

Application of Pipeline Calibration

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What are we talking about?

- What to do after a Pipeline Processing Complete email.
- Extra flagging, re-derive & apply calibration.
- Rerun for known issues that cause problems.
- Apply existing pipeline calibration to raw data.
- Things to consider when running on your own.
- Remote access to NRAO computing.

Extra flagging

The pipeline may not flag everything needed ...

Antenna hardware issues – RFI – may flag good data – ~~Alien signals*~~

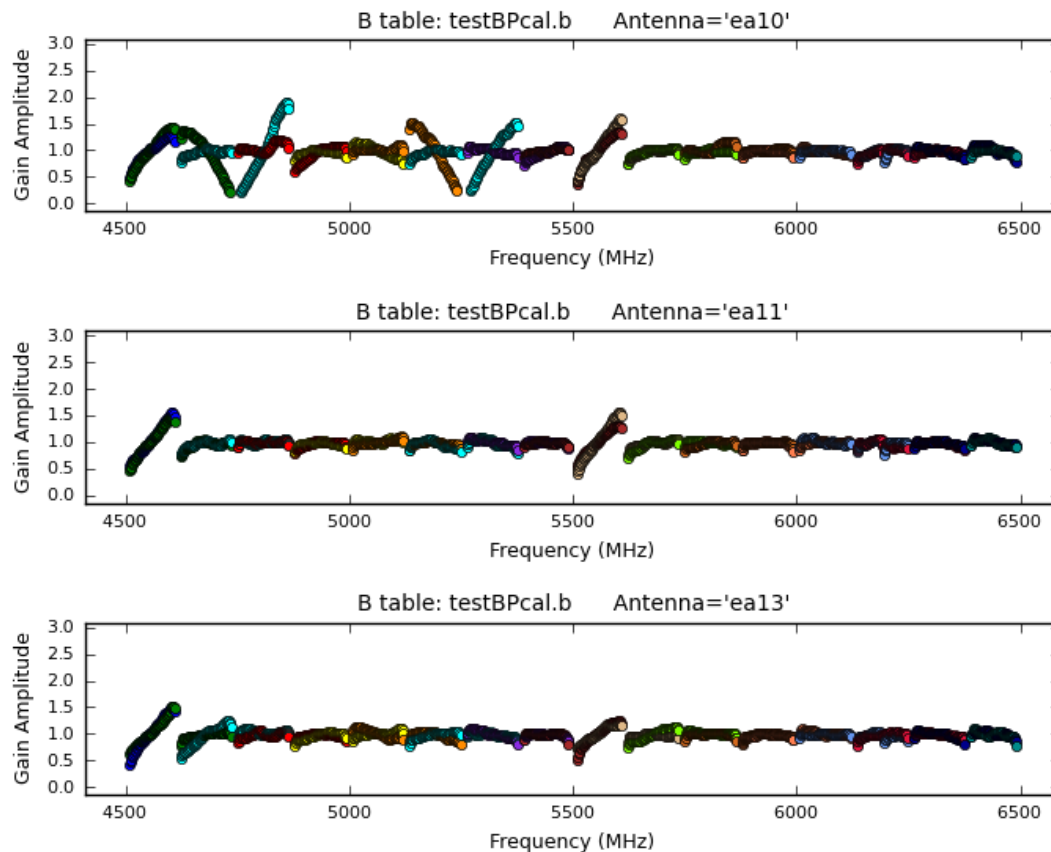


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*Theoretical only at this time

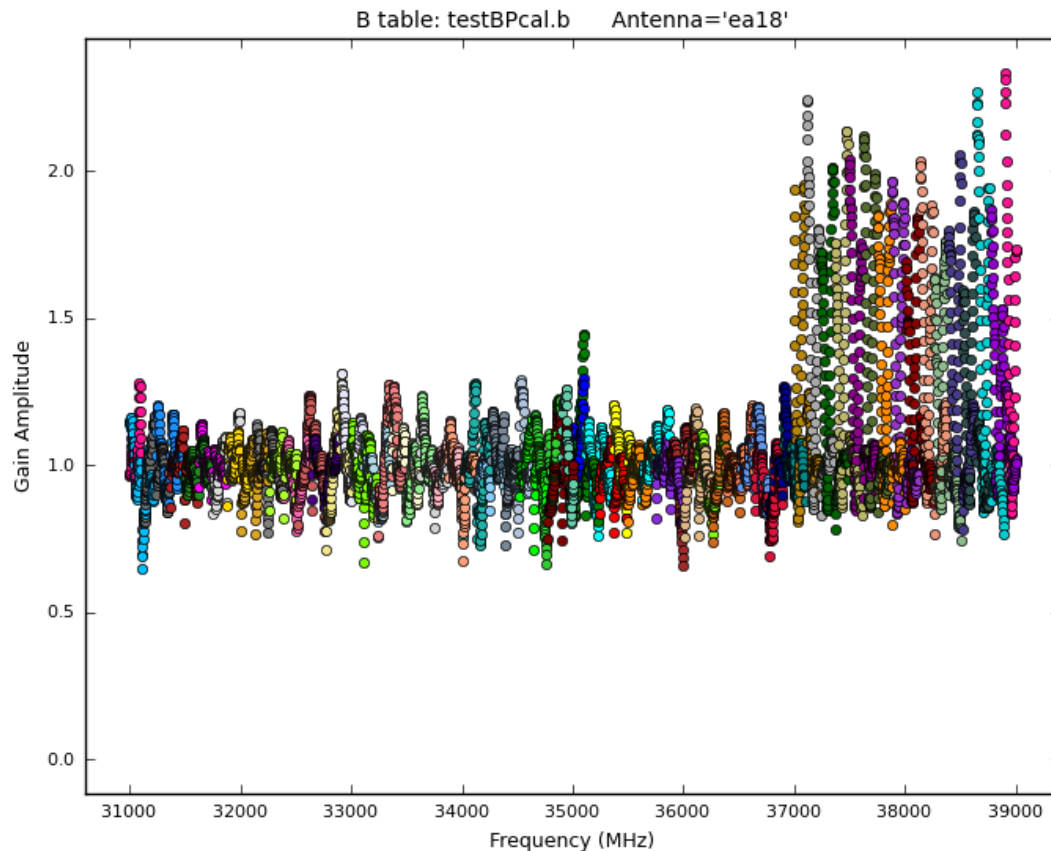
Additional flagging

ea10 bandpass, bad data (DTS issue); ea11, ea12 OK



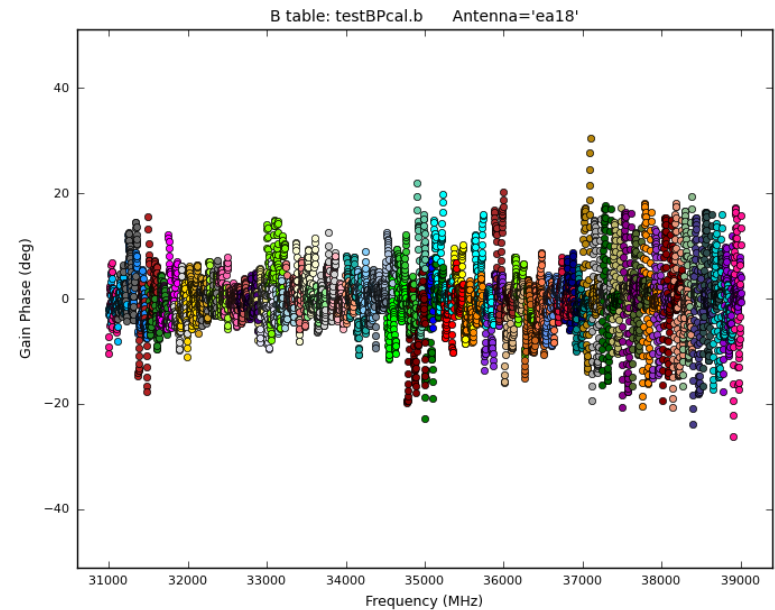
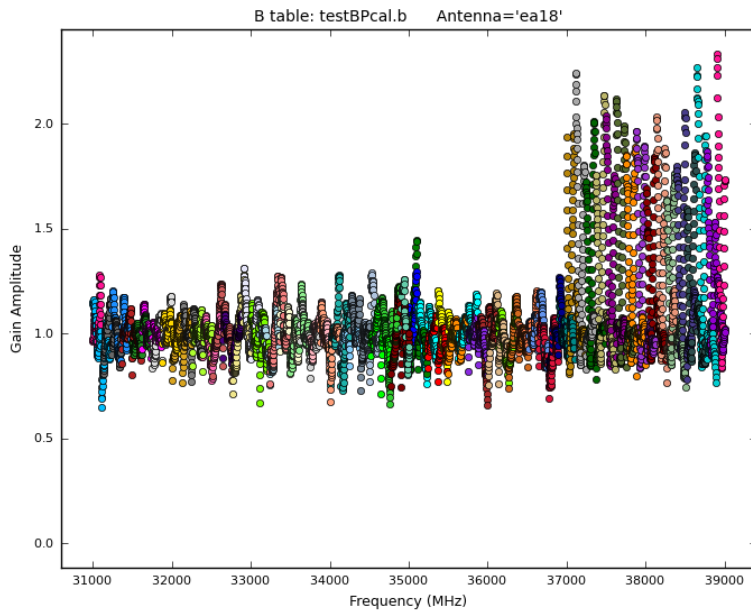
Additional flagging

ea18 bandpass, bad data (DTS issue for 37-39GHz)



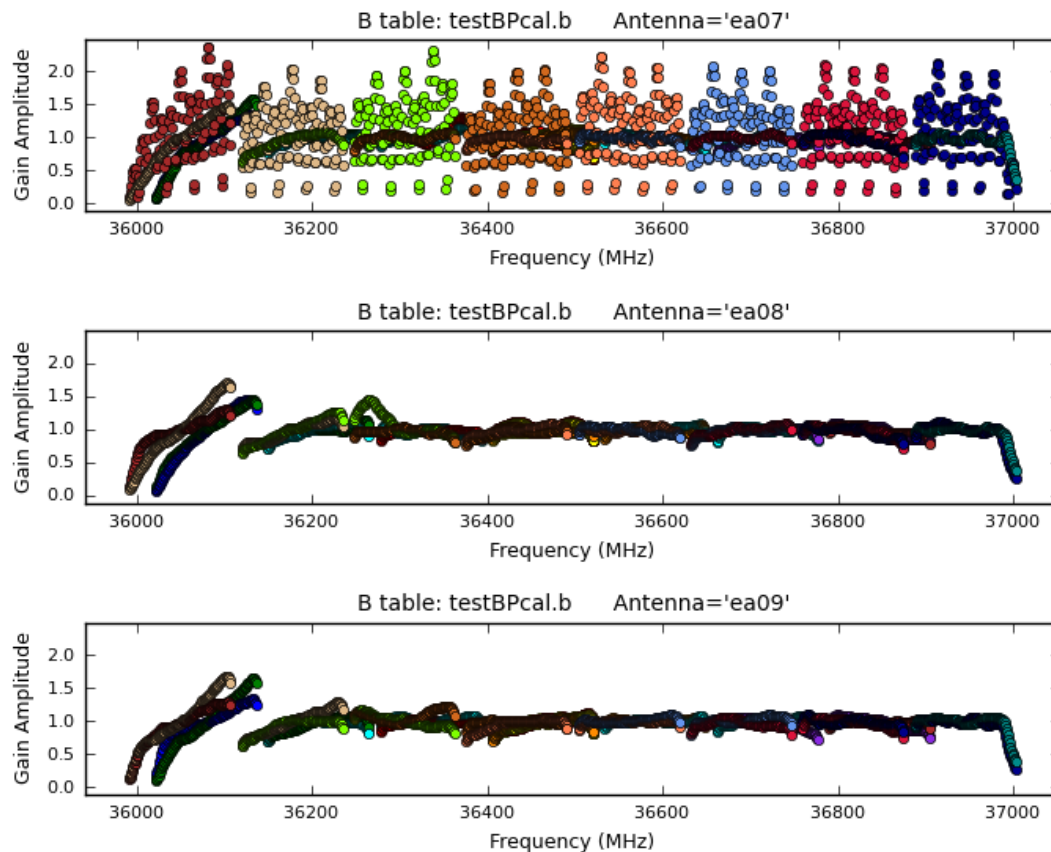
Additional flagging

ea18 bandpass **and** phase affected, bad data (DTS issue)



Additional flagging

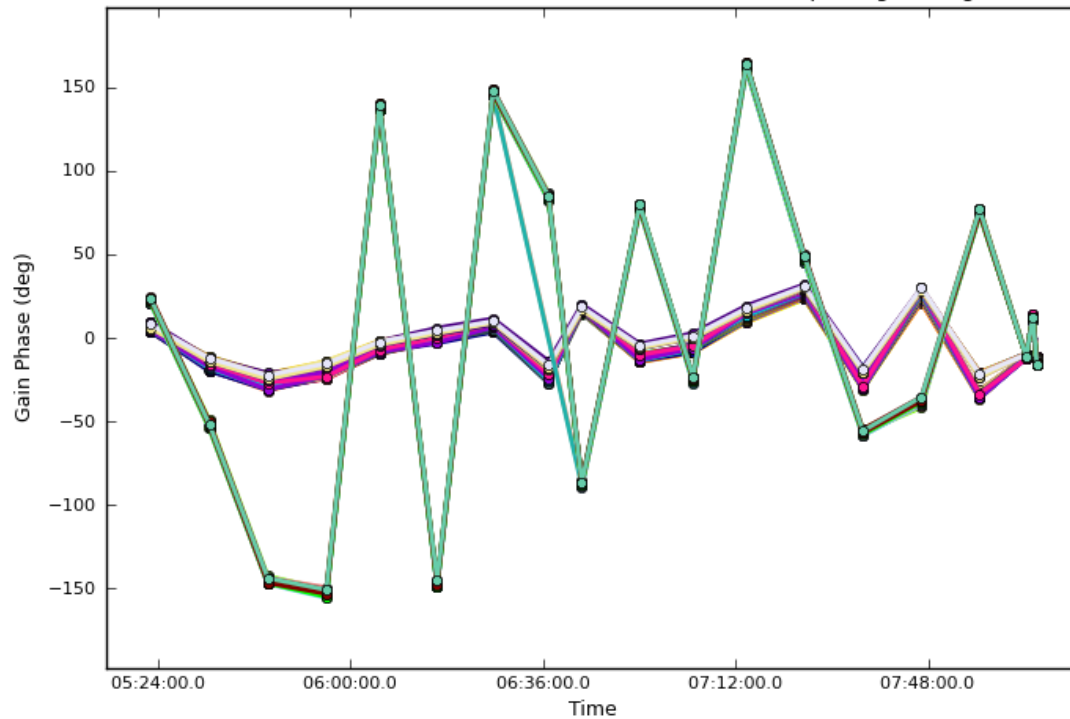
ea07 bandpass, bad data (DTS issue); ea08, ea09 OK



Additional flagging

ea02 phase jumps for some spws

G table: 13A-398.sb17165245.eb19476558.56374.213876608796.ms.finalphasegaincal.g Antenna='ea02'



Additional flagging

- Carefully check your data and the calibration from the pipeline.
- If extra flagging **ONLY** on your science target(s), no recalibration of your data required: use CASA as usual.
- If any of your **calibrator sources** require additional flagging, you should re-derive the calibration with your additional flagging included.
- To recalibrate your data using the VLA Calibration Pipeline, there are two good starting points:
 - Pipeline calibrated MS
 - Raw data (SDM-BDF) >> (see Scripted Pipeline webpage)

Additional flagging: Cal'd MS

1. Create a pipeline recipe **without** Hanning smoothing:
 - Navigate to your CASA installation with the pipeline.
 - Go to /pipeline/pipeline/recipes inside your CASA installation.
 - Copy hifv.py and **rename** to hifv_nohanning.py.
 - Comment out call to hifv_hanning (# symbol)
 - Save changes.
2. Inspect the calibrated MS and flag as needed in CASA.
3. Create a **new directory** and copy **ONLY** the calibrated and fully flagged MS to this new directory. **No** other files should be copied.

Additional flagging: Cal'd MS

4. From your new directory with the flagged MS, start CASA:

`casa --pipeline` (be sure to start the same version where you were editing a new recipe!)

5. Clear the *calibration* using **clearcal** with `addmodel=False`.
*See the pipeline web page (special topics) for details.

6. Next, run the **clearstat** task in CASA.

5. `import pipeline.recipes.hifv_nohanning as hifv_nohanning`

6. `hifv_nohanning.hifv(['MSname'])`

7. Wait again while the pipeline runs.

Rerun for known issues (scripted pipeline)

The pipeline may choose the worst possible option ...

- Problems during the flux or delay calibration scan(s)
- Bad reference antenna or setup issues
- Run the Scripted Pipeline (see Scripted Pipeline webpage)



Rerun for known issues (scripted pipeline)

1. From the SMD-BDF, create a MS and apply online flags. You may request the online-flags-applied MS from the NRAO archive.
2. Carefully inspect your data to determine what additional flagging is required.
3. Using the **flagdata** task in CASA, you will create a text file (e.g. additional_flags.txt) for all additional flags needed (no need to apply these flags now!) We recommend:

```
flagdata(action='none',mode='manual',savepars=True,outfile='additional_flags.txt', antenna='ea##,ea##', scan='scan#,scan#')
```

4. Rerun the above until all additional flags have been added. Through this approach, all the flags will be accumulated in a single text file that may be applied on the data through one application once all extra flags have been recorded.

Rerun for known issues (scripted pipeline)

5. Make a copy of the file "EVLA_pipeline.py" and rename it.
6. Edit your renamed copy of EVLA_pipeline.py to include an extra step of the CASA task **flagdata** just after 'flagall.py' is called.

```
flagdata(vis=ms_active,mode='list', inpfiler='additional_flags.txt',  
action='apply', flagbackup=True)
```

Please take care to **notice indentation** amounts and to set your CASA flagdata command to the same indentation as the other "execfile()" commands.

7. In a new directory with the SDM-BDF, start the pipeline by calling your edited version of EVLA_pipeline.py

```
execfile('/path/on/computer/EVLE_pipeline_edit.py')
```

8. Wait ... wait longer ... just come back the next day.

Apply to Raw Data

May only have pipeline calibration & flag tables, no MS

- Calibrated MS held by NRAO for only 15 days:
Calibration tables, flag tables, weblog archived!
- Local storage limitations:
Reduced storage needs by only keeping the tables.



Apply to Raw Data

1. Download the correct CASA version with the pipeline.
2. You will need the following
 - SDM-BDF
 - Session_1.caltables.tar.gz
 - *.ms.flagversions.tar.gz
 - *.ms.calapply.txt
 - casa_piperestorescript.py
3. Make a directory called “restoration”.
4. cd to restoration, and create three more directories inside:
rawdata, working, & products << names must be exact!

Apply to Raw Data

5. Put your SDM-BDF into the “rawdata” directory.
6. Put all the *.tar.gz files and *.txt files into the “products” directory.
7. Put casa_piperestorescript.py into the “working” directory.
8. Go to the “working” directory and edit casa_piperestorescript.py:
 - Insert “../rawdata/” before the SDM-BDF name (mySDM) in the call to hifv_restoreddata.
 - Save your changes.

Apply to Raw Data

9. From the “working” directory, start CASA with the pipeline

```
casa --pipeline
```

10. Execute the casa_piperestorescript.py file:

```
execfile('casa_piperestorescript.py')
```

11. Enjoy calibrated data once the process completes.

Considerations

Scan intents correct?

Hanning Smoothing?

Computing time?

Disk space – 3-4X raw size(!)

PL version differences.

CASA version differences.

NRAO cluster available for
remote Access



Remote Access: Accounts

Use your **visitor account** (what you're using now)

- Remote processing
- Data staging for download
- Short term work, **NOT** long term storage.

Use your account's "**data**" directory:

- Archive deliveries (future archive will use this)
- Pipeline data requests
- Don't change permissions of this directory!

Remote Access: Node Request

Login with your account username

- `ssh nm-####@login.aoc.nrao.edu`

Go to `nmport-master` and request a node

- `ssh nm-####@nmport-master`
- `nodescheduler --request 14 1`

If you get no email, you are probably queued

- Don't keep requesting more nodes

Exit `nmport-master`, then ssh to your assigned node

- `ssh nm-####@nmport###`

Remote Access

Interact with your data for reduction and analysis

- SSH and VNC available for working with your data.

Download your data:

- RSYNC, SFTP, SCP, LFTP available.

Too much data to download?

- Use hard disk shipping option.

Need help?

- <https://science.nrao.edu/facilities/vla/docs/manuals/computing-resources>

Questions?

- VLA CASA Calibration Pipeline information at:

<https://science.nrao.edu/facilities/vla/data-processing/pipeline>

- CASA Integrated Pipeline & Scripted Pipeline available
- Have Questions?
- Need Help?
- Use the **NRAO HelpDesk**: **<https://help.nrao.edu/>**
- Submit your ticket under the **Pipeline Department**.
- Please include specific details when submitting HelpDesk tickets.
(Project code, SB number, CASA/PL versions, errors, etc.)