



RM-SYNTHESIS DEMO

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LOFAR

Overview

- Tutorial topics
 - General notes about RM-synthesis code
 - Input data preparation & inspection
 - Output data inspection
- Tutorial goals
 - Inspection of input/output data
 - Editing (both manual and automatic) of imaged data
 - Setup of an RM-synthesis run
 - Wide field imaging of data

CEP3 login and usage

- 6 working nodes reserved: lof006 lof010 lof012 lof[015 016 017]
- 25 active user accounts: lods01...lods27
- 1 active reservation with id=lofar_school2018_114
- Work in couples!
 - Username=lodsXX | working node=lof0XX
 - Accounts from lods01 to lods05 => lof006
 - Accounts from lods06 to lods10 => lof010
 - Accounts from lods06 to lods10 => lof012
 - Accounts from lods11 to lods15 => lof015
 - Accounts from lods16 to lods20 => lof016
 - Accounts from lods21 to lods25 => lof017

Log in to the LOFAR portal: **> ssh -Y lodsXX@portal.lofar.eu**

Log in to the head node: **> ssh -Y lodsXX@lhdhead.offline..lofar**

Activate a dummy session using the reservation id and the your node:

srun -A lofar_school2018 -reservation=lofar_school2018_114 -N 1 -w lof0XX -u bash -i

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Now, open a new terminal tab (keep the previous one open too!) and type: > **ssh -Y lodsXX@portal.lofar.eu**

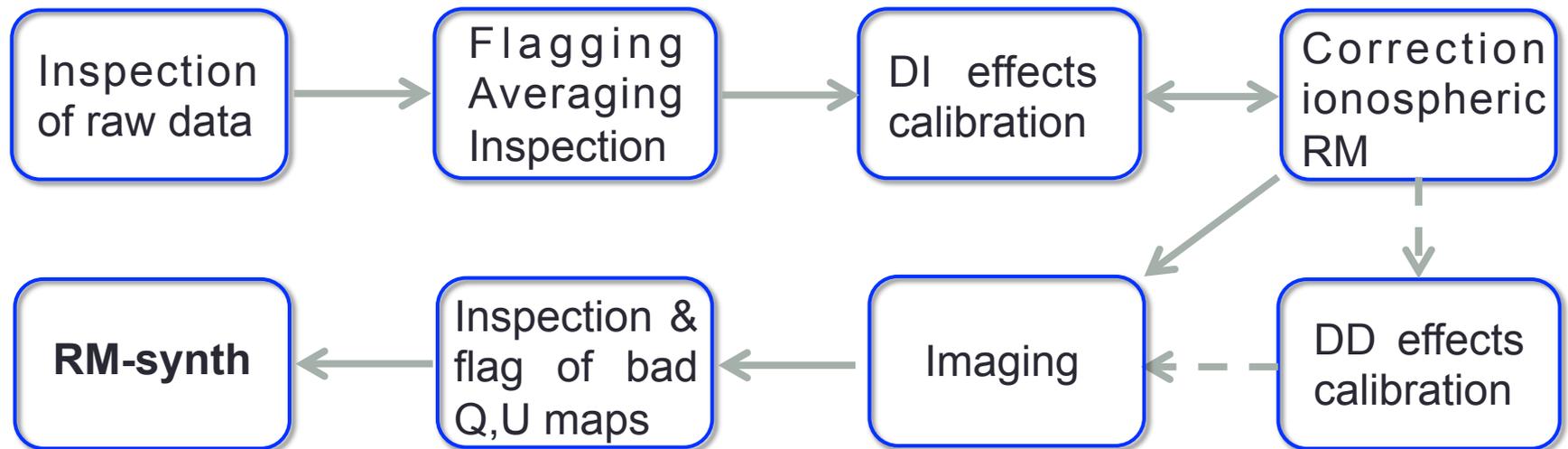
> **ssh -Y lodsXX@lhdhead.offline.lofar**

> **ssh -Y lodsXX@lof0XX**

- Verify that graphics forwarding works: > **geany**

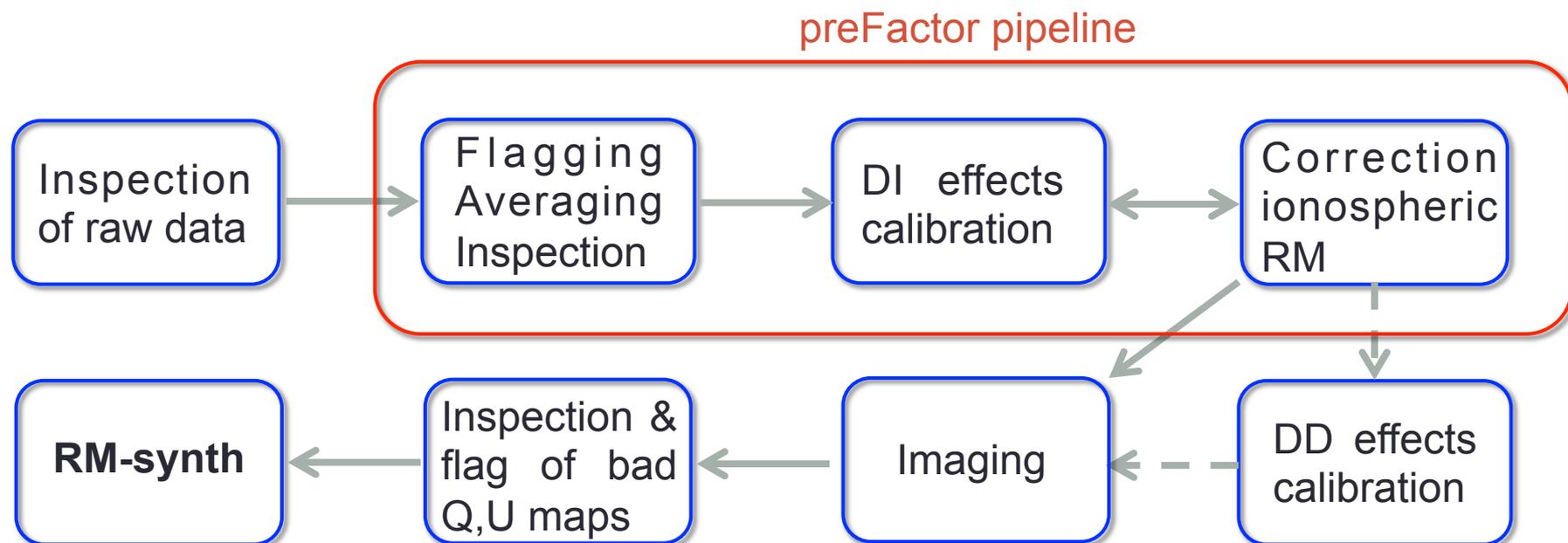
Input data preparation

- How to get there . .



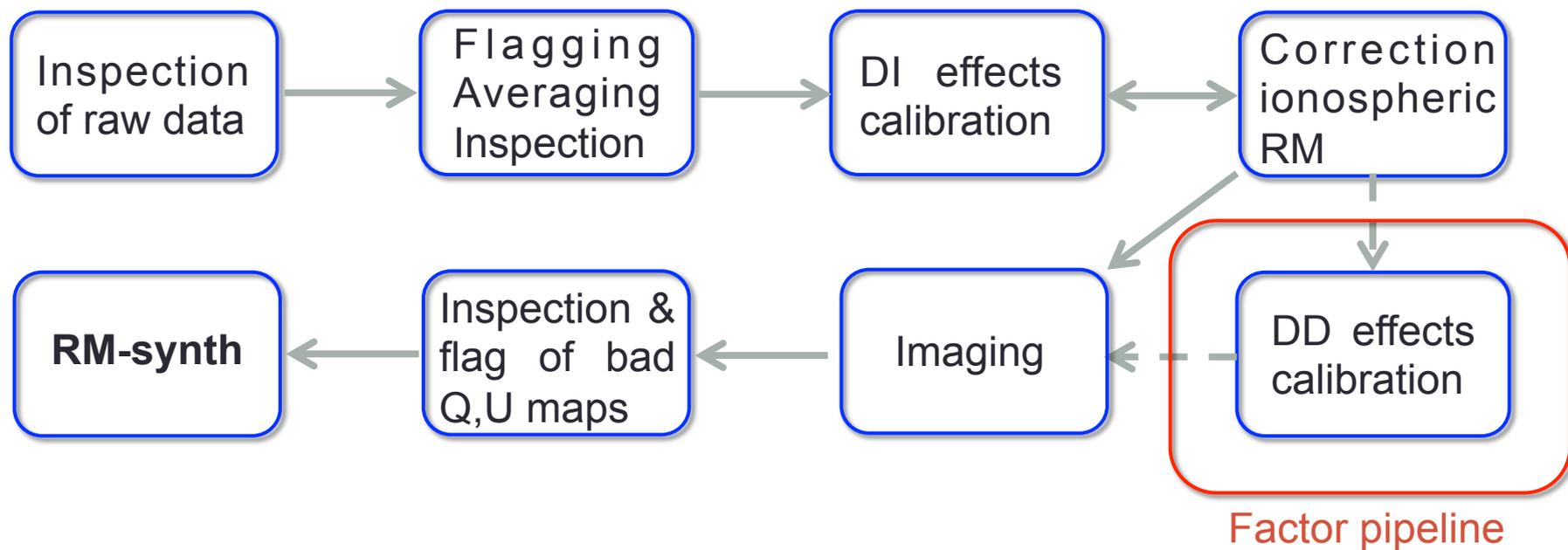
Input data preparation

- How to get there . .



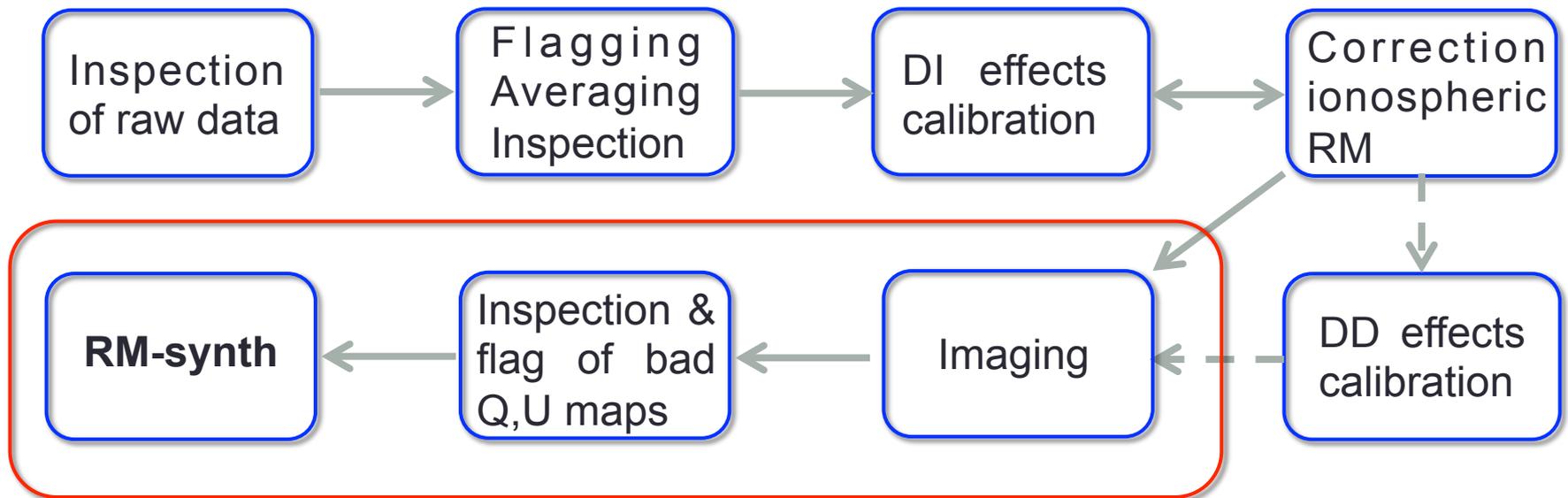
Input data preparation

- How to get there . .



Input data preparation

- How to get there . .

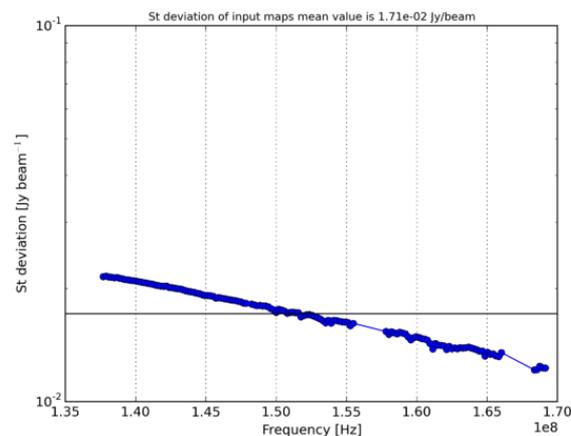
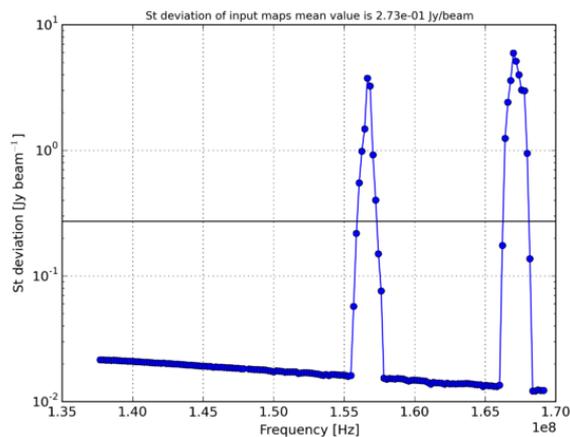


Input data preparation: imaging

- See imaging lecture L9 and tutorial T4
 - Which Stokes parameters ? **I Q U V**
 - Which parameters ? Npix, cellsize, pol clean
- Simplest way, a single parameter with no polarization cleaning
 - > **wsclean -name 10_deg_SB221_StokesU -size 1200 1200 -scale 30asec -weight briggs 0.0 -pol U -niter 0 -threshold 0.005 -mgain 0.8 -beamsize 165asec ../L123685_SB221_uv.DPPP.postionRMcorr**
 - Cleaning is performed in sum over channels of $Q_{ch}^2 + U_{ch}^2$. When doing RM-synthesis, this is the most sensible option.
 - > **wsclean -pol QU -join-polarizations -join-channels -squared-channel-joining -channels-out 100 ...**

Input data preparation: inspection

- Flagging of bad maps
 - Removal of **Q U** maps with anomalous variance is an option
 - Which Stoke parameter refer the plots below ?
 - Why a non flat variance ?



- Weighting of maps
 - Assign a weight to each **Q U** maps based on the inverse of their variance is also an option

General notes about pyrmsynth code

- Several software tools available. In this tutorial we will use some of the main software tool developed for LOFAR and low frequency data processing
 - <https://github.com/mrbell/pyrmsynth> A package that performs fast RM Synthesis and RM CLEAN imaging while still providing the flexibility of a Python interface.
 - The code works on sets of FITS files.
 - Each FITS file contains images from a single sub-band, or some other subset of the observed frequencies.
 - As a default, the code assumes all Stokes parameters (IQUV) to be saved in one FITS file. However it can also handle separately saved Q and U FITS files.
 - Collect all FITS files in a directory so the software will read them all in, stack them into a single data cube, and perform RM synthesis along each line of sight.

General notes about pyrmsynth code

- Command-line tool, input as a parset, output as feedback on screen.
- To source main software packages, type:
 - > **module load lofar**
 - Which packages you loaded ?
 - > **python rmsynthesis.py rmsynth.parset**

```
[iacobelli@lhd001]$ python pyrmsynth.py -h  
Usage: rmsynth.py <input parameter file>
```

Options:

- version show program's version number and exit
- h, --help show this help message and exit
- p, --plot_rmsf Plot the RMSF as soon as it is computed.
- V, --stokes_v Produce a Stokes V cube after reading the fits files.
- s, --separate_stokes
Indicate that the Stokes Q and U input images are stored in separate FITS files.
- f, --freq_last Indicate that NAXIS4 is the frequency axis.
- r, --rest_freq Indicate that the frequency for an image is given in the RESTFREQ header keyword.

General notes about pyrmsynth code

- How to set it up

Useful to focus on objects

```
% parameter file for rmsynthesis python code
% capable of running standard RM Synthesis as well as RM Clean

% Parameter file format:
% Comments can be added on their own lines, starting with a %, this must be followed by a space
% Parameters are given as keyword value pairs, with spaces as delimiters
```

```
% ra and dec min and max of the subimage to process, given in pixels
% a value of -1 means to use the bound of the image
dec_min -1 %200
dec_max -1 %400
ra_min -1 %200
ra_max -1 %400
```

```
% Define the phi axis, dphi in rad/m/m
phi_min -25 %-100
nphi 100 %200
dphi 0.25 %1
```

Useful to speed up runs

```
% Mask file. Pixels with non-zero values in the image will be used for RM Synthesis
% Comment the following line out if you don't wish to use a mask
% Mask image must have the same number of pixels as the Stokes-Q and U images
% irrespective of the ra, dec min and max values. The mask file should be
% located in the input_dir
%imagemask ./mask.fits
```

```
% Clean parameters. gain is the loop gain, niter is the number of clean iterations,
% cutoff sets the value of the max of the residual image at which point the procedure stops, defined in Jy
do_clean False
gain 0.1
niter 100
cutoff 0.0001
```

Useful to optimize S/N

```
% weighting parameter. Give the name of the weight file (located in the input_dir).
% If you leave it out, all channels will be given a weight of 1.0.
%do_weight weight.txt
```

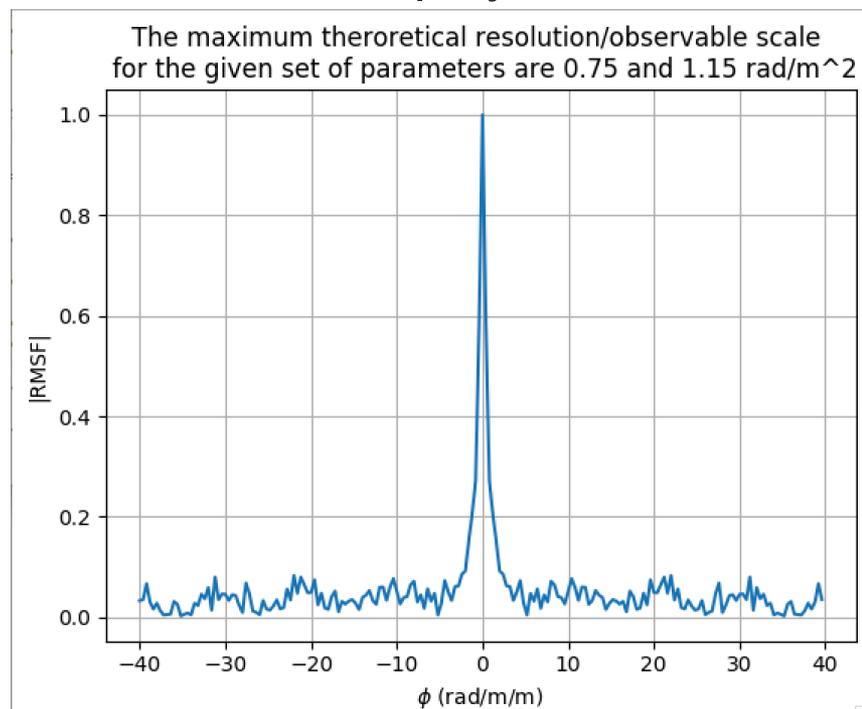
```
% spectral index option. Give directory or global average value. If wanted,
% specify reference frequency.
% alpha 0
%ref_freq
```

```
% Detection threshold on polarized intensity map
%threshold 0.1
```

```
% output file
outputfn /data/scratch/iacobelli/M33_DATA_CsOnly/RMSYNTHESIS/
% directory where the input fits file can be found
input_dir /data/scratch/iacobelli/M33_DATA_CsOnly/RMSYNTHESIS
```

Running RM-synthesis

- Data and scripts are in /home/iacobelli/RMDEMO so first make your working directory and enter it:
 - > `cd /data/scratch/lodsXX`
 - > `python myrmsynth.py -p -s rmsynth.par`
- While running a few useful outputs are saved and displayed:
 - the maximum theoretical resolution
 - the maximum observable scale
 - the RMSF values and profile
 - How the frequency coverage affects the RMSF side lobes ?



RM-synthesis: outputs inspection

- Data are in /home/iacobelli/RMDEMO

- Make and access your working directory:

> **cd /data/scratch/lodsXX**

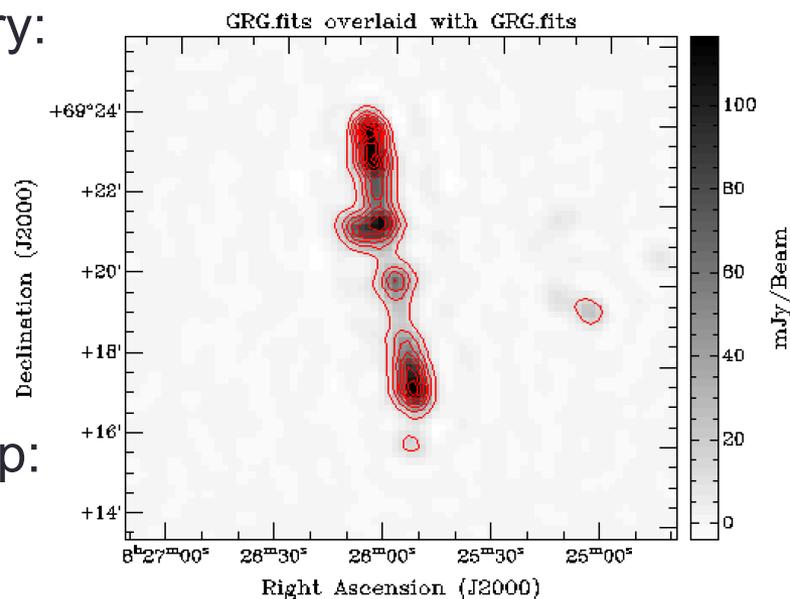
- To source fits viewer packages, type:

> **module load ds9** or

> **module load karma** (for kvis)

- First let us open the target Stokes I map:

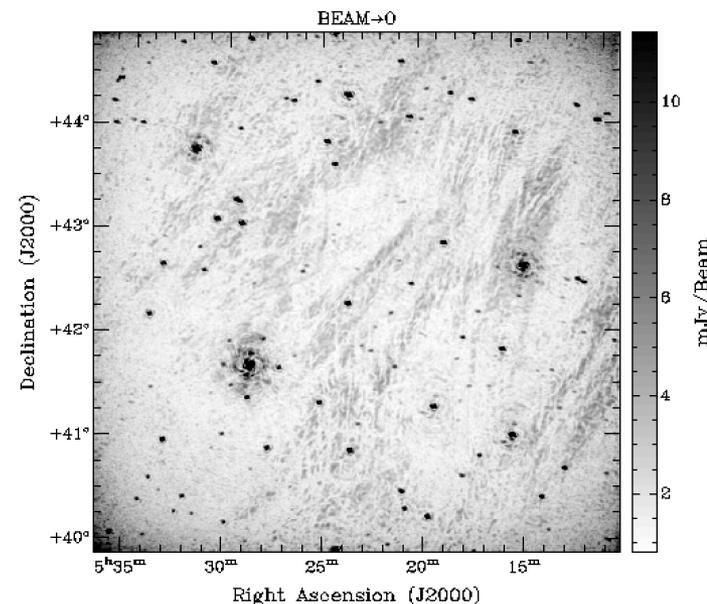
> **ds9 GRG.fits**



- Now let us check a fits cube to find the Faraday depth range selected in the experiment.
 - Check noise distributions; Gaussian for Q,U but not for PI: why ?
 - Find the instrumental polarization component and quantify it
 - Find the GRG in both RA,DEC and FD

RM-synthesis: outputs inspection

- Data are in /home/iacobelli/RMDEMO
 - Make and access your working directory:
 - > `cd /data/scratch/lodsXX`
 - To source fits viewer packages, type:
 - > `module load ds9` or
 - > `module load karma` (for kvis)
- Collapsing RM cubes allows to portrait the content of structures in both FD and PI. Let us open a map showing the peaks of PI:
 - > `ds9_polint.fits`
- Now let us check a fits cube showing Galactic foreground . .
 - > `ds9_di_p.fits`
 - Can you recognize how Stokes I looks like ?
 - Find the instrumental polarization component
 - Can you find any background polarized source ?





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