Using the IDIA MeerKAT Pipeline

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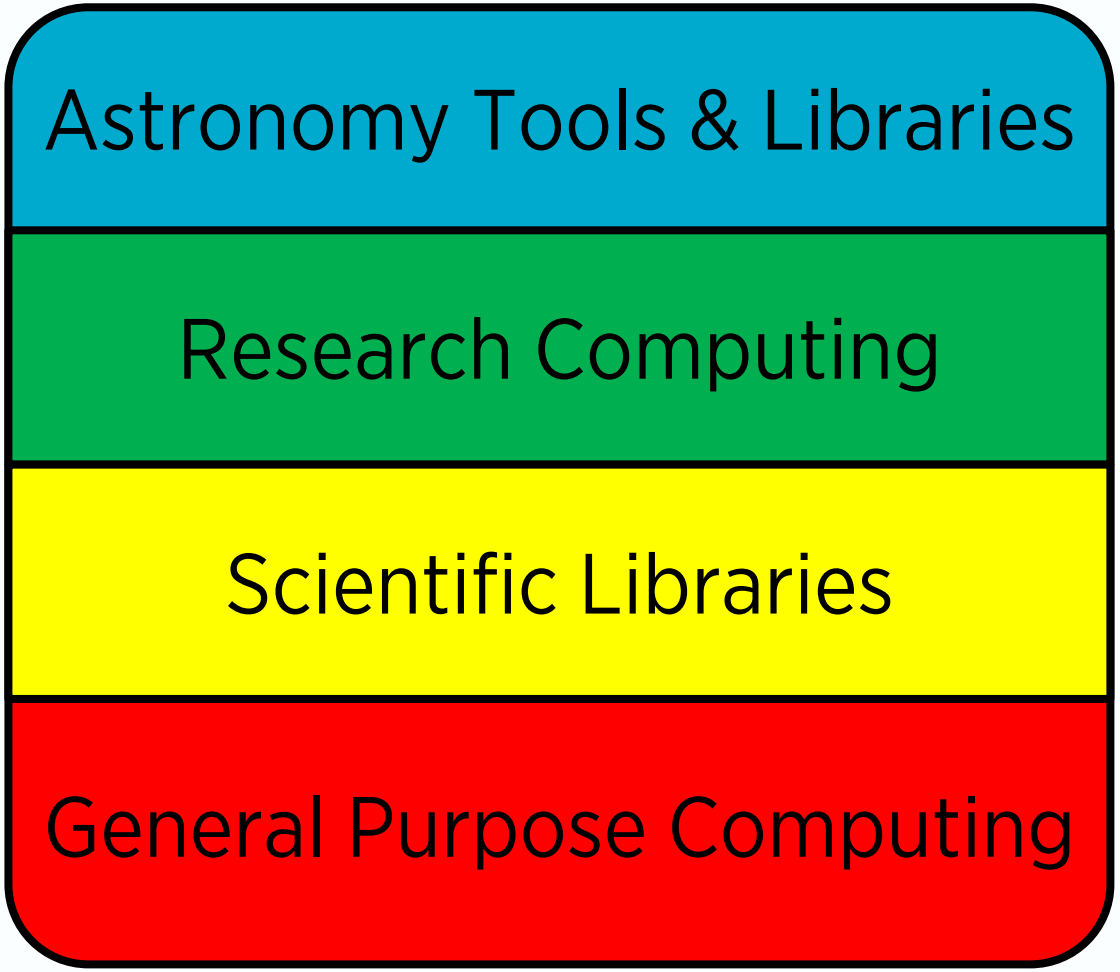
Jordan Collier | 20 Feb 2019 | MIGHTEE workshop

Our Systems: Software Stack



CASA

slurm workload manager

Open MPI

jupyterhub

Linux

openstack CLOUD SOFTWARE

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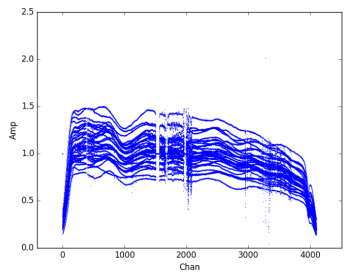
ssh access

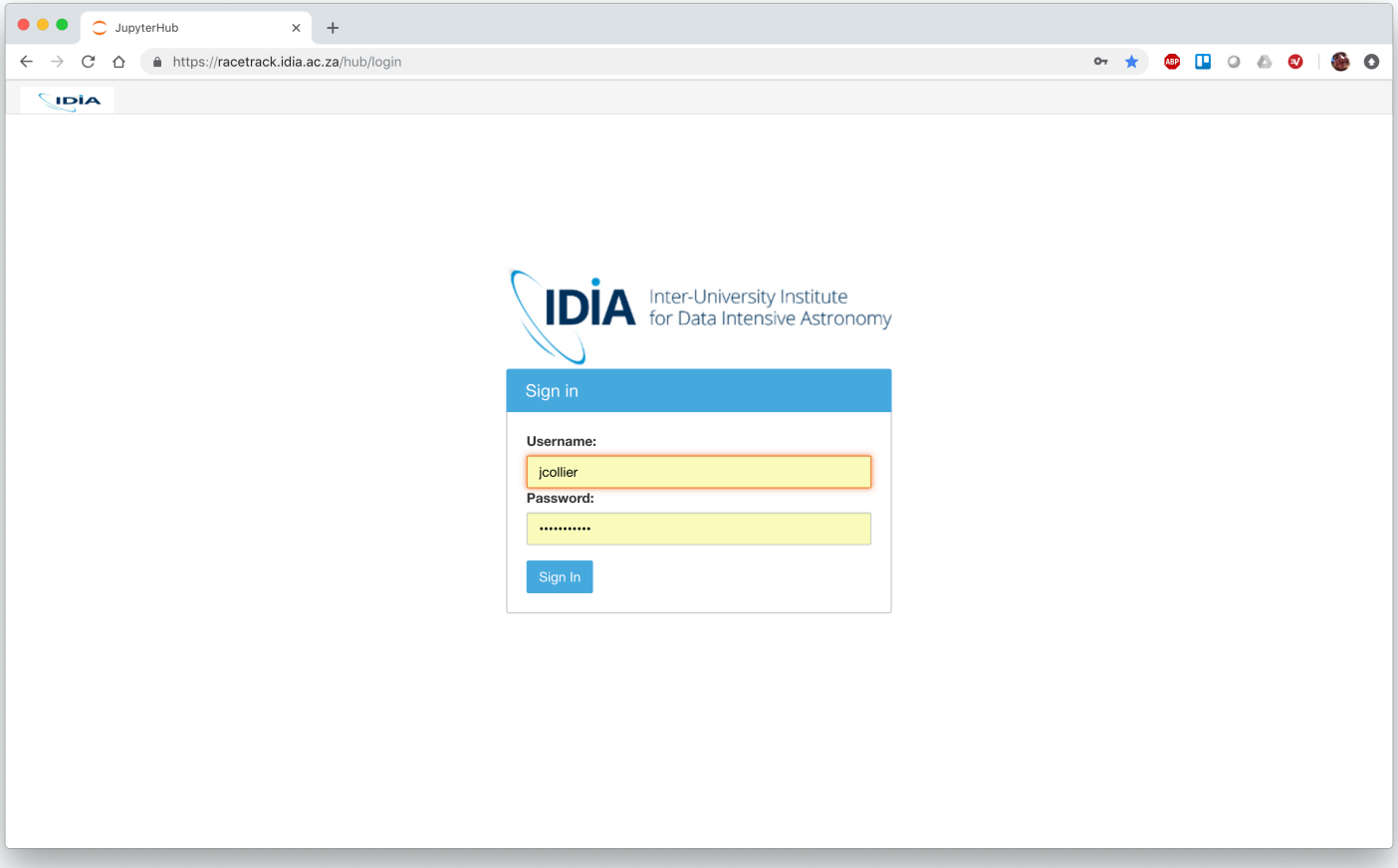
1. ssh

```
Warning: No xauth data; using fake authentication data for X11 forwarding.
Welcome to Ubuntu 16.04.5 LTS (GNU/Linux 4.4.0-137-generic x86_64)

 * Documentation:  https://help.ubuntu.com
 * Management:     https://landscape.canonical.com
 * Support:        https://ubuntu.com/advantage

  Get cloud support with Ubuntu Advantage Cloud Guest:
    http://www.ubuntu.com/business/services/cloud

58 packages can be updated.
3 updates are security updates.

New release '18.04.1 LTS' available.
Run 'do-release-upgrade' to upgrade to it.


*** System restart required ***
Last login: Mon Nov 26 02:19:05 2018 from 137.154.142.200
jcollier@racetrack:~$ ls data/CASA_testing/plots/
jcollier@racetrack:~$ ls data/CASA_testing/plot
plotcal.last  plotms.last   plots/
jcollier@racetrack:~$ ls data/CASA_testing/*/plots/
1491550051_bpass_gain.png   1491550051_bpass_real_imag.png                  1491550051_fastplot_phase.png  1491550051_gain_phase.png   1491550051_target_spectrum.png
1491550051_bpass_phase.png  1491550051_fastplot_amp.png                     1491550051_gain_amp.png        1491550051_target_spec.pdf
1491550051_bpass.png        1491550051_fastplot_phase_benchmarking_helo.png  1491550051_gain_Amp.png        1491550051_target_spec.png
jcollier@racetrack:~$ ls data/CASA_testing/*/plots/1491550051_bpass_gain.png
data/CASA_testing/1491550051/plots/1491550051_bpass_gain.png
jcollier@racetrack:~$ imgcaty data/CASA_testing/*/plots/1491550051_bpass_gain.png
imgcaty: command not found
jcollier@racetrack:~$ imgcat data/CASA_testing/*/plots/1491550051_bpass_gain.png
```

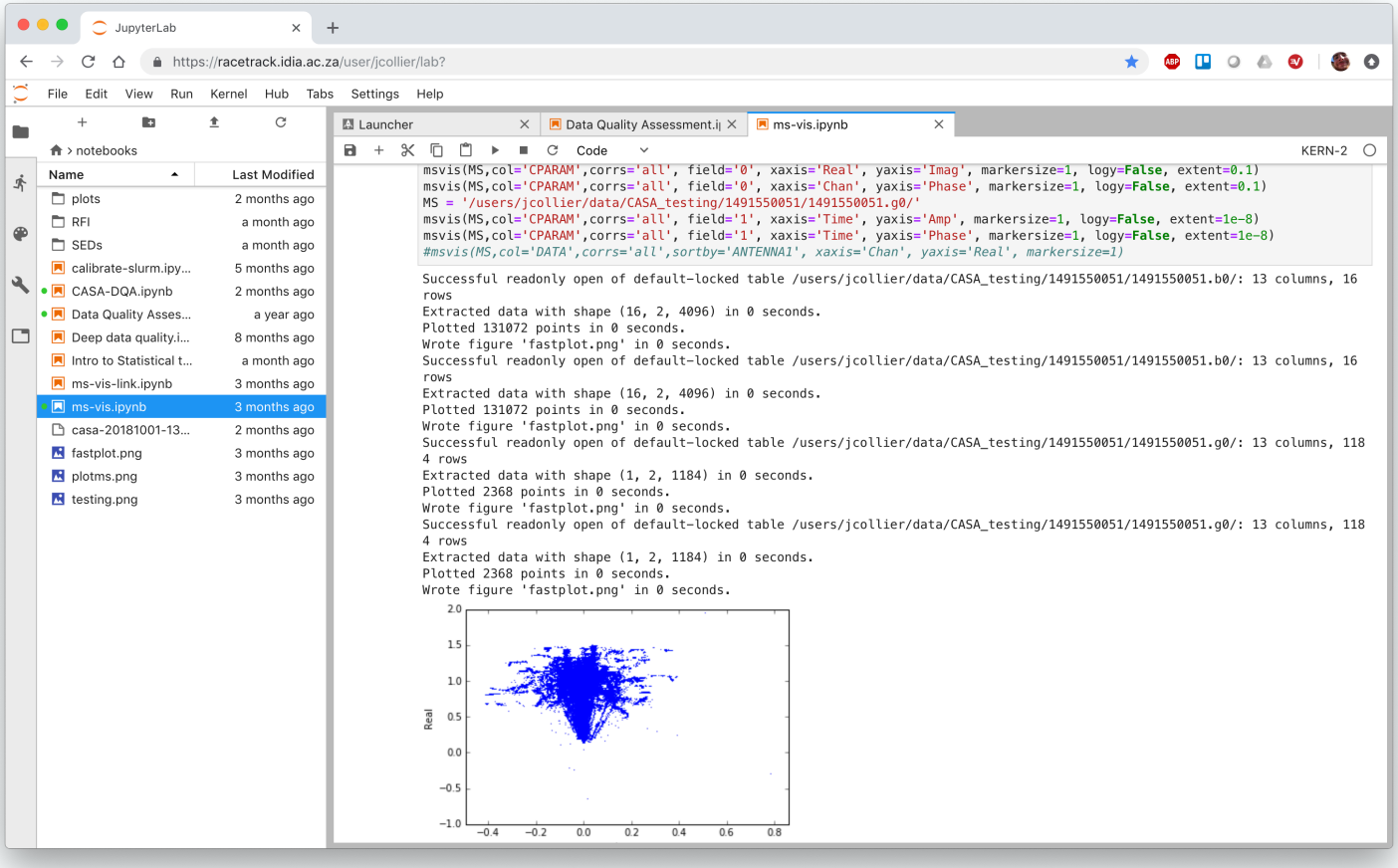



```
jcollier@racetrack:~$
```

JupyterHub


JupyterHub
https://racetrack.idia.ac.za/hub/login
IDIA
Inter-University Institute
for Data Intensive Astronomy
Sign in
Username:
jcollier
Password:
Sign In

JupyterHub


JupyterLab
https://racetrack.idia.ac.za/user/jcollier/lab?
File Edit View Run Kernel Hub Tabs Settings Help
notebooks
Name
Last Modified
plots 2 months ago
RFI a month ago
SEDs a month ago
calibrate-slurm.ipy... 5 months ago
CASA-DQA.ipynb 2 months ago
Data Quality Asses... a year ago
Deep data quality.i... 8 months ago
Intro to Statistical t... a month ago
ms-vis-link.ipynb 3 months ago
ms-vis.ipynb 3 months ago
casa-20181001-13... 2 months ago
fastplot.png 3 months ago
plotms.png 3 months ago
testing.png 3 months ago
Launcher
Data Quality Assessment.i
ms-vis.ipynb
Code
KERN-2
msvis(MS,col='CPARAM',corrs='all', field='0', xaxis='Real', yaxis='Imag', markersize=1, logy=False, extent=0.1)
msvis(MS,col='CPARAM',corrs='all', field='0', xaxis='Chan', yaxis='Phase', markersize=1, logy=False, extent=0.1)
MS = '/users/jcollier/data/CASA_testing/1491550051/1491550051.g0/'
msvis(MS,col='CPARAM',corrs='all', field='1', xaxis='Time', yaxis='Amp', markersize=1, logy=False, extent=1e-8)
msvis(MS,col='CPARAM',corrs='all', field='1', xaxis='Time', yaxis='Phase', markersize=1, logy=False, extent=1e-8)
#msvis(MS,col='DATA',corrs='all',sortby='ANTENNA1', xaxis='Chan', yaxis='Real', markersize=1)
Successful readonly open of default-locked table /users/jcollier/data/CASA_testing/1491550051/1491550051.b0/: 13 columns, 16 rows
Extracted data with shape (16, 2, 4096) in 0 seconds.
Plotted 131072 points in 0 seconds.
Wrote figure 'fastplot.png' in 0 seconds.
Successful readonly open of default-locked table /users/jcollier/data/CASA_testing/1491550051/1491550051.b0/: 13 columns, 16 rows
Extracted data with shape (16, 2, 4096) in 0 seconds.
Plotted 131072 points in 0 seconds.
Wrote figure 'fastplot.png' in 0 seconds.
Successful readonly open of default-locked table /users/jcollier/data/CASA_testing/1491550051/1491550051.g0/: 13 columns, 1184 rows
Extracted data with shape (1, 2, 1184) in 0 seconds.
Plotted 2368 points in 0 seconds.
Wrote figure 'fastplot.png' in 0 seconds.
Successful readonly open of default-locked table /users/jcollier/data/CASA_testing/1491550051/1491550051.g0/: 13 columns, 1184 rows
Extracted data with shape (1, 2, 1184) in 0 seconds.
Plotted 2368 points in 0 seconds.
Wrote figure 'fastplot.png' in 0 seconds.
Real
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−0.4
−0.2
0.0
0.2
0.4
0.6
0.8

Singularity Containers

- Led by Jeremy Smith (IDIA, UWC) – slide courtesy
- Containers are a software package that contains everything required to run an application/workflow
  - files, environmental variables, libraries and dependencies
- Allows for interchangeable environments and configurations at runtime and portability between systems

Singularity Containers

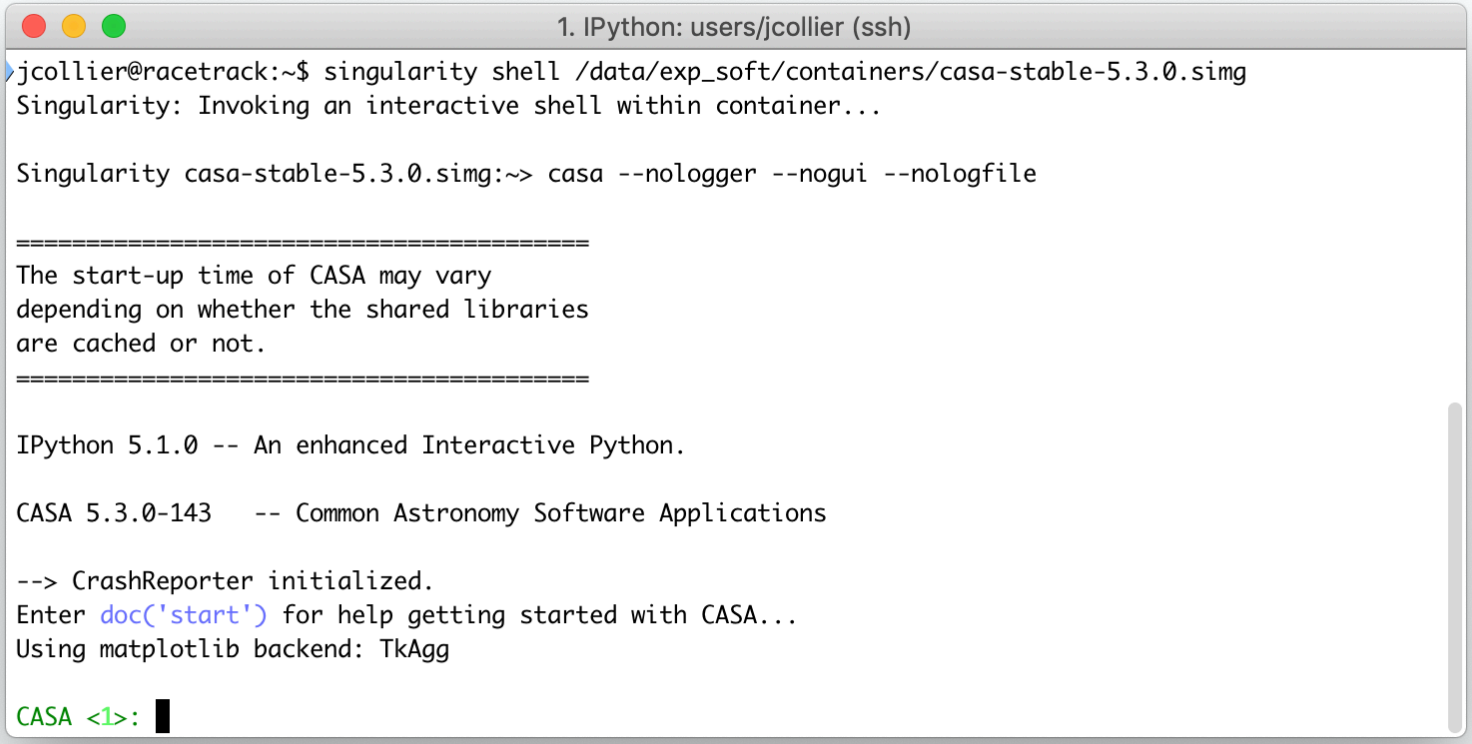
sourcefinding_py3_update.sh — ~/Research/MeerKAT/containers

sourcefinding_py3_update.sh

```
1  Bootstrap: localimage
2  From: /mnt/containers/sourcefinding_py3_update/sourcefinding_py3-2018-08-30-tmp.simg
3  Include: software-properties-common
4
5  %environment
6
7          export INSTALLER_PATH=/installer
8          ...
9
10 %setup
11         #Copy patch file for blobcat
12         cp blobcat_patch.diff $SINGULARITY_ROOTFS
13
14 %post
15
16         #Installation of BLOBCAT
17         export BLOBCAT_PATH=$INSTALLER_PATH/BLOBCAT
18         mkdir $BLOBCAT_PATH
19         cd $BLOBCAT_PATH
20         wget -O blobcat-v1.2.tar.bz2 http://sourceforge.net/projects/blobcat/files/blobcat-v1.2.tar.bz2/download
21         tar xvjf blobcat-v1.2.tar.bz2
22
23         #Patch blobcat to run python2.7 and fix error on one line
24         patch blobcat.py /blobcat_patch.diff
25
26         ...
27
28         #Upgrade pip
29         pip install --upgrade pip
30
31         #Installation of PASTA
32         pip install git+https://github.com/bwkeller/PASTA
33
34         ...
35
36         #Install eog (and dependencies)
37         apt-get -y install eog
```

sourcefinding_py3_update.sh 1:1 — 1 LF UTF-8 Shell Script 0 files 4 updates

Singularity Containers



Singularity Containers

- Exec

```
:~$ singularity exec /data/exp_soft/containers/sourcefinding_py3.simg python myscript.py
```

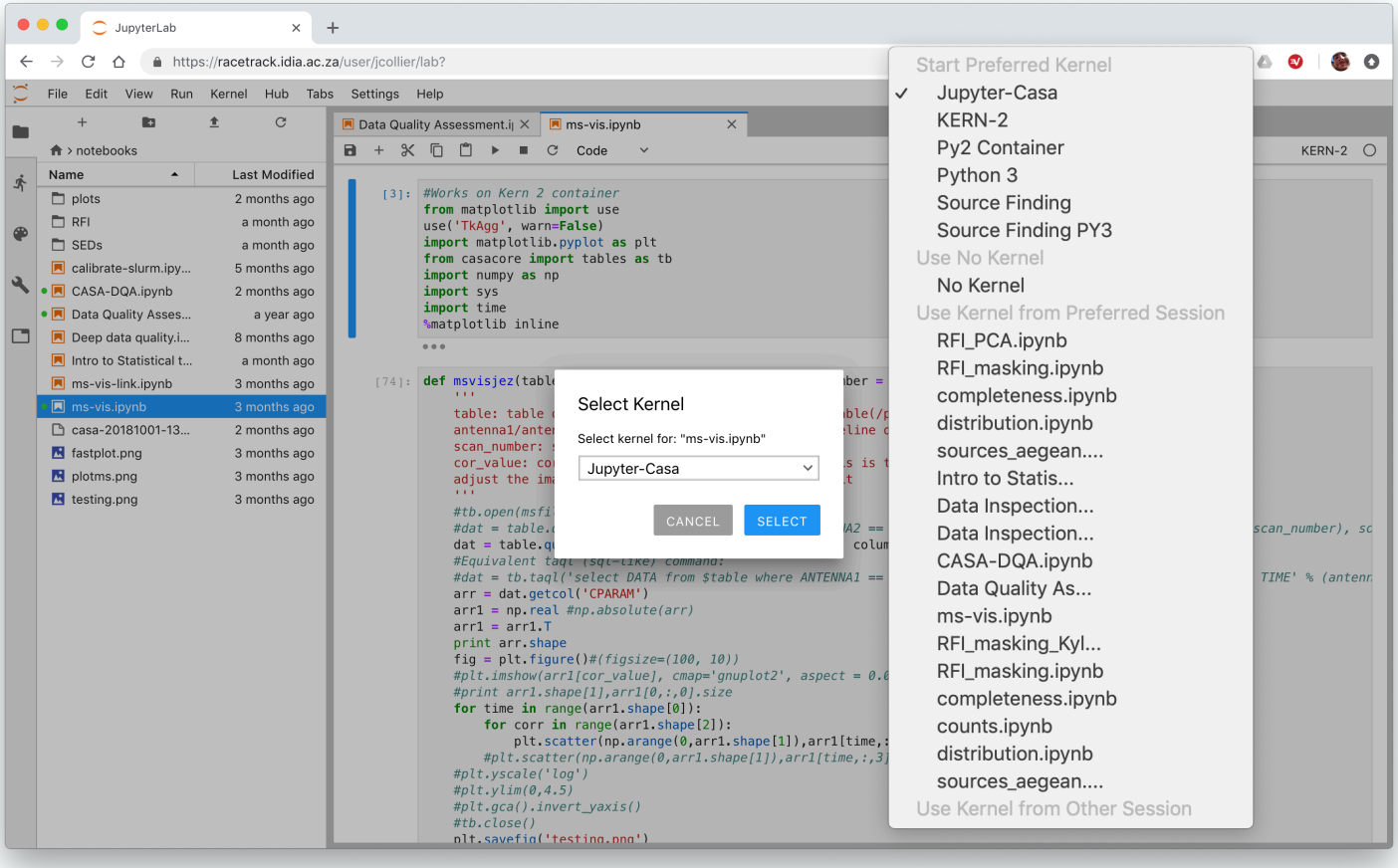
- MPICASA with SLURM using batch file (“my-casa-job.sh”)

```
#!/bin/bash
#SBATCH --nodes=15
#SBATCH --ntasks-per-node=8
#SBATCH --cpus-per-task=3
#SBATCH --mem=98304
#SBATCH --job-name=tclean
#SBATCH --distribution=plane=4
#SBATCH --output=logs/tclean-%j.out
#SBATCH --error=logs/tclean-%j.err

/path/to/mpicasa /usr/bin/singularity exec /data/exp_soft/pipelines/casameer-5.4.0.simg "casa"
--nologger --nogui --logfile logs/tclean-${SLURM_JOB_ID}.casa -c tclean.py --config .config.tmp
```

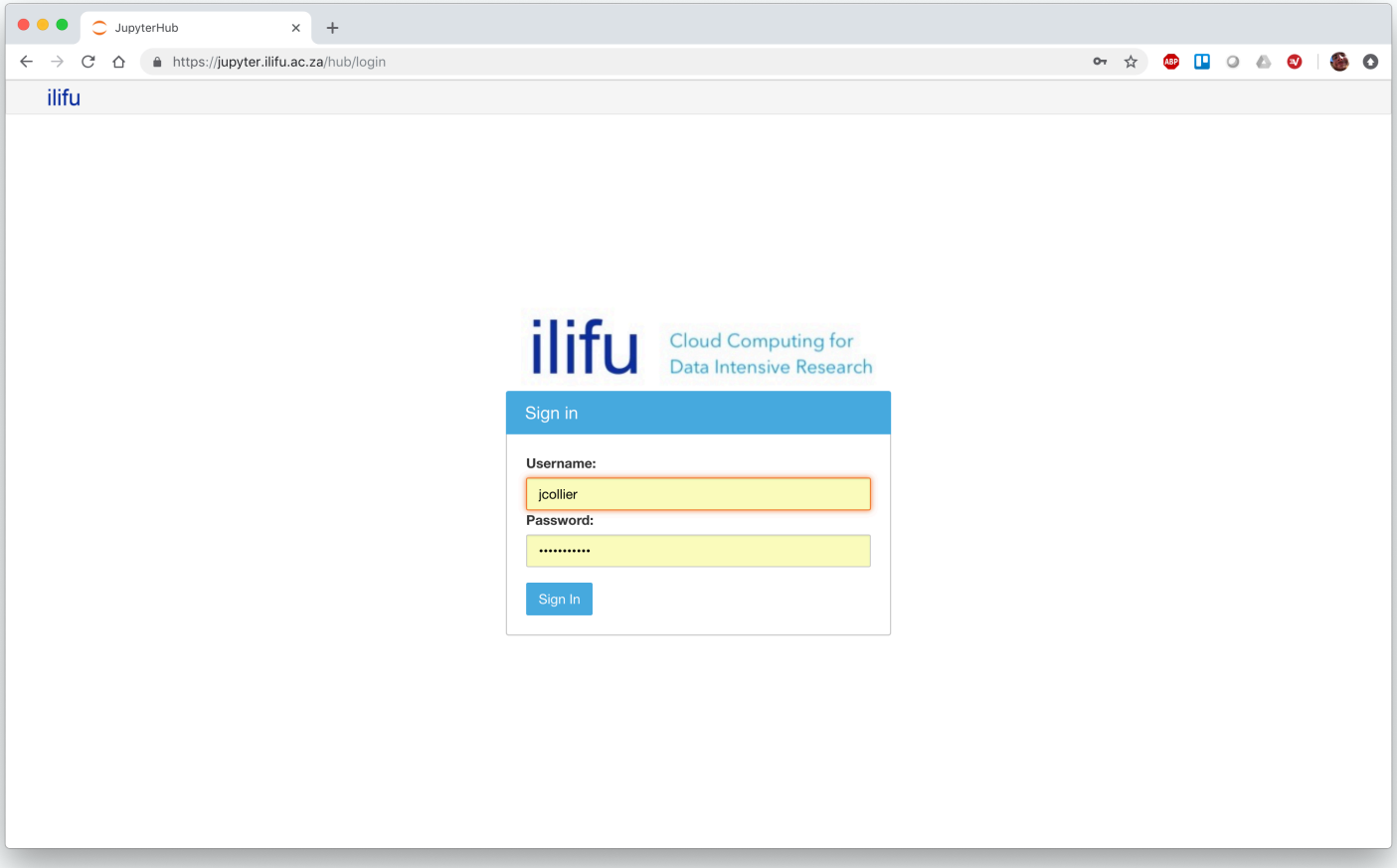
- Terminal on SLURM cluster:

```
user@master001:~> sbatch my-casa-job.sh
```

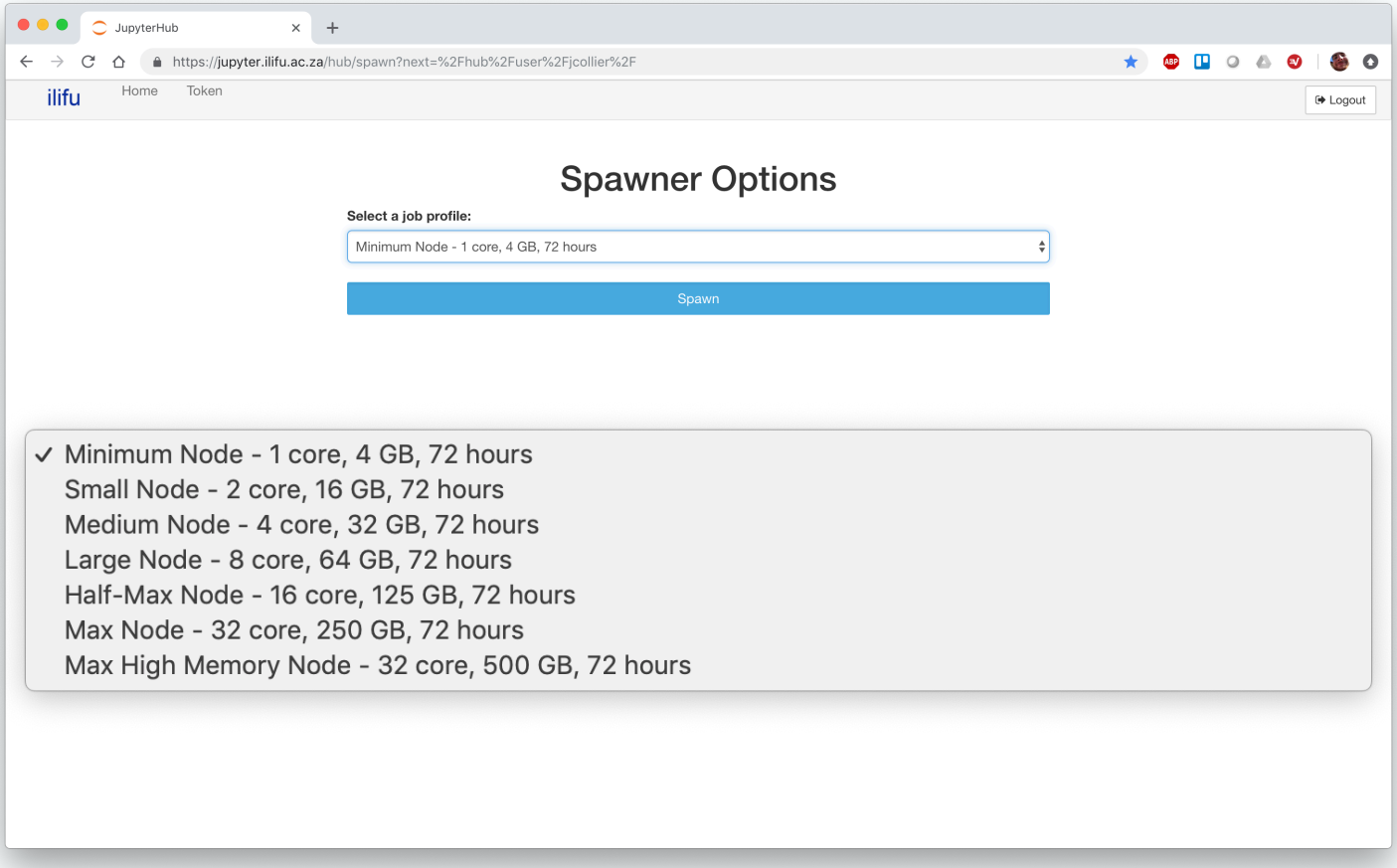
JupyterHub: Singularity Containers



JupyterHub Spawner



JupyterHub Spawner



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processMeerKAT.py

- Builds and submits pipeline job on IDIA / Ilifu SLURM cluster
- Launch pipeline
  - Input measurement set
  - Build your config file
  - OR run your config file (i.e. build job scripts)
  - Request resources
  - Optionally insert your own scripts
  - Specify containers and MPI wrappers

1. bash

```
MC02RTBRAG8WP:~ jordan$ processMeerKAT.py -h
usage: /Users/jordan/Research/MeerKAT/IDIA/pipelines/processMeerKAT/processMeerKAT.py
       [-h] [-M path] [-C path] [-N num] [-t num] [-P num] [-m num] [-p name]
       [-T time] [-S script threadsafe container] [-w path] [-c path] [-l]
       [-s] [-v] (-B | -R | -V)

Process MeerKAT data via CASA measurement set. Version: 1.0

optional arguments:
  -h, --help            show this help message and exit
  -M path, --MS path    Path to measurement set.
  -C path, --config path
                        Path to config file.
  -N num, --nodes num   Use this number of nodes [default: 8; max: 35].
  -t num, --ntasks-per-node num
                        Use this number of tasks (per node) [default: 4; max:
                        128].
  -P num, --plane num   Distribute tasks of this block size before moving onto
                        next node [default: 2; max: ntasks-per-node].
  -m num, --mem num     Use this many GB of memory (per node) for threadsafe
                        scripts [default: 236; max: 236.
  -p name, --partition name
                        SLURM partition to use [default: 'Main'].
  -T time, --time time  SLURM partition to use [default: 'Main'].
  -S script threadsafe container, --scripts script threadsafe container
                        Run pipeline with these scripts, in this order, using
                        this container (3rd value - empty string to default to
                        [-c --container]). Is it threadsafe (2nd value)?
  -w path, --mpi_wrapper path
                        Use this mpi wrapper when calling threadsafe scripts
                        [default: '/data/exp_soft/pipelines/casa-
                        prerelease-5.3.0-115.el7/bin/mpicasa'].
  -c path, --container path
                        Use this container when calling scripts [default:
                        '/data/exp_soft/pipelines/casameer-5.4.1.xvfb.simg'].
  -l, --local           Build config file locally (i.e. without calling srun)
                        [default: False].
  -s, --submit          Submit jobs immediately to SLURM queue [default:
                        False].
  -v, --verbose         Verbose output? [default: False].
  -B, --build           Build config file using input MS.
  -R, --run             Run pipeline with input config file.
  -V, --version         Display the version of this pipeline and quit.
```

The Build Step

- `processMeerKAT.py -B -C myconfig.txt –M /data/projects/mightee/MS/1524147354_sdp_l0.full.full_pol.MS`
- Builds your config file
  - Reads MS and automatically extracts field IDs based on intent
    - Extracts total flux, bandpass, phase cal, and targets
  - Compares resources you've requested to the number of scans / sub-MSs
    - We partition by scan, so it prevents requesting too many / few threads
  - Ensures reference antenna exists

Config Files

- [crosscal]
  - Select spectral window / freqs (e.g. for HI)
  - Other parameters affecting calibration
- [slurm]
  - Resource parameters
  - List of scripts to run, and whether to call MPI
  - etc
- Path to MS stored
- Field IDs stored

myconfig.txt

```
[crosscal]
minbaselines = 4                    # Minimum number of baselines to use while calibrating
specavg = 1                         # Number of chans to avg after calibration
timeavg = '8s'                      # Time interval to avg after calibration
spw = '0:860~1700MHz'               # spw to use for calibration
calcrefant = True                   # Calculate reference antenna in program (overwrites 'refant')
refant = 52                         # Reference antenna name/number
standard = 'Perley-Butler 2010'     # Flux density standard for setjy
badants = [2, 3, 5, 13, 19, 53]     # List of bad antenna numbers in the MS
badfreqranges = [ '944~947MHz',     # List of bad frequency ranges
                  '1160~1310MHz',
                  '1476~1611MHz',
                  '1670~1700MHz']

[slurm]
nodes = 8
ntasks_per_node = 2
mem = 236
plane = 1
submit = False
container = '/data/exp_soft/pipelines/casameer-5.4.1.xvfb.simg'
mpi_wrapper = '/data/exp_soft/pipelines/casa-prerelease-5.3.0-115.el7/bin/mpicasa'
partition = 'Main'
time = '12:00:00'
verbose = False
scripts = [('setjy.py', True, ''),
           ('xx_yy_solve.py', False, ''),
           ('xx_yy_apply.py', True, ''),
           ('flag_round_2.py', True, ''),
           ('setjy.py', True, ''),
           ('xy_yx_solve.py', False, ''),
           ('xy_yx_apply.py', True, ''),
           ('split.py', True, ''),
           ('plot_solutions.py', False, ''),
           ('quick_tclean.py', True, '')]

[data]
vis = '/data/projects/mightee/MS/1523541036_sdp_l0.full.full_pol.ms'

[fields]
bpassfield = '0'
fluxfield = '0'
phasecalfield = '1'
targetfields = '2'
```

The Run Step

- `processMeerKAT.py -R -C myconfig.txt`
- Builds SLURM sbatch files (one for each unique python script)
  - `Wrote sbatch file "validate_input.sbatch"`
  - `Wrote sbatch file "partition.sbatch"`
  - `Wrote sbatch file "calc_refant.sbatch"`
  - `Wrote sbatch file "flag_round_1.sbatch"`
  - `Wrote sbatch file "setjy.sbatch"`
  - `Wrote sbatch file "xx_yy_solve.sbatch"`
  - `Wrote sbatch file "xx_yy_apply.sbatch"`
  - `...`
  - `Wrote sbatch file "split.sbatch"`
  - `Wrote sbatch file "plot_solutions.sbatch"`
  - `Wrote sbatch file "quick_tclean.sbatch”`
- Writes master submission (bash) script

Sbatch Files

- Calls script within CASA container, within MPI wrapper
- Logs with job ID written for stdout, stderr, and CASA
- User can choose to submit these jobs one at a time
- OR use master submission script

partition.sbatch

```
#!/bin/bash
#SBATCH --nodes=8
#SBATCH --ntasks-per-node=2
#SBATCH --cpus-per-task=1
#SBATCH --mem=236GB
#SBATCH --job-name=partition
#SBATCH --distribution=plane=1
#SBATCH --output=logs/partition-%j.out
#SBATCH --error=logs/partition-%j.err
#SBATCH --partition=Test02
#SBATCH --time=12:00:00

export OMP_NUM_THREADS=1

/data/exp_soft/pipelines/casa-prerelease-5.3.0-115.el7/bin/
mpicasa singularity exec /data/exp_soft/pipelines/
casameer-5.4.1.xvfb.simg xvfb-run -d casa --nologger --nogui
--logfile logs/partition-${SLURM_JOB_ID}.casa -c
/data/exp_soft/pipelines/dev/processMeerKAT/cal_scripts/
partition.py --config .config.tmp
```

Master Submission Script

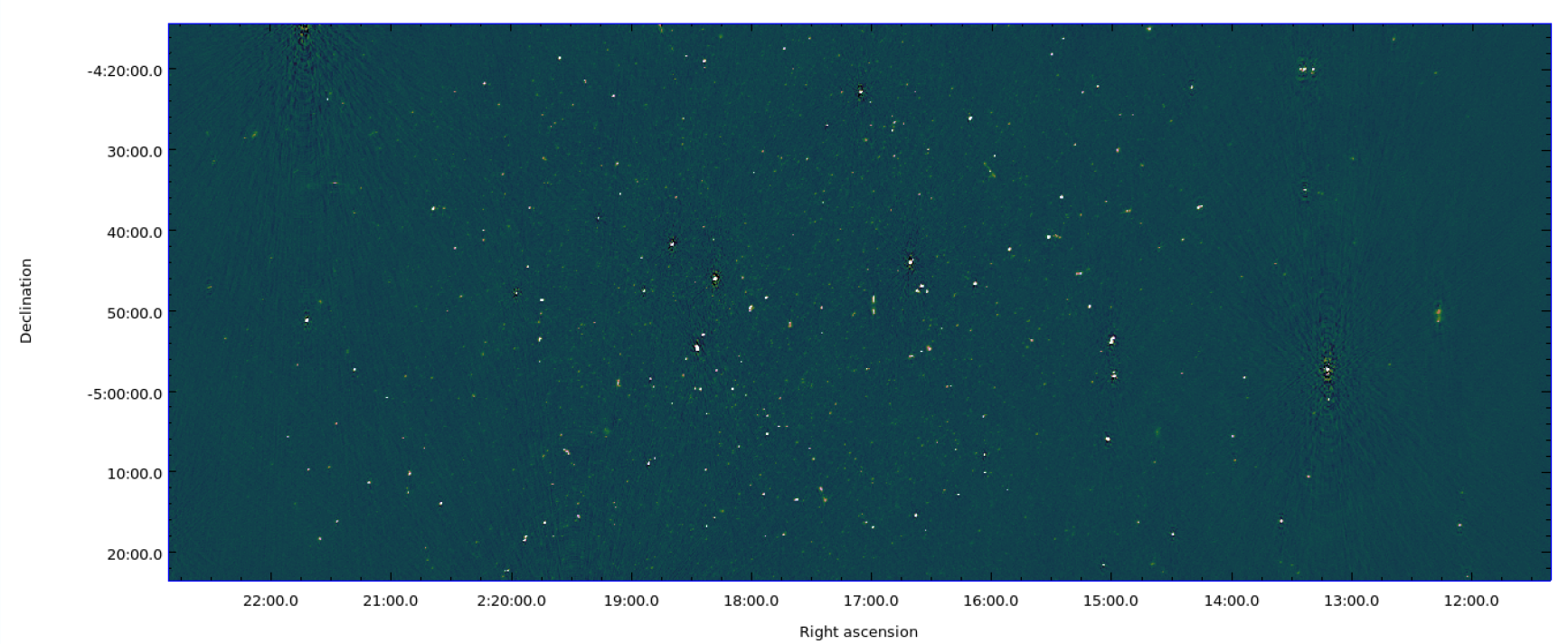
- Bash script that submits sbatch files to SLURM queue each time it's run
- Builds ancillary bash scripts that interact with pipeline run
  - Summarise progress
  - Kill all jobs
  - Find errors (after pipeline run)
  - Display runtimes (after pipeline run)

submit_pipeline.sh

```
#!/bin/bash
cp myconfig.txt .config.tmp

#setjy.sbatch
IDs=$(sbatch setjy.sbatch | cut -d ' ' -f4)

#xx_yy_solve.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes xx_yy_solve.sbatch | cut -d ' ' -f4)

#xx_yy_apply.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes xx_yy_apply.sbatch | cut -d ' ' -f4)

#flag_round_2.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes flag_round_2.sbatch | cut -d ' ' -f4)

#setjy.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes setjy.sbatch | cut -d ' ' -f4)

#xy_yx_solve.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes xy_yx_solve.sbatch | cut -d ' ' -f4)

#xy_yx_apply.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes xy_yx_apply.sbatch | cut -d ' ' -f4)

#split.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes split.sbatch | cut -d ' ' -f4)

#plot_solutions.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes plot_solutions.sbatch | cut -d ' ' -f4)

#quick_tclean.sbatch
IDs+=,$(sbatch -d afterok:$IDs --kill-on-invalid-dep=yes quick_tclean.sbatch | cut -d ' ' -f4)

#Output message and create jobScripts directory
echo Submitted sbatch jobs with following IDs: $IDs
mkdir -p jobScripts
```

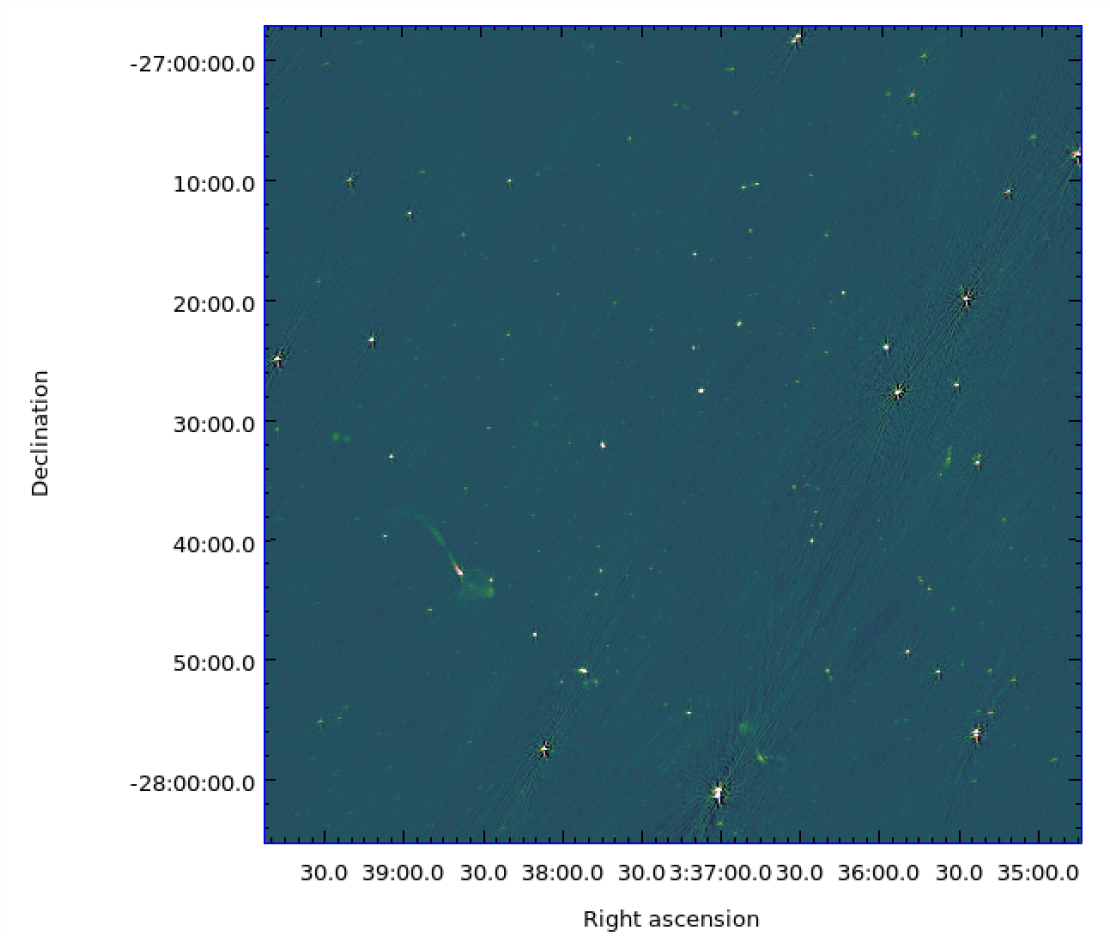
Initial QA: Quick Look Images

- Very quick and dirty imaging for QA purposes
  - No selfcal, no w-projection, no thresholding, no multi-scale, etc
  - XMM-LSS field: RMS ~10 uJy / beam



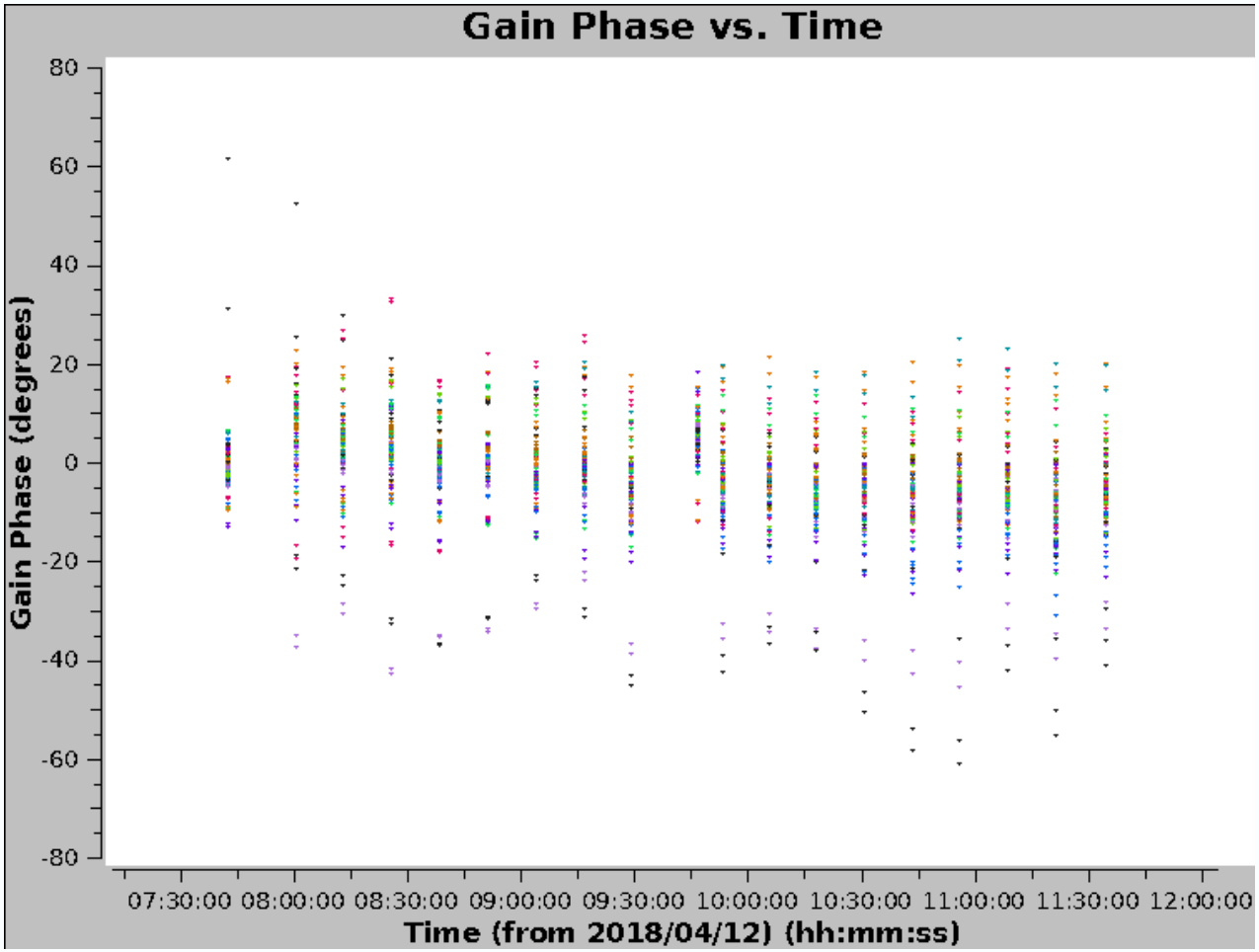
Initial QA: Quick Look Images

- CDFS field
- ~8 hours, 10 MHz spw
- RMS ~80 uJy / beam
- Scales to ~10 uJy over whole band (assuming ~100 MHz flagged out)



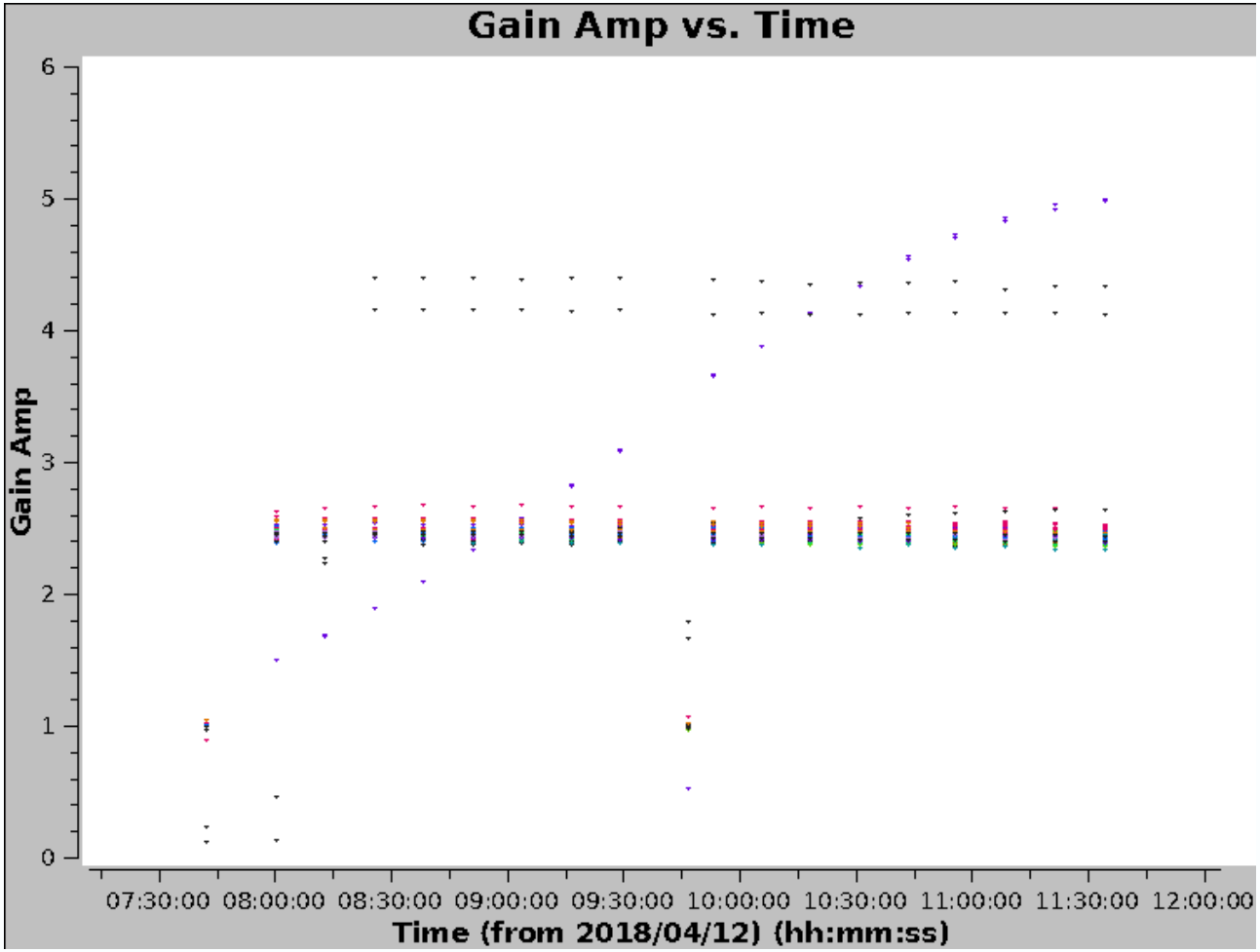
Initial QA: Calibration Solutions

- Phase cal solutions



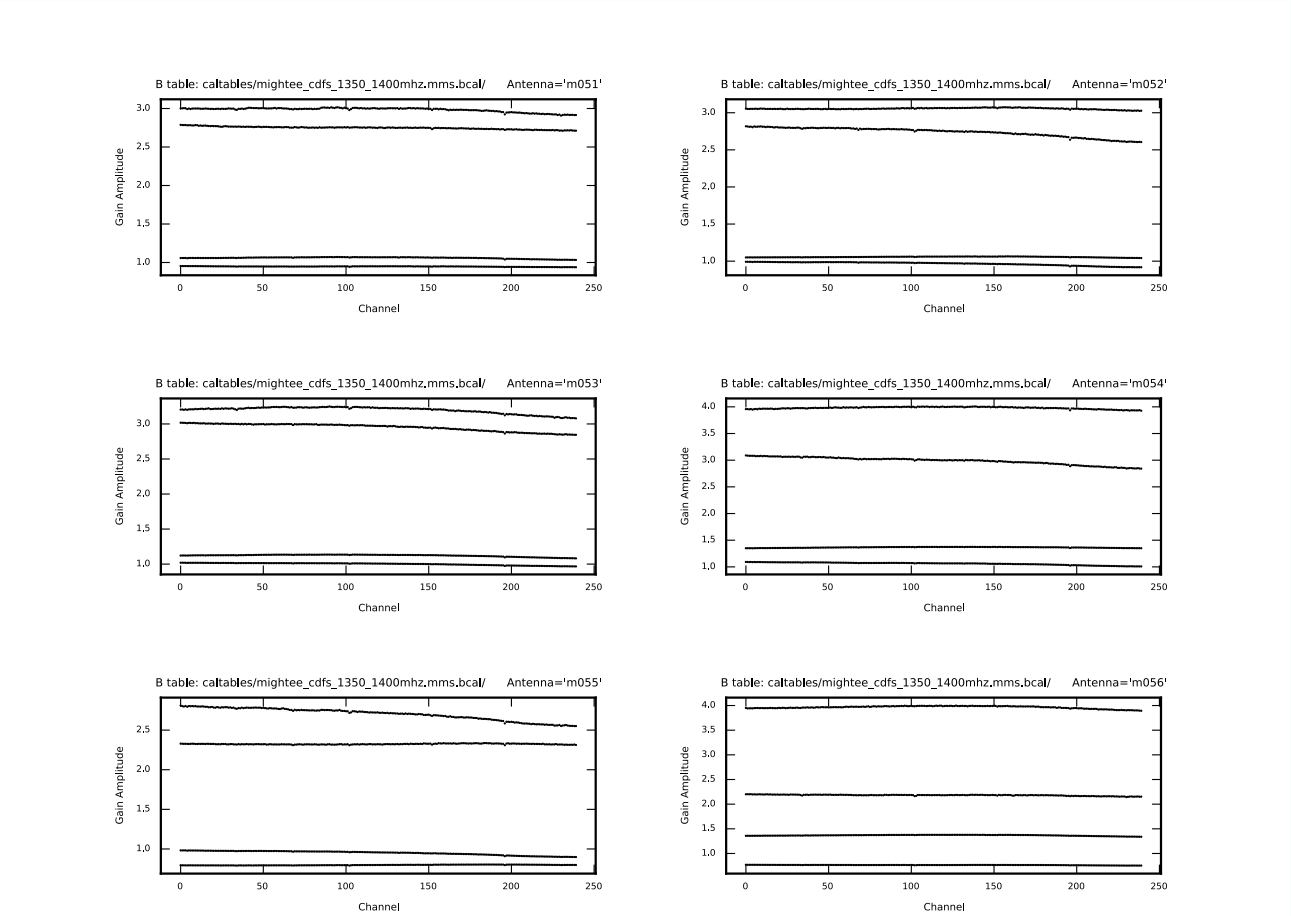
Initial QA: Calibration Solutions

- Phase cal solutions



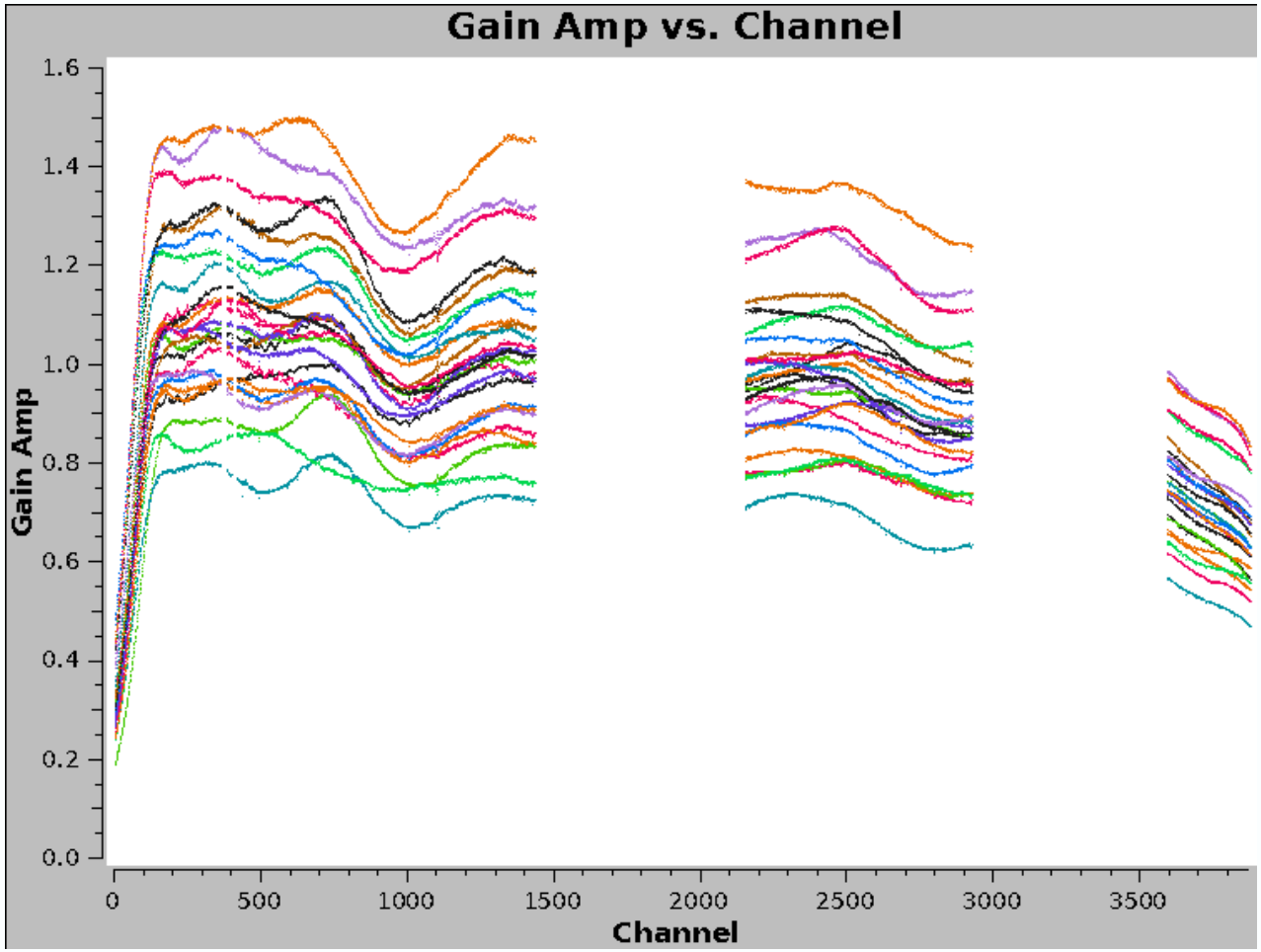
Initial QA: Calibration Solutions

- Phase cal solutions



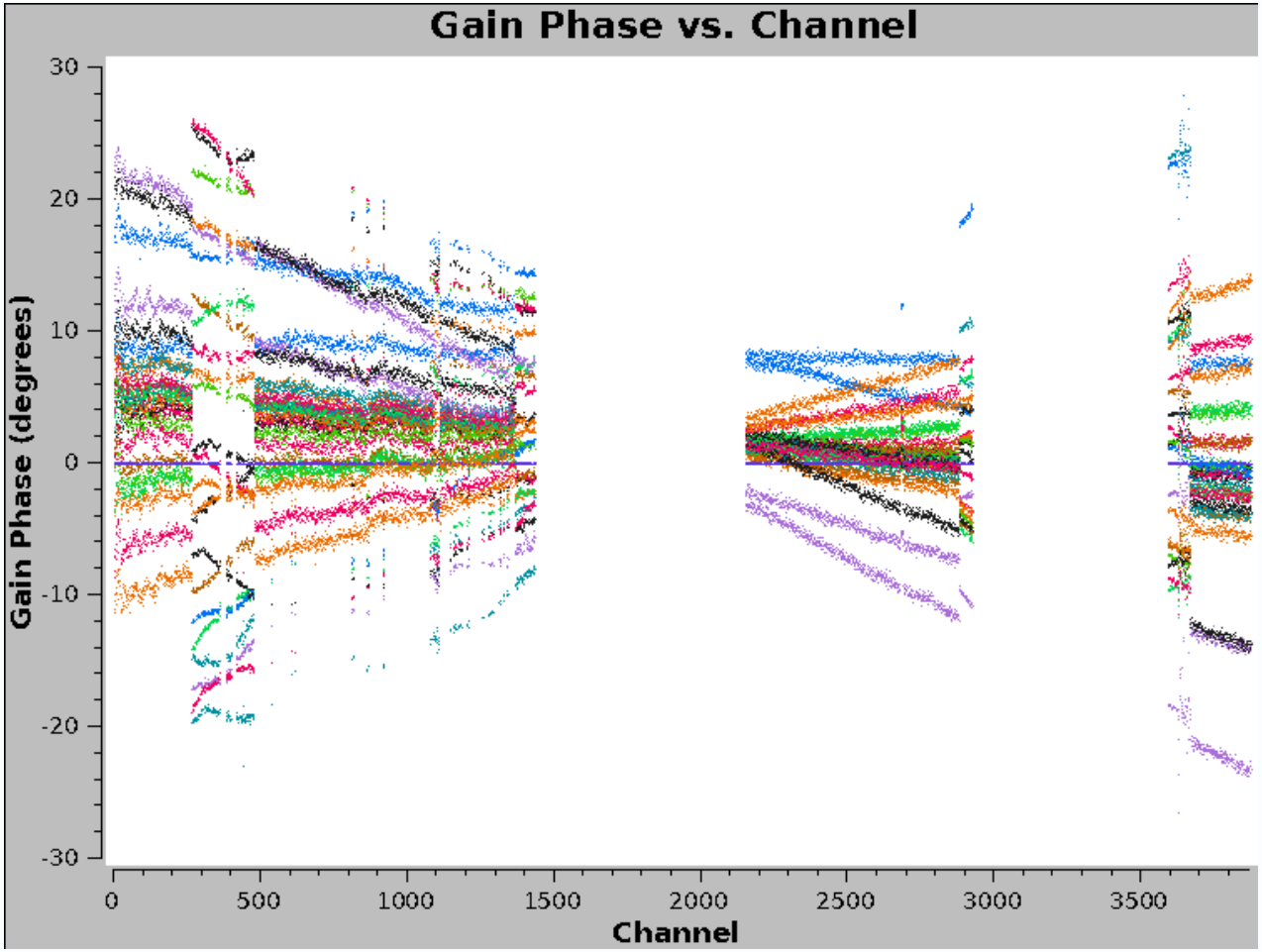
Initial QA: Calibration Solutions

- Bandpass solutions



Initial QA: Calibration Solutions

- Bandpass solutions
- And more...



A Good Framework

- Pipeline outputs calibrated MMS + MSs/MMSs split by field
- You insert your scripts at start, middle or end (e.g. WSClean)
- Each job/script is a logical step that does/doesn't use MPI, and optionally uses a different container
- HPC-friendly – dynamically uses resources & submits to queue
- Use cases we currently support
  - Full stokes calibration + Stokes I only calibration (minimal speedup)
  - Narrow band (spectral line) calibration, full-band calibration
  - Single MS (speedup for small BW), multi-MS
  - Inserting your own scripts (hard-coded or read config file)
- https://idia-pipelines.github.io/docs/processMeerKAT

Future Development

- Selfcal and AW Projection (see Krishna's talks)
- Optimisation of resources / performance
  - Currently takes ~1 day to process 64 dish 4k data (~2 TB)
  - Split data by intent during beginning, and simultaneously calibrate / flag
  - More dynamic use of threads and memory per script per intent (based on benchmarking)
    - Partitioning (IO), flagging (RAM), imaging (CPU)
  - Will see a significant speedup, necessary for arrival of 32k data
- Comprehensive data quality assessment

Inter-University Institute
for Data Intensive Astronomy

Data Quality Assessment

UNIVERSITY OF CAPE TOWN
IYUNIVESITHI YASEKAPA • UNIVERSITEIT VAN KAAPSTAD

WESTERN SYDNEY
UNIVERSITY

Jordan Collier | 20 Feb 2019 | MIGHTEE workshop

Data Quality Assessment

- Already implemented: quality assessment of data products out of processing pipeline
  - i.e. scientific DQA at end of workflow
- End-to-end analysis of MeerKAT (continuum) image / catalogue
- Presents validation report
- Produces several tables summarising data and DQA metrics
- Runs in jupyter notebook on IDIA cloud with interactive plots

MeerKAT Continuum Data Validation Report

Image

File: 'DEEP_2_mfs.sc7.image.tt0.fits'

Date	Field Centre	Central Frequency (MHz)	Synthesised Beam (arcsec)	Median r.m.s. (uJy)	Image peak (Jy)	Dynamic Range (image peak / worst r.m.s.)	Sky Area (deg^2)
2017-04-04T07:48:03.798989	04:13:26.8223 -69:00:01.1	1273.88	4.4 x 4.3	14	0.03	5E+02	6.27

Catalogue

File: 'DEEP_2_mfs.sc7.image.tt0_aegean_comp.fits'

Source Finder	Flux Type	Number of sources (≥5.0σ)	Multi-component islands	Sum of image flux vs. sum of catalogue flux	Median spectral index	Source Counts X_{red}^2
aegean	integrated	2769	113	1.1 Jy vs. 1.4 Jy		191.10

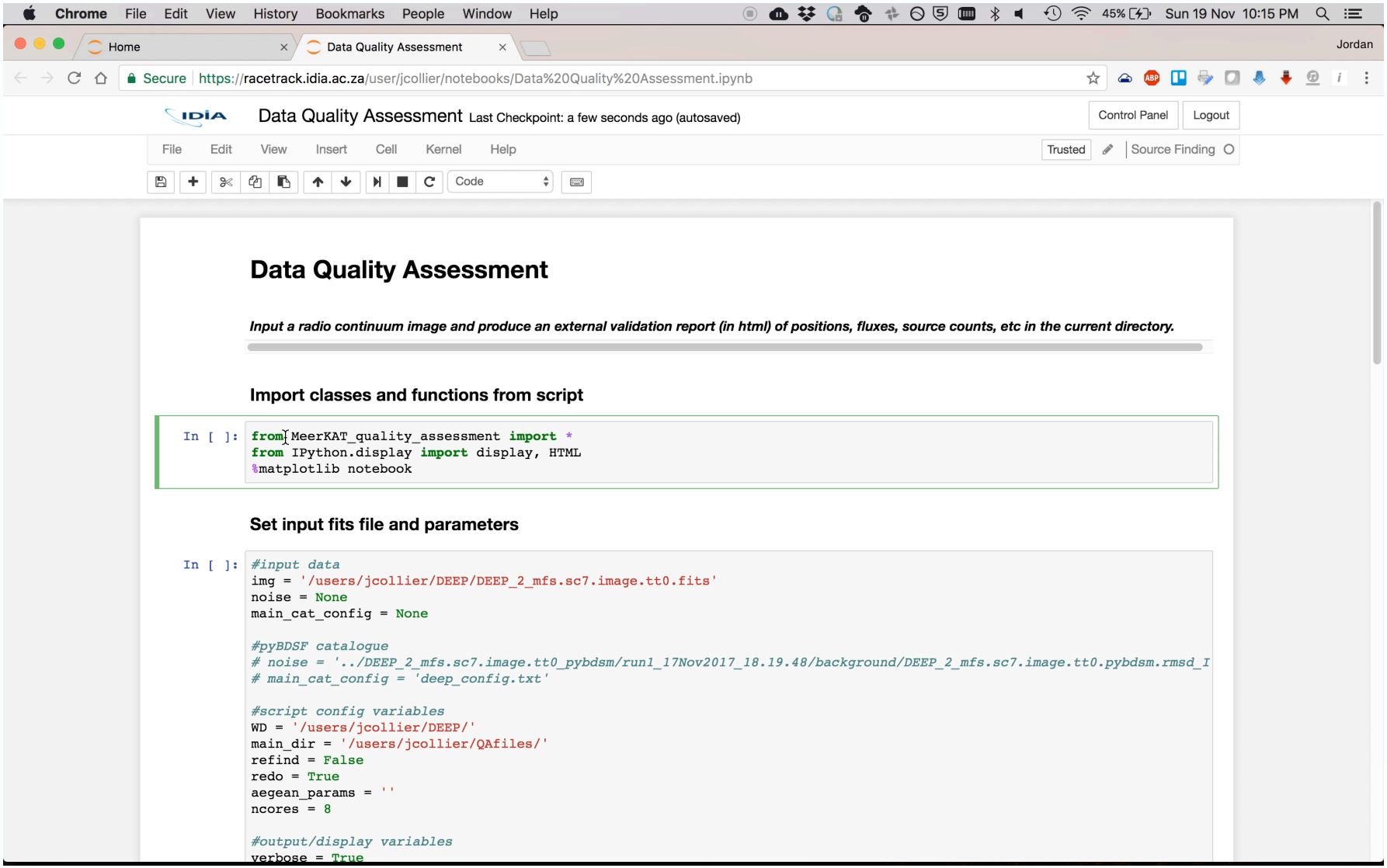
Cross-matches

Survey	Frequency (MHz)	Cross-matches	Median offset (arcsec)	Median flux ratio	Median spectral index
SUMSS	843.0	35	-0.63 ± 2.19 (RA) 0.03 ± 2.85 (Dec)	0.15 ± 0.14 (extrapolated)	-5.40 ± 2.46
GLEAM	201.0	14	-5.72 ± 4.42 (RA) 6.78 ± 10.30 (Dec)	0.04 ± 0.03 (extrapolated)	-2.59 ± 0.52

MeerKAT continuum validation metrics

Flux Ratio (MeerKAT / SUMSS-extrapolated)	Flux Ratio Scatter (MeerKAT / SUMSS-extrapolated)	Positional Offset (arcsec) (MeerKAT — SUMSS)	Positional Offset Scatter (arcsec) (MeerKAT — SUMSS)	Resolved Fraction from int/peak Flux (MeerKAT)	Spectral Index (MeerKAT-SUMSS)	Source Counts X_{red}^2 (MeerKAT)
0.15	0.14	0.63	3.60	0.21	-5.40	191.10

Data Quality Assessment



Data Quality Assessment

- Have started looking at DQA of early pipeline data products
- In close collaboration with MeerKAT SDP.
  - Framework to measure quality of pipelines.
  - Standard set of metrics between all pipelines
    - Defer processing to most efficient pipelines?
  - Mapping science requirements from LSPs to technical requirements for pipelines

Data Quality Assessment

Table 1: Selected requirements and specifications from various MeerKAT imaging LSPs

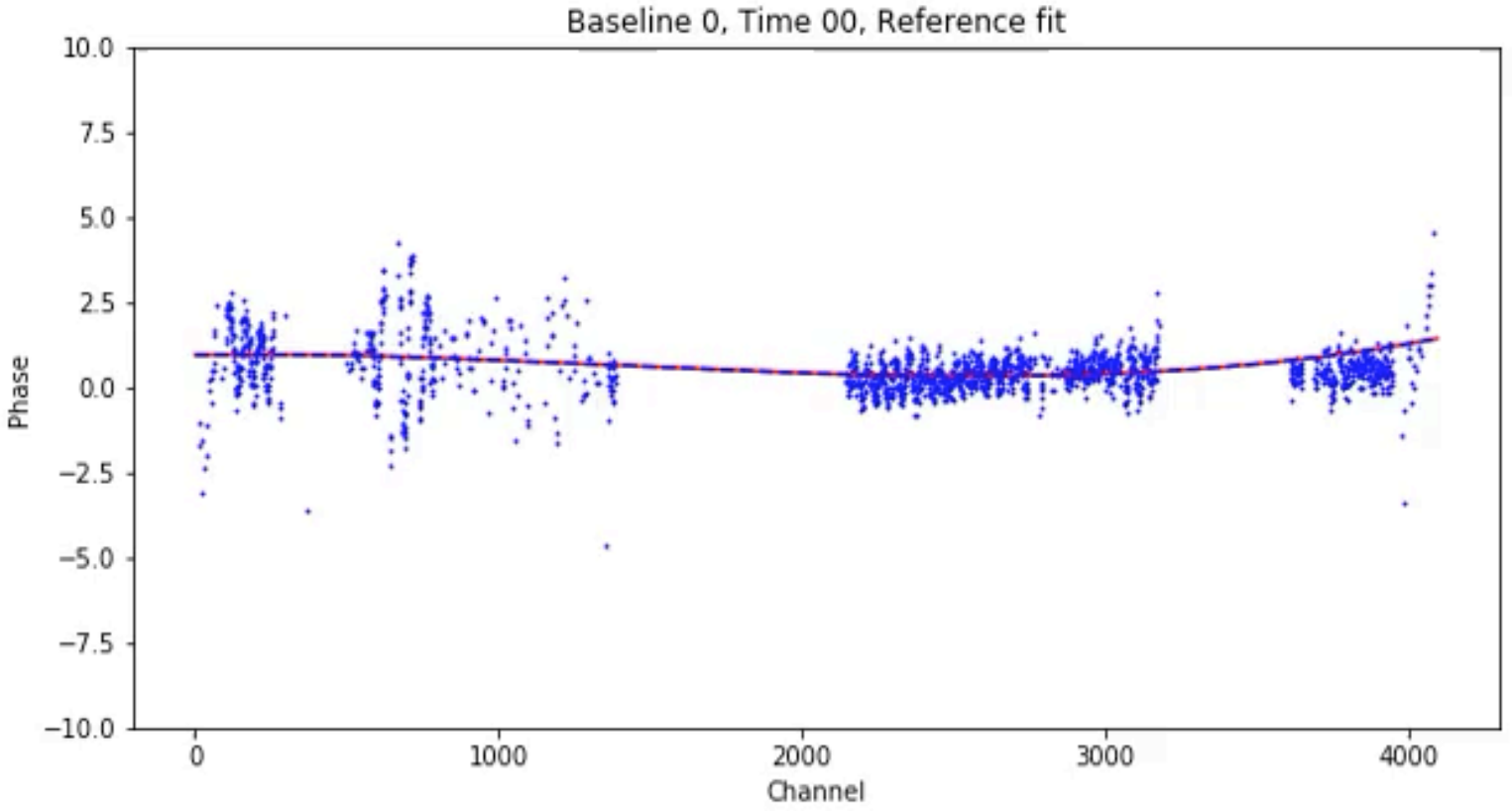
LSP	Sensitivity (μJy beam^{-1})	Dynamic Range	Velocity Resolution (km s^{-1})	Redshift Range	Area (deg^2)	RM precision (rad m^{-2})	Column Density Sensitivity	H I Mass ($M_\odot$)
MIGHTEE (L-band)	2^a / 90^b	$\geq 10^{5c}$	6^d	$z \lesssim 0.5$	20	$\sim$1	$\sim 1\ M_\odot\ pc^{-2^e}$	$\gtrsim 2 \times 10^{9f}$
MIGHTEE (S-band)	1^g	$\geq 10^{5c}$	-	$z \lesssim 0.5$	5.5	$\sim$1	-	-
MIGHTEE (UHF-band)	6^a	$\geq 10^{5c}$	-	$z \lesssim 0.5$	3.5	$\sim$1	-	-
LADUMA (L-band)	45^b	-	6^d	$0 \leq z \leq 0.58$	0.9–2.2	-	-	$\sim 10^{7.5-10.5}$
LADUMA (UHF-band)	26^h	-	8^i	$0.42 \leq z \leq 1.45$	1.8–5.4	-	-	$\sim 10^{9.2-10.5}$
FORNAX	100^j	-	$\sim$1	$z \sim 0$	$\sim$12	-	$\sim 0.1^k - 5^l \times 10^{19}$cm^{-2}	$\gtrsim 5 \times 10^{5l}$
MHONGOOSE	74^j	-	16	$z \sim 0$	$>45^m$	$\lesssim 1$	$0.55^n - 7.5^o \times 10^{18}$cm^{-2}	$\sim 10^{6-11}$
MALS (L-band)	3 / 500^p	-	$\sim$5	$0 < z < 0.85^q$	1000^r	-	$> 10^{19}$cm^{-2}	$\sim 5 \times 10^4$
MALS (UHF-band)	3 / 600^p	-	$\sim$5	$0.4 < z < 1.87^q$	700^r	-	$> 10^{19}$cm^{-2}	$\sim 5 \times 10^4$
THUNDERKAT	$\sim$1–1000^s	-	-	-	-	-	-	-

Data Quality Assessment

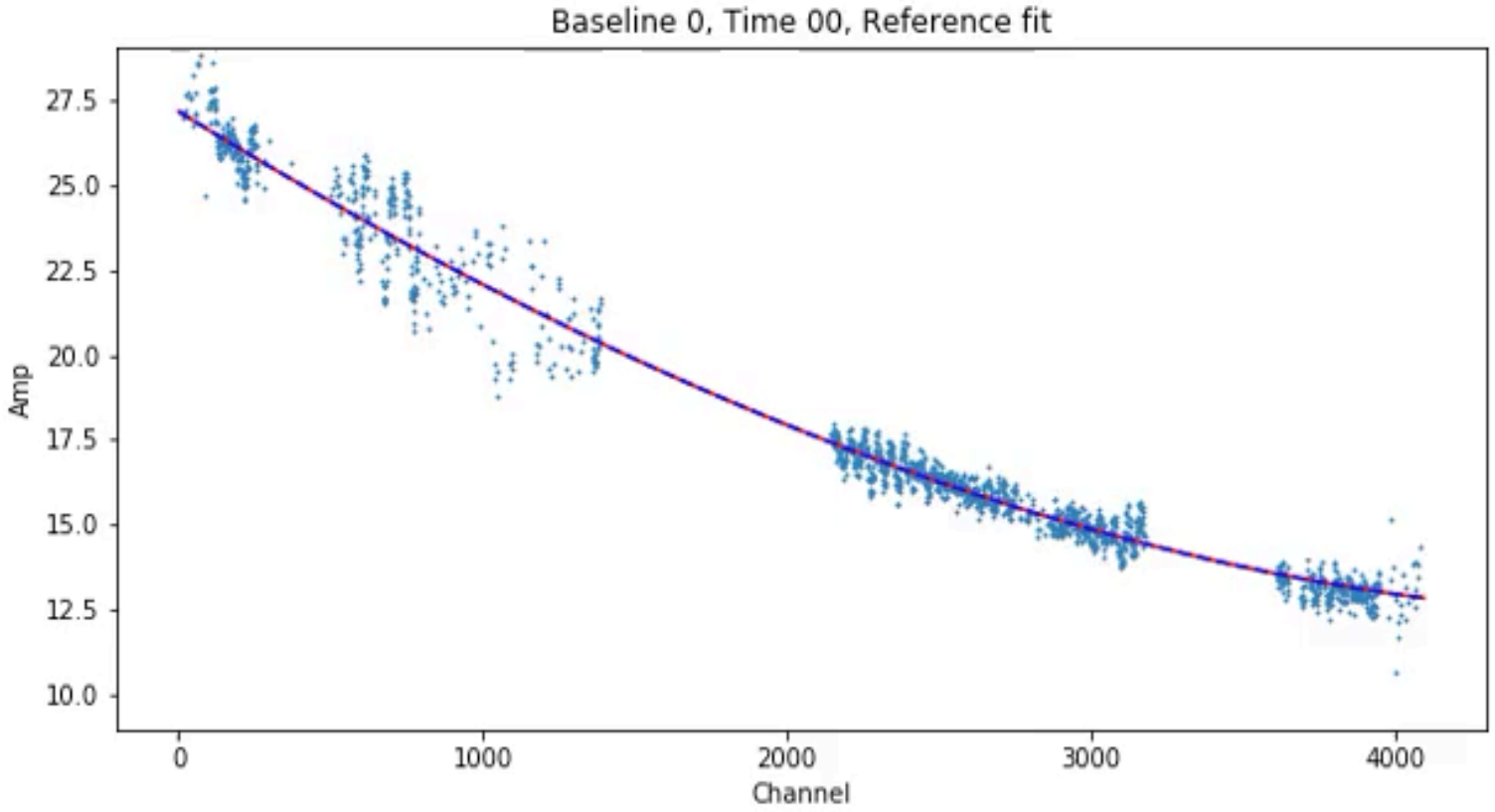
Table 2: Data quality metrics for a simple calibration pipeline. This table is not exhaustive, but represents a selection of metrics we have drafted.

Step	Distribution / plot	Statistic / Metric	Computation	Tolerance
Bandpass calibration	Calibrated bandpass (amplitude and phase) as a function of frequency (per timestamp)	Residual of polynomial fit	Mean reduced χ squared of polynomial fit to each timestamp, compared to reference timestamp[a]	< 10
	Calibrated bandpass (amplitude and phase) as a function of frequency	Normalised median absolute deviation	Maximum normalised absolute deviation from polynomial fit	< 5
Bandpass flagging	Calibrated amplitude as a function of frequency	Fraction of channels flagged	Number of channels with >50% of visibilities flagged divided by total number of channels	$< 10\%$
Phase calibration	Complex gain solutions as a function of time	Outlier metric	Running median?	?
	Calibrated amplitude as a function of frequency	Normalised median absolute deviation	Normalised median absolute deviation from polynomial fit	< 5
Phase cal. flagging	Calibrated amplitude as a function of time	Fraction of timestamps flagged	Number of timestamps with >50% of visibilities flagged divided by total number of timestamps	$< 10\%$
Target flagging	Amplitude as a function of frequency	Fraction of data flagged	Fraction of visibilities flagged divided by total visibilities	$< 20\%$
	Amplitude as a function of frequency	Normalised median absolute deviation	Normalised median absolute deviation	< 5

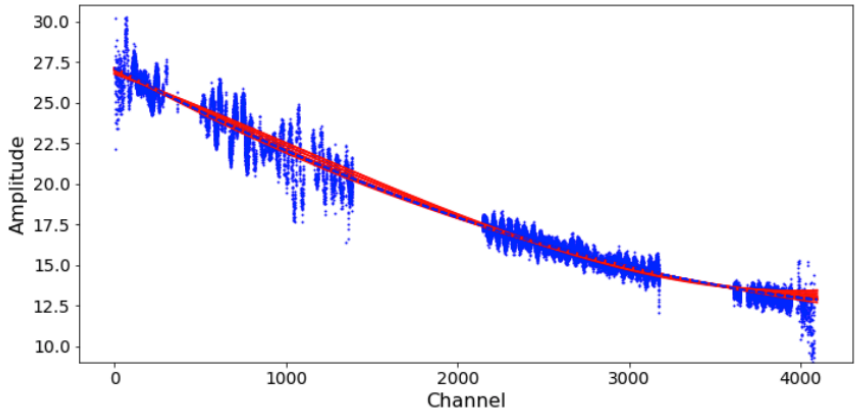
Data Quality Assessment

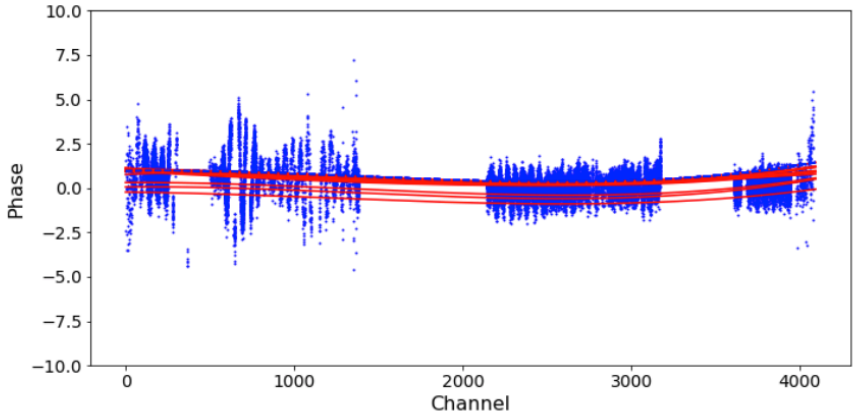


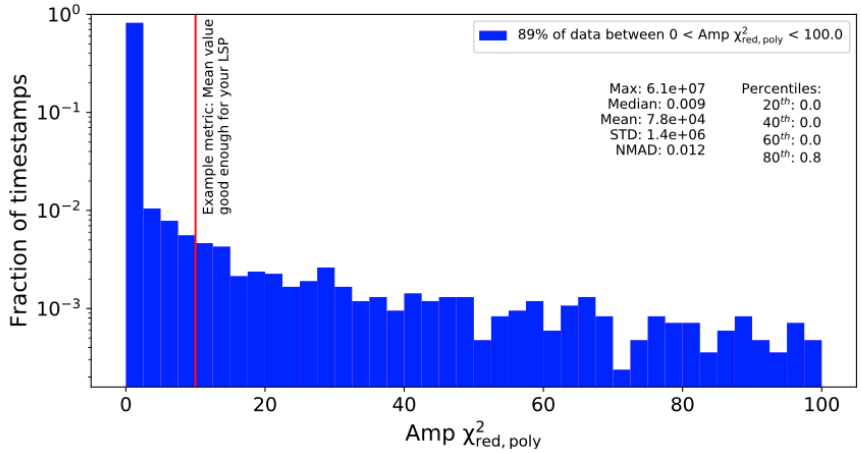
Data Quality Assessment

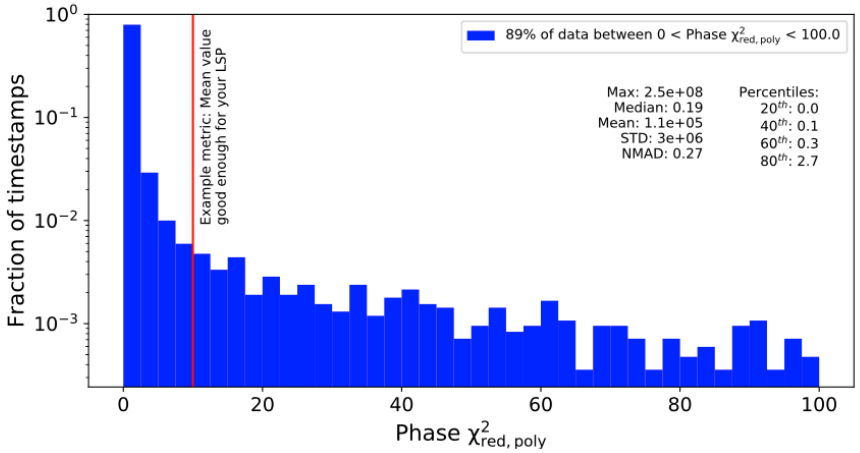


Data Quality Assessment









Data Quality Assessment

- DQA being discussed amongst SKA pathfinders in general, within SPARCS DQA WG
  - http://spacs.pbworks.com/w/page/126067640/dataquality
- Drafted list of metrics between us
  - https://docs.google.com/spreadsheets/d/1l8x0doPW6LhoWZMLeB5W4tcwzdgw18FWdh5DxnK6_vs/edit#gid=0

Summary

- The IDIA “processMeerKAT” pipeline is an efficient, user-friendly pipeline, that is widely tested and documented
- It runs on the Ilifu cluster, making dynamic use of resources, and containers, and presents a good framework for pipelines
- Many use cases are supported, incl. inserting your own scripts
- Even the quick look images are good
- Coming soon: selfcal, AW projection, optimisation/speedup

THANK YOU

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