KOKOKOK	0707070	XOXOXO
		200	nv/
0202020	000000		

```
> inp clean
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> clean?
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			000000				00000		000000	0000000	00000		000000	
# c	lean ::	Invert	and	deconvolve	images	wit	h se	electe	d al	gorit	:hm			
vis			=	1.1	7	# N	ame	of in	put	visit	oilit	y fil	e	
imag	ename		=		7	# P	re-1	name o	f ou	tput	imag	jes		
outl	ierfile		=		7	# т	ext	file	with	imag	je na	mes,	size	es,
					7	¥	cent	ters f	or o	utlie	ers			
fiel	d		=		7	# F	iel	d Name	or	id				
spw			=		7	# S	pect	tral w	indo	ws e.	.g. '	0~3',		is
					7	¥	all							
sele	ctdata		-	True	7	⊭ Ο	the	r data	sel	ectio	on pa	iramet	ers	
	timeran	ge	=		7	# R	ange	e of t	ime	to se	elect	: from	n dat	a
	uvrange	1	=		7	# S	ele	ct dat	a wi	thin	uvra	inge		
	antenna		=		7	# S	ele	ct dat	a ba	sed o	on an	tenna	i/bas	el
	scan		=		7	# S	can	numbe	r ra	nge				
	observa	tion	=		7	⊭ Ο	bse	rvatio	n ID	rang	je			
	intent		=		7	# S	can	Inten	t(s)					
mode	1		-	'mfs'	7	# S	pect	tral g	ridd	ing t	:ype	(mfs,	cha	inn
					7	¥	vel	ocity,	fre	quenc	ey)			
	nterms		=	1	7	# N	umb	er of	Tayl	or co	effi	cient	s to	
					7	¥	mode	el the	sky	freq	quenc	y der	pende	enc
	reffreq		=		7	# R	efe	rence	freq	uency	/ (nt	erms	> 1)	1
					7	ŧ	use	s cent	ral	data-	-freq	fuency	7	

× vis

The value of the vis parameter is either the name of a single MS, or a list of strings containing the names of multiple MSs, that should be processed to produce the image. For example, vis = 'ngc5921.ms' set a single input MS, while vis = ['ngc5921_day1.ms', 'ngc5921_day2.ms'] points to two separate measurement sets that will be gridded

together to form the image.

field

The field parameter selects the field indexes or names to be used in imaging. Unless you are making a mosaic, this is usually a single index or name: field = '0' # First field (index 0) field = 'NGC1234' # source name field = '0~10' # 11 pointings to mosaic

Spw

The spw parameter selects the spectral windows that will be used to form the image, and possibly a subset of channels within these windows.

The spw parameter is a string with an integer, list of integers, or a range, e.g.

- spw = '1' # select spw 1
- **spw = '0,1,2,3'** # select spw 0,1,2,3
- spw = '0~3' # same thing using ranges
- spw = '1:10~30' # select channels 10-30 of spw 1
- spw = '0:5~55,3:5;6;7' # chans 5-55 of spw 0 and 5,6,7 of spw 3

imagename

The clean task will produce a number of output images based on the root name given in imagename. These include: <imagename>.image # the restored image <imagename>.iflux # the effective response (e.g. for pbcor) <imagename>.iflux.pbcoverage # the PB coverage (ftmachine='mosaic' only) <imagename>.model # the model image <imagename>.residual # the residual image <imagename>.psf # the synthesized (dirty) beam

imagename

WARNING: If an image with that name already exists, it will in general be overwritten. Beware using names of existing images. However, if the clean is run using an imagename where <imagename>.residual and <imagename>.model already exist then clean will continue starting from these (effectively restarting from the end of the previous clean). Thus, if multiple runs of clean are run consecutively with the same imagename, then the cleaning is incremental.

The cell parameter defines the pixel size in the x and y axes for the output image. If given as floats or integers, this is the cell size in arc seconds, e.g.

cell=[0.5,0.5]

make 0.5" pixels. You can also give the cell size in quantities, e.g.

cell=['1arcmin', '1arcmin']

If a single value is given, then square pixels of that size are assumed.

Usually cell = 1/5 of the minor axis of the clean beam

imsize

The image size in numbers of pixels on the x and y axes is set by imsize. For example,

imsize = [640, 640]

makes a square image 640 pixels on a side. In general, the best performance is obtained with image sizes that are even and factorizable to 2,3,5,7 only. An easy rule of thumb would be $2n \times 10$ where n is an integer number, like 160, 320, 640, 1280, 2560, etc.

Usually imsize= FOV (1.22 λ /D) or mosaic area+half FOV each side

phasecenter

The phasecenter parameter indicates which of the field IDs should be used to define the phase center of the mosaic image, or what that phase center is in RA and Dec. The default action is to use the first one given in the field list. For example:

phasecenter='5' # field 5

phasecenter='J2000 19h30m00 -40d00m00' # specified

position

mode (expandable)

The mode parameter defines how the frequency channels in the synthesis MS are mapped onto the image. The allowed values are:

mfs # multi-frequency-synthes
for continuum imaging
channel # image the specified
velocity # for line imaging
frequency # grid in frequency

e.g. LSRK, BARY

ode		=	'velocity'
	nchan	-	-1
	start	-	• •
	annels width	-	
	interpolation	-	'linear'
	resmooth	-	False
	chaniter	-	False
-	outframe	-	• •
	veltype	-	'radio'

Spectral gr velocity, Number of c image; -1 Velocity of '0.0km/s'(SpW of MS) Channel wid (''=width SpW of MS) Spectral in linear, cu Re-restore beam when # Clean each (True), or (False) spectral re image; ... Velocity de

imagermode

This choose the mode of operation of clean, either as single-field deconvolution using image-plane major and minor cycles only (imagermode=''), single-field deconvolution using Cotton-Schwab (CS) residual visibilities for major cycles (imagermode='csclean'), or multi-field mosaics using CS major cycles (imagermode='mosaic').

In the CS mode, cleaning is split into minor and major cycles. For each field, a minor cycle is performed using the PSF algorithm specified in psfmode. At major-cycle breakpoints, the points thus found are subtracted from the original visibilities.

psfmode

The psfmode parameter chooses the "algorithm" that will be used to calculate the synthesized beam for use during the minor cycles in the image plane. The value types are strings. Allowed choices are 'clark' (default) and 'hogbom'.

s gain

The gain parameter sets the fraction of the flux density in the residual image that is removed and placed into the clean model at each minor cycle iteration. The default value is gain = 0.1 and is suitable for a wide-range of imaging problems. Setting it to a smaller gain per cycle, such as gain = 0.05, can sometimes help when cleaning images with lots of diffuse emission.

threshold

The threshold parameter instructs clean to terminate when the maximum absolute residual reaches this level or below. Note that it may not reach this residual level due to the value of the niter parameter which may cause it to terminate early.

threshold = (0.1 mJy)

niter

The niter parameter sets the maximum total number of minorcycle CLEAN iterations to be performed during this run of clean. niter = 2000

weighting

To determine how we weight the visibilities.

For weighting='natural', visibilities are weighted only by the data weights, which are calculated during filling and calibration and should be equal to the inverse noise variance on that visibility. It gives best signal-to-noise ratio, worst angular resolution.

For weighting = 'uniform', the data weights are calculated as in 'natural' weighting. The data is then gridded to a number of cells in the uv-plane, and after all data is gridded the uv-cells are re-weighted to have "uniform" imaging weights. It gives best angular resolution, worst signal-to-noise ratio.

The weighting = 'briggs' mode is an implementation of the flexible weighting scheme. The robust keyword takes value between -2.0 (close to uniform weighting) to 2.0 (close to natural).

restfreq

The value of the restfreq parameter, if set, will overwrite the rest frequency in the header of the first input ms to define the velocity frame of the output image. For example:

restfreq='115.2712GHz',

will set the rest frequency to that of the CO 1-0 line.

ALERT: Setting restfreq explicitly here in clean is good practice, and may be necessary if your ms has been concatenated from different files for different spectral windows

mask

To limit the clean algorithm where to look for clean components. Default is the full image plane. Putting a mask is sometimes useful for poor dirty beam and for somehow known source structure. Check manual for more detail.

pbcor

The pbcor parameter (True or False) controls whether the final .image is scaled to correct for the Primary Beam of the array or not. Check the task impbcor as well.

vis field

SPW

imagename

cell

imsize

phasecenter phasecenter magennode imagennode

check manual or type "clean?" for more information

Dsfmode gain

niter

threshold

restfreq

mask bbcor Weighting



Before we start

- Demo dataset: "continuum.ms" and "molecular_line.ms", both come from the IRAS16293-2422 Band4 science verification (SV) dataset (<u>https://almascience.nao.ac.jp/</u> <u>almadata/sciver/IRAS16293Band4/</u>)
- IRAS16293-2422: a low-mass protostellar binary system with a rich amount of molecular line emission



The contents of a MS: listobs

CASA <2>: listobs vis='continuum.ms'

- With "listobs", figure out
 - How many fields are observed?
 - How many spectral windows (spw)in the ms?
 - Center frequency?
 - Bandwidth?
 - Number of channels?
 - Channel width?
 - How many antennae in the array?

CASA <2>: listobs vis='continuum.ms'

• • •	Log Me	essages (:/Users/	kswang/Do	cuments/sy	nthesis_ima	aging_tutorial2	016/demo_	dataset	/casapy-20160818	3-021652.log)		
		Searc	h Message:			<i>i</i> *6	Filter:	Time	•		\Box	C
Message												
CASA Version	4.6.0-REL (r36590)										
Tagged on:	2016-04-01	11:11:00 -0400	(Fri, 01	Apr 2016)								
**********	**********	*************	****									
##### Begin	Task: listob	s #	####									
listobs(vis=	"continuum.m	s", selectdata=	Frue, spw="	",field="	", antenna=	·",						
uvra	inge="",timer	ange="",correl	ation="",s	can="",in	tent="",							
feed	="",array=""	, observation="	verbose=	True, list	file="",							
list	unfl=raise,c	achesize=50,ov	erwrite=ra	use)			_					
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Observer	satoko	Project: uid:/	/2002/2866	6bf/Xc			-					
Observation:	ALMA	rioject. ulu.,	AUUL/AUUU	001/20								
Data records	: 166980	Total elaps	ed time =	4550.98 s	econds							
Observed	from 11-Ju	1-2014/04:11:3	6.5 to	11-Jul-2	014/05:27:	27.5 (UTC)						
Compute subs	can properti	es										
Observati	onID = 0	ArrayID =	0									
Date	Timerange	(UTC)	Scan Fld	Id FieldN	ame	nRows	SpwId	s Ave	erage Interval(s) ScanInt	ent	
11-Jul-201	4/04:11:36.5	- 04:17:56.8	9	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	04:18:56.3	- 04:25:16.6	11	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	04:26:37.0	- 04:32:57.2	14	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	04:34:03.9	- 04:40:24.2	16	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	04:41:46.9	- 04:48:07.2	19	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	04:49:06.5	- 04:55:26.7	21	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	04:56:45.0	- 05:03:05.3	24	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	05:04:12.9	- 05:10:33.2	26	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	05:11:57.0	- 05:18:17.2	29	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	05:19:17.2	- 05:25:37.4	31	0 IRAS16	293-2422	165	60 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
	05:26:57.3	- 05:27:27.5	34	0 IRAS16	293-2422	13	80 [0]	[6.05]	[OBSERVE_TARGET	#ON_SOURCE]		
((nRows = Total number of rows per scan)											
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ID Code	Name	RA 16.22.1		Deci	Бро	och Srcia	nRows					
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Antennas: 23	1	14005										

The contents of ms: listobs

CASA <2>: listobs vis='continuum.ms'

command line mode, scripting style

- With "listobs", figure out
 - How many fields are observed? [one, ID = 0]
 - How many spectral windows (spw)in the ms? [one, ID = 0]
 - Center frequency? [146044.6047 MHz TOPO frame]
 - Bandwidth? [1.875 GHz]
 - Number of channels? [192]
 - Channel width? [-9.765625 MHz]
 - How many antennae in the array? [23]

Set up "CLEAN"

CASA <9>: inp clean interactive mode

# clean :: Invert	and	deconvolve	mages with selec	ted algorithm
vis	=	1.1	# Name of	input visibility file
imagename	=		<pre># Pre-name</pre>	of output images
outlierfile	=		# Text file	e with image names, sizes,
			# centers	for outliers
field	=		<pre># Field Na</pre>	me or id
spw	=		# Spectral	windows e.g. '0~3', '' is
			# all	
selectdata	-	True	# Other da	ta selection parameters
timerange	=		<pre># Range of</pre>	time to select from data
uvrange	=		# Select d	ata within uvrange
antenna	=		# Select d	ata based on antenna/baseline
scan	=		# Scan num	ber range
observation	=		# Observat	ion ID range
intent	=		# Scan Int	ent(s)
mode	-	'mfs'	# Spectral	gridding type (mfs, channel,
			<pre># velocit</pre>	y, frequency)
nterms	=	1	# Number o	f Taylor coefficients to
			<pre># model t</pre>	he sky frequency dependence
reffreq	=		# Referenc	e frequency (nterms > 1),''
			# uses ce	ntral data-frequency
gridmode	-		# Gridding	kernel for FFT-based
			# transfo	rms, default='' None
niter	=	500	# Maximum	number of iterations
gain	=	0.1	# Loop gai	n for cleaning

CASA <14>: clean?

user guide for help

Set up "CLEAN" (continuum imaging)

CASA <**9**>: inp clean

- Essential (necessary) keywords
 - vis = 'continuum.ms'
 - imagename = '116293_cont'
 - field = '0'
 - **s**pw = '0'
 - mode = 'mfs' # mfs means multi-frequency synthesis. We use this mode for continuum imaging.
 - **niter = 3000**
 - gain = 0.1 # this is the default, usually a good number
 - threshold = '0.9mJy'

Set up "CLEAN" (continuum imaging)

- Essential (necessary) keywords
 - psfmode = 'clark'
 - imagermode = 'csclean'
 - cell = ['0.15arcsec', '0.15arcsec']
 - imsize = [256,256]
 - weighting = 'natural' / weighting = 'uniform'
 - pbcor = False
 - when you are ready, type 'go'

Image visualization



Image visualization

CASA <**9**>: viewer



X

IRAS 16293-2422 Continuum



Set up "CLEAN" (line imaging)

CASA <8>: listobs vis='molecular_line.ms' CASA <9>: inp clean

- Essential (necessary) keywords
 - vis = 'molecular_line.ms'
 - imagename = 'OCS_12_11'
 - **field** = '0'
 - **s**pw = '0'
 - mode = 'velocity'
 - niter = 3000
 - **a** gain = 0.1
 - threshold = '5mJy'

Set up "CLEAN" (line imaging)

- Essential (necessary) keywords
 - psfmode = 'clark'
 - imagermode = 'csclean'
 - outframe = 'LSRK'
 - restfreq = '145946.815MHz'
 - cell = ['0.15arcsec','0.15arcsec']
 - imsize = [256,256]
 - weighting = 'natural' / weighting = 'uniform'
 - pbcor = False
 - when you are ready, type 'go'

IRAS 16293-2422 OCS 12-11



Excellent reference to read

- CASA cookbook, Chapter 5 (<u>https://casa.nrao.edu/</u> <u>casa_cookbook.pdf</u>)
- "CLEAN" algorithm (Högborn 1974; http://adsabs.harvard.edu/abs/1974A%26AS...15..417H)
- "Bible"
 - Interferometry and Synthesis in Radio Astronomy (<u>http://as.wiley.com/WileyCDA/WileyTitle/productCd-0471254924.html</u>)
 - Synthesis Imaging in Radio Astronomy II (<u>http://</u> <u>www.aspbooks.org/a/volumes/table_of_contents/?book_id=292</u>)

Quiz

- How to generate dirty image with CLEAN?
- Try to image "m51_mockObs.ms"
 - input hints
 - field='0~22'
 - imsize=[512,512]
 - cell = ['0.5arcsec','0.5arcsec']
 - phasecenter = '11'

