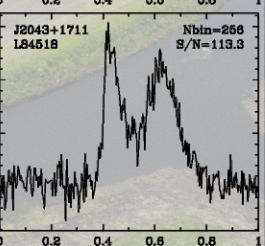
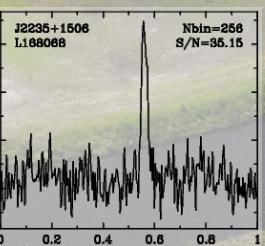
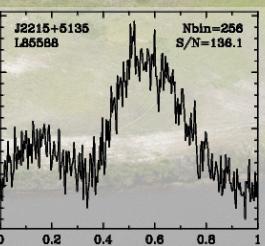
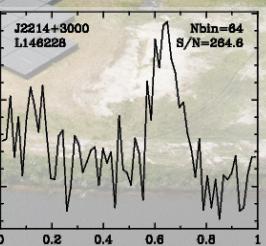
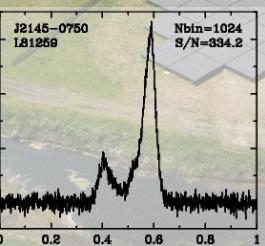
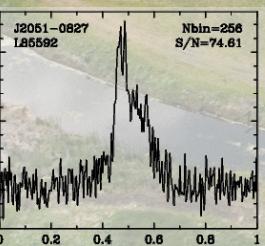
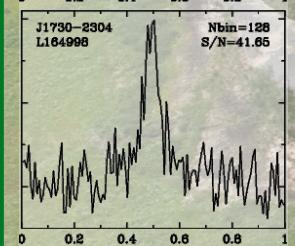
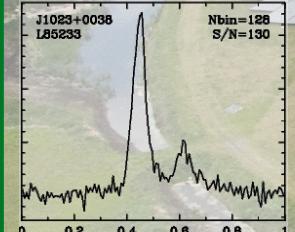
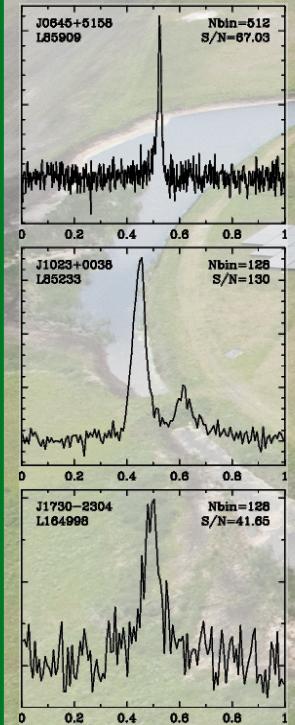


LOFAR Pulsar Pipeline (PulP)

Vlad Kondratiev
(ASTRON)



Pulsar visualisation credit: Alessandro Ridolfi

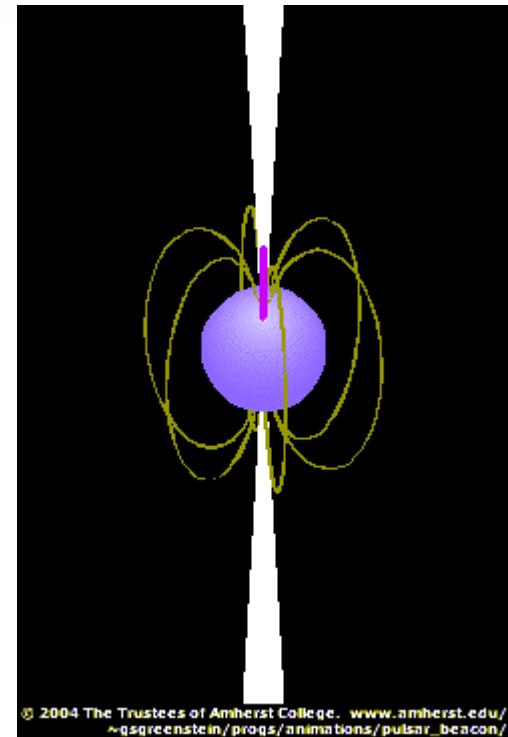
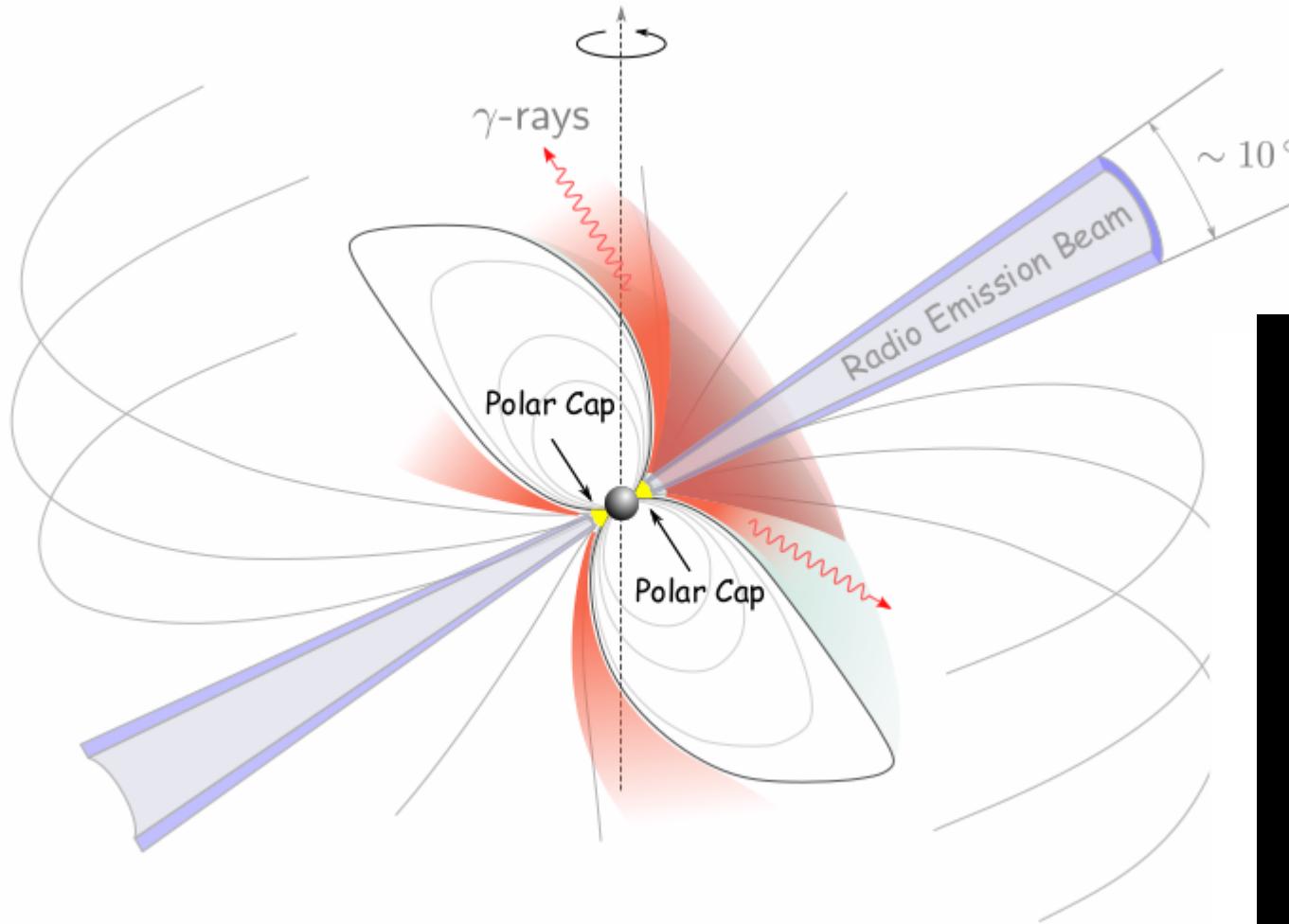
What is PulP?

PulP is LOFAR Pulsar Pipeline for *known pulsars*. The essential goal of the PulP is to get the average profile of the pulsar(s) and provide a user with freq/time/phase/pol data cubes for further analysis. It is *not* the *search* pipeline, i.e. you can not do periodicity and single-pulse searches for a large range of dispersion measure trials. However, PulP can provide both PSRFITS/filterbank data and raw data converted to 8-bit for further searches.

Disclaimer: In no way PulP can guarantee the optimal use of all the processing parameters for *every* observation/pulsar. The default parameters are only meant to provide the good results for most observations based on our current knowledge in the automated way. And serves to give the first diagnostics of the quality of the data.

What is pulsar?

rapidly rotating
highly-magnetised neutron star,
«electric lighthouse»



Outline:

- Pulp overview
 - briefly: Pulp implementation
- In a nutshell about (de-)dispersion, folding
- Pulp flowchart
 - DSPSR pipeline
 - PRESTO pipeline
- Pulp output data
- Pulp in the NorthStar
- Pulp options

PulP overview

- Bookkeeping, service functions
 - Logging
 - Cluster configuration/settings
 - User options
 - Where input data are?
 - Observing setup (HDF5 metadata / *parset*)
 - Coordination of processing data for different TABs/frequency parts
 - Feedback files for LTA ingest
- The actual data processing
- Diagnostic summaries and pipeline output data products

PulP overview

- Bookkeeping, service functions
 - Logging
 - Cluster configuration/settings
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 - Where input data are?
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 - Coordination of processing data for different TABs/frequency parts
 - Feedback files for LTA ingest
- **The actual data processing**
- **Diagnostic summaries and pipeline output data products**

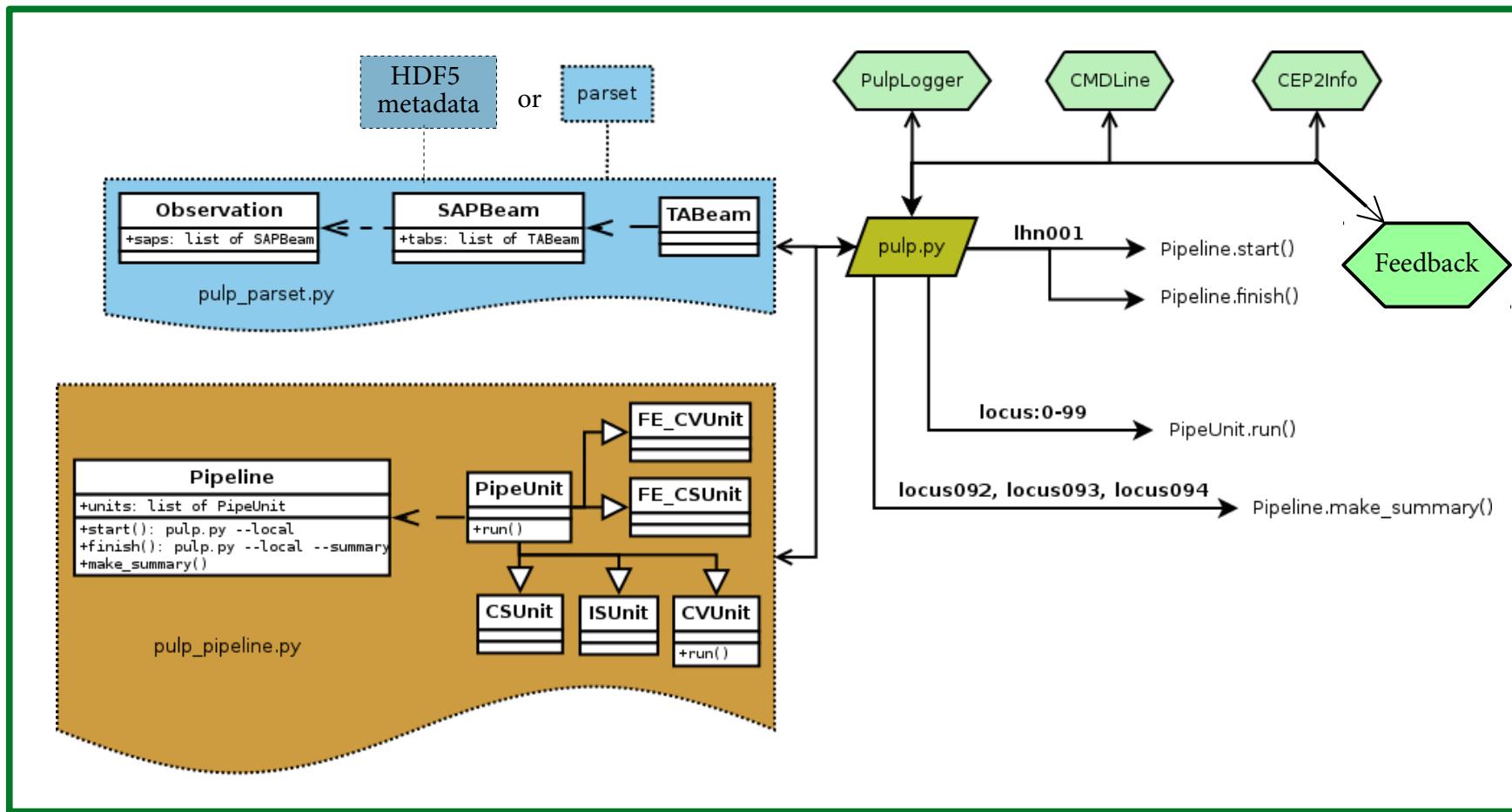
PulP overview

- Bookkeeping, service functions
 - Logging
 - Cluster configuration/settings
 - User options
 - Where input data are?
 - Observing setup (HDF5 metadata / *parset*)
 - Coordination of processing data for different TABs/frequency parts
 - Feedback files for LTA ingest

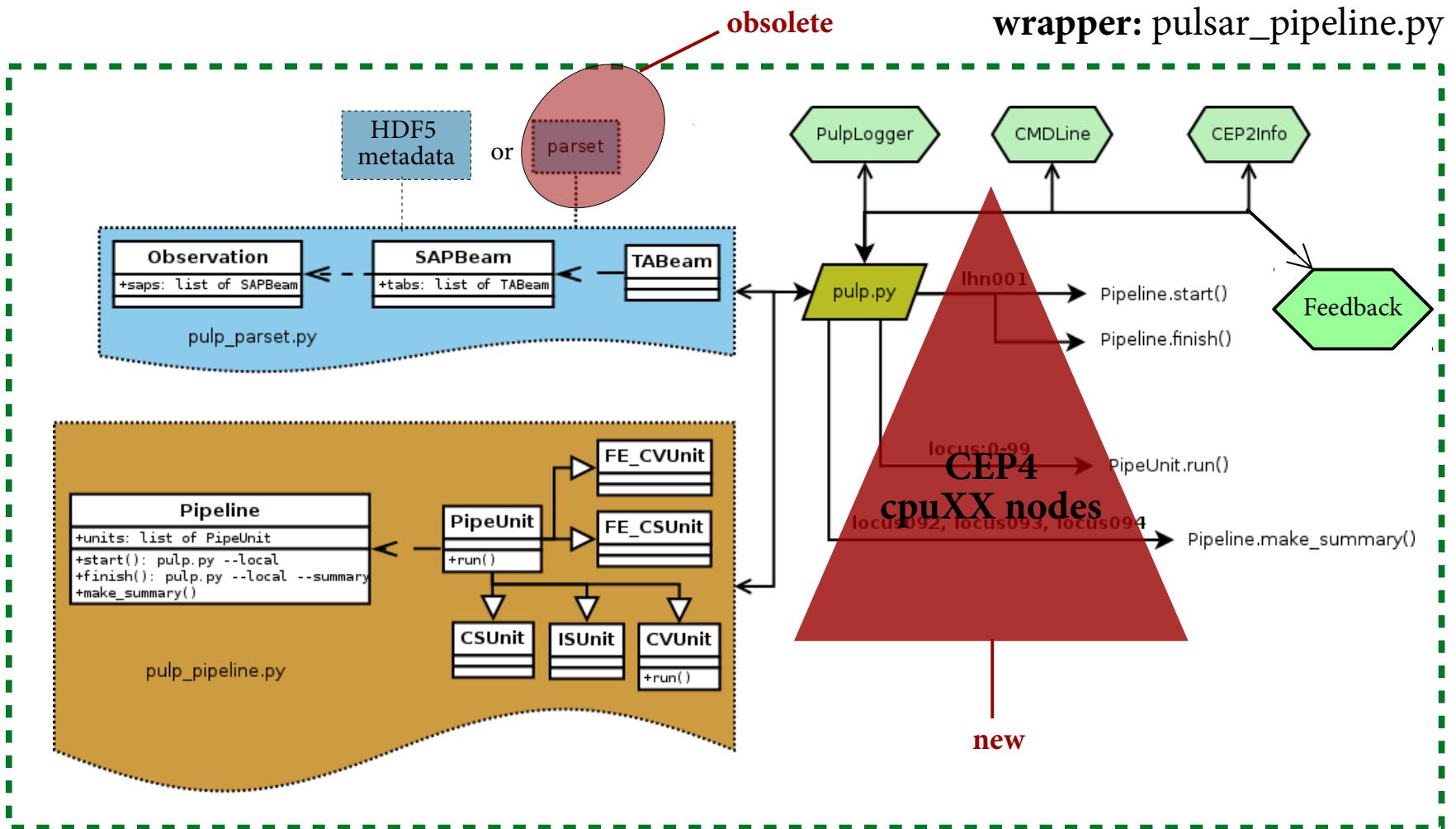
- The actual data processing
- Diagnostic summaries and pipeline output data products

NB: PulP is still implemented in the LOFAR system as a «black box» via the *wrapper* that calls the original PulP itself. This slows down bug fixing and further improvements quite significantly, and the plan is to truly incorporate PulP into the central system as it is the case for imaging pipeline(s).

PulP implementation

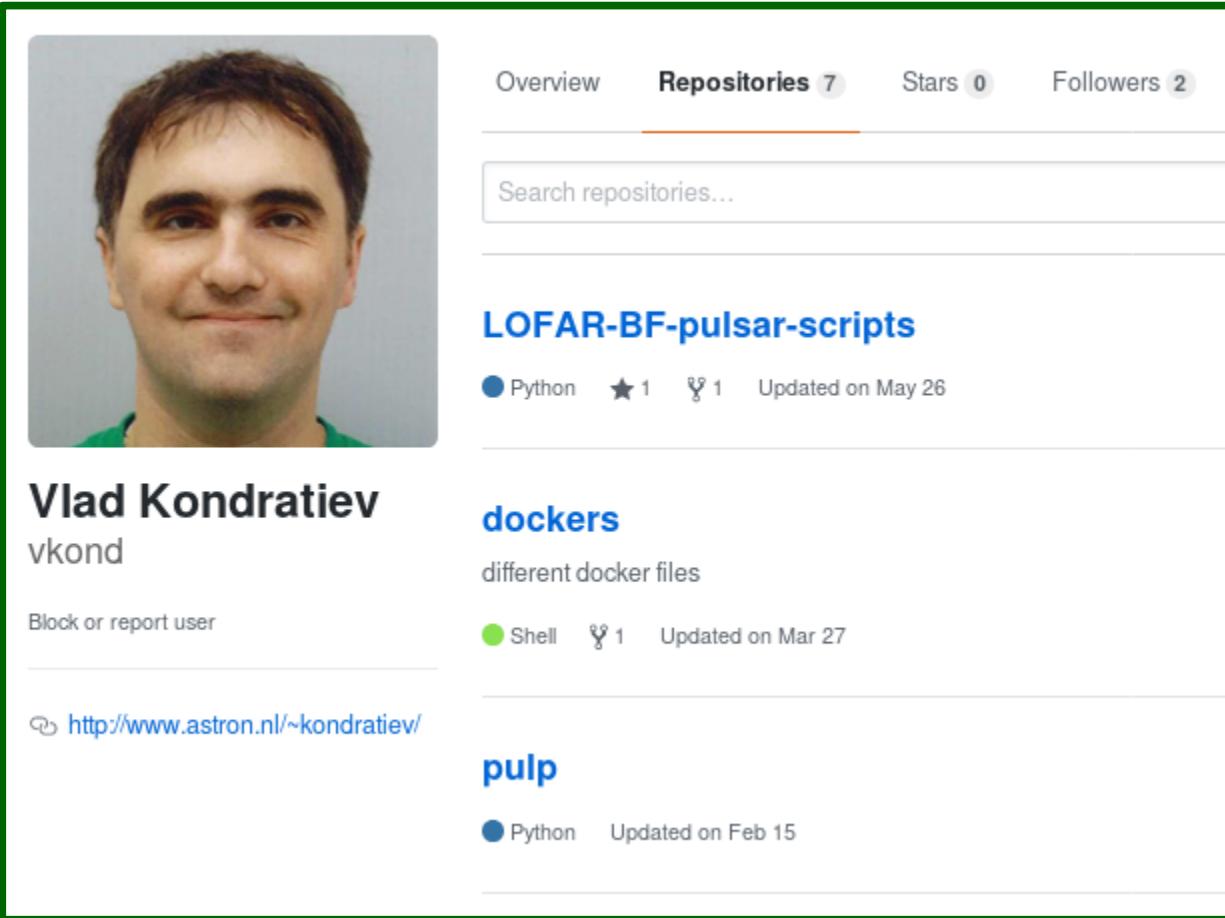


PulP implementation



@github

<https://github.com/vkond>



The screenshot shows Vlad Kondratiev's GitHub profile. At the top, there is a large photo of him, followed by his name "Vlad Kondratiev" and handle "vkond". Below this, there are links to "Block or report user" and a link to his website "http://www.astron.nl/~kondratiev/". The main section is titled "Repositories" and shows three repositories:

- LOFAR-BF-pulsar-scripts**: Python, 1 star, 1 fork, updated on May 26. Description: "different docker files".
- dockers**: Shell, 1 star, 1 fork, updated on Mar 27. Description: "different docker files".
- pulp**: Python, updated on Feb 15.

Various scripts

<https://github.com/vkond/LOFAR-BF-pulsar-scripts>

Dockerfile for LOFAR

<https://github.com/vkond/dockers>

PulP:

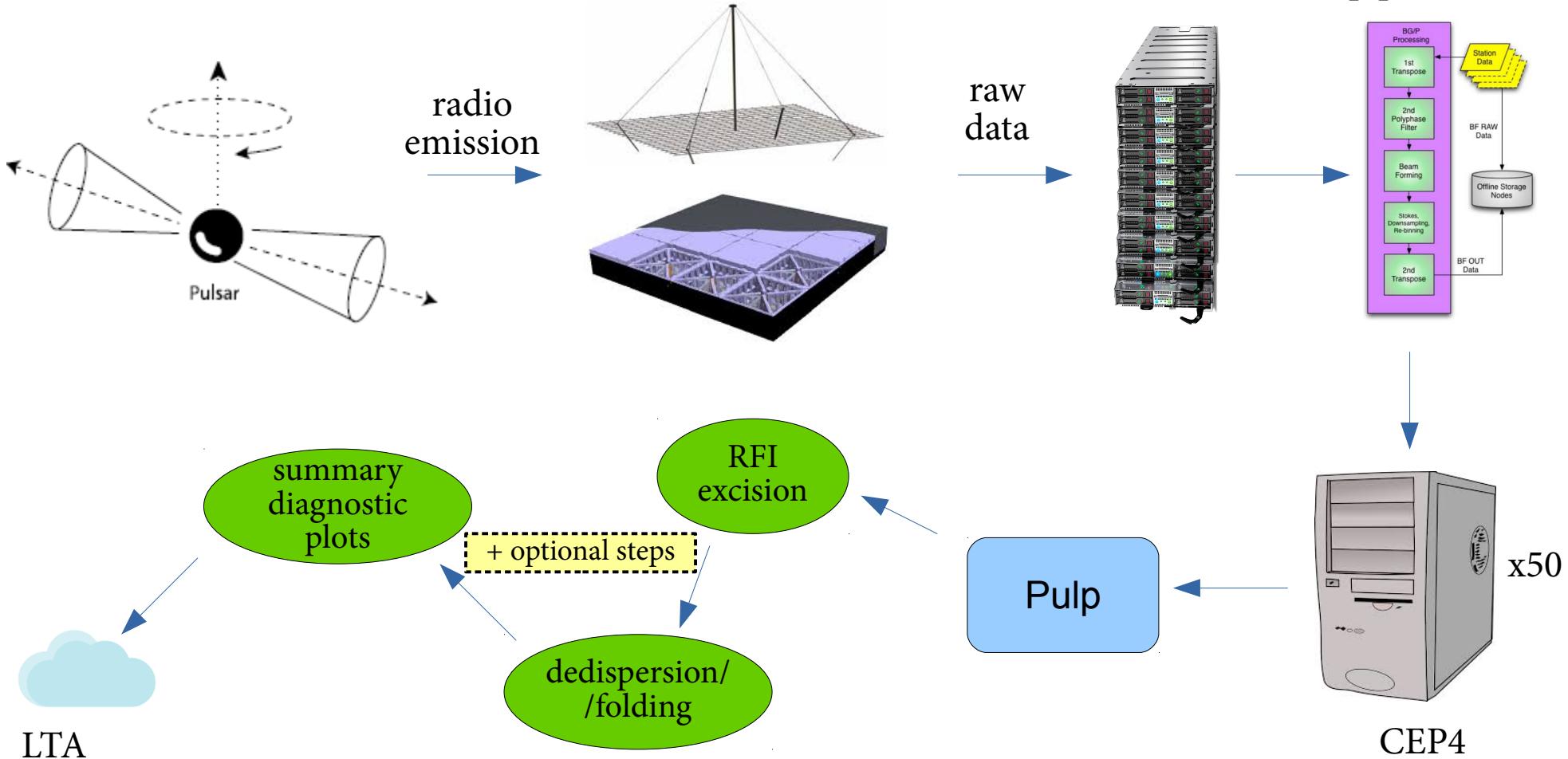
<https://github.com/vkond/pulp>

Pulsar software (needed by Pulp)

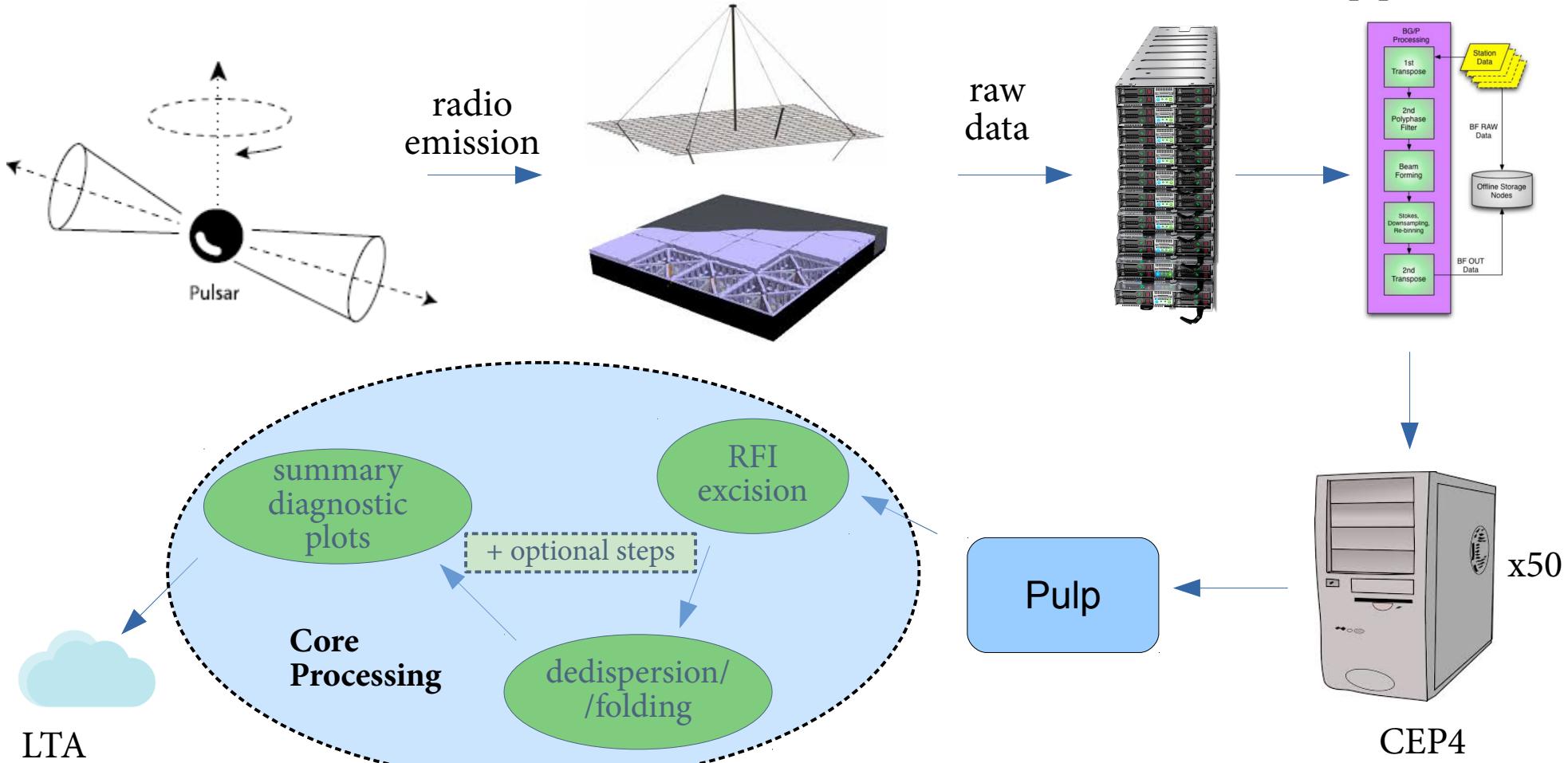
- FFTW
- PGPLT, + python bindings
- TEMPO
- TEMPO2
- psrcat
- Sigproc
- PRESTO
- psrdada
- PSRCHIVE
- DAL
- DSPSR
- COAST_GUARD (written by Patrick Lazarus, for RFI excision)
- LOFAR-BF-pulsar-scripts
- in the future (needed for pulsar flux calibration):
 - casacore
 - python-casacore
 - mscorpol

Lofar-pulp Docker container has all this software installed:
<https://github.com/vkond/dockers>

Data flow



Data flow



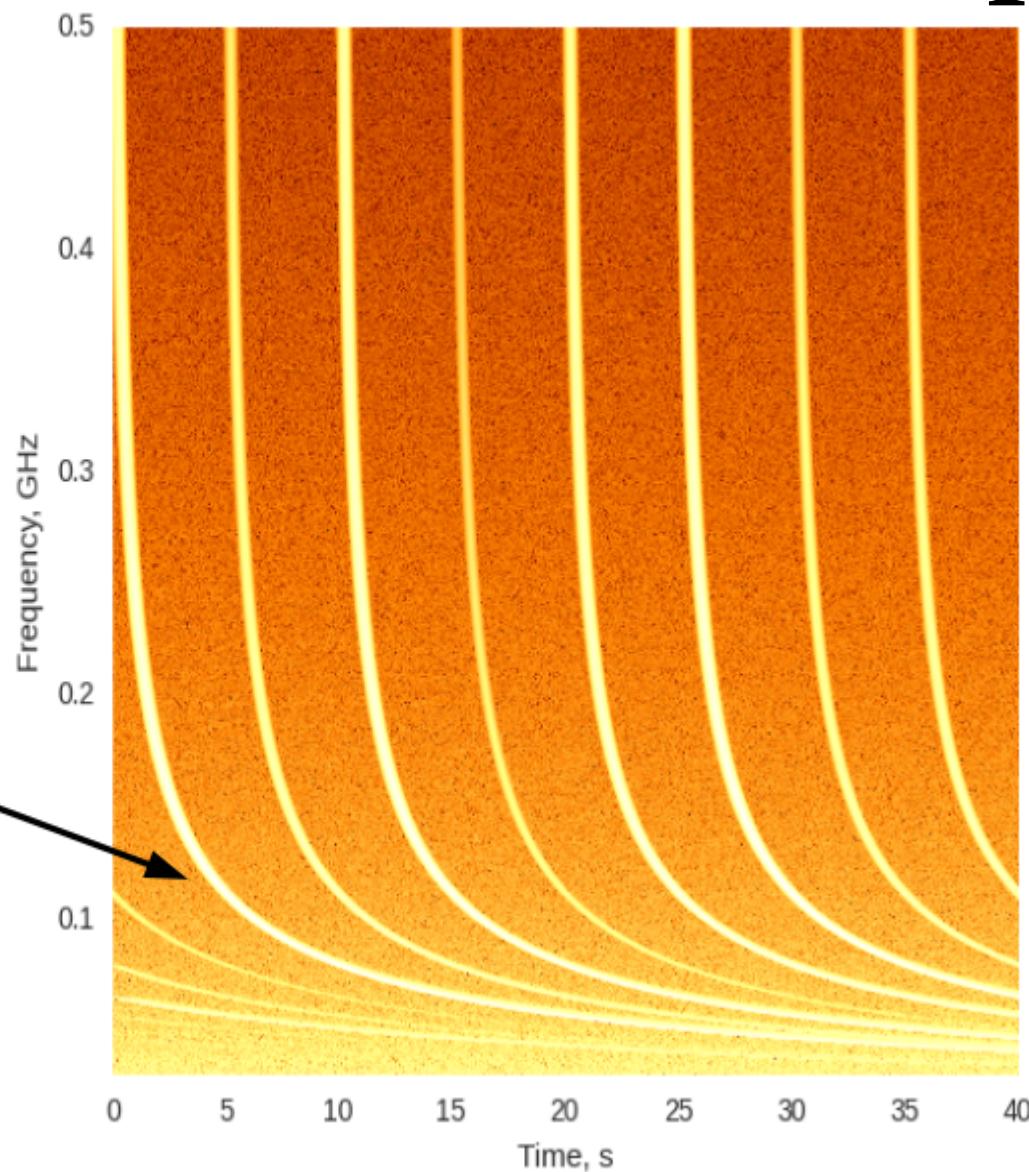
Dispersion

Simulated ultra-broadband pulse recording

$DM = 15 \text{ pc cm}^{-3}$
 $P = 5 \text{ s}$

Dispersive delay

$$\delta t \sim DM/v^2$$



Credit: Anya Bilous

Dispersion

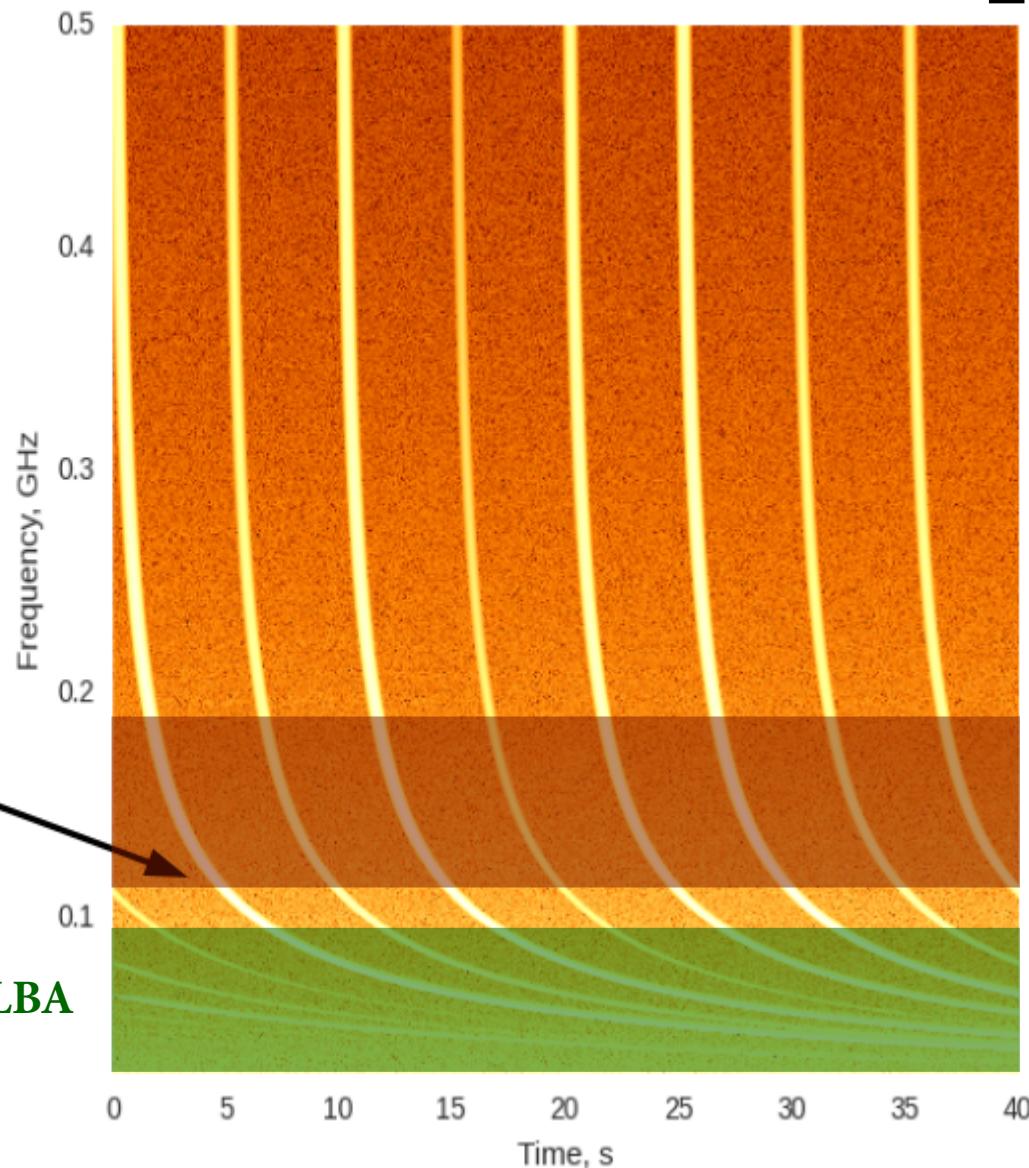
Simulated ultra-broadband pulse recording

$DM = 15 \text{ pc cm}^{-3}$
 $P = 5 \text{ s}$

Dispersive delay

$$\delta t \sim DM/v^2$$

LOFAR LBA



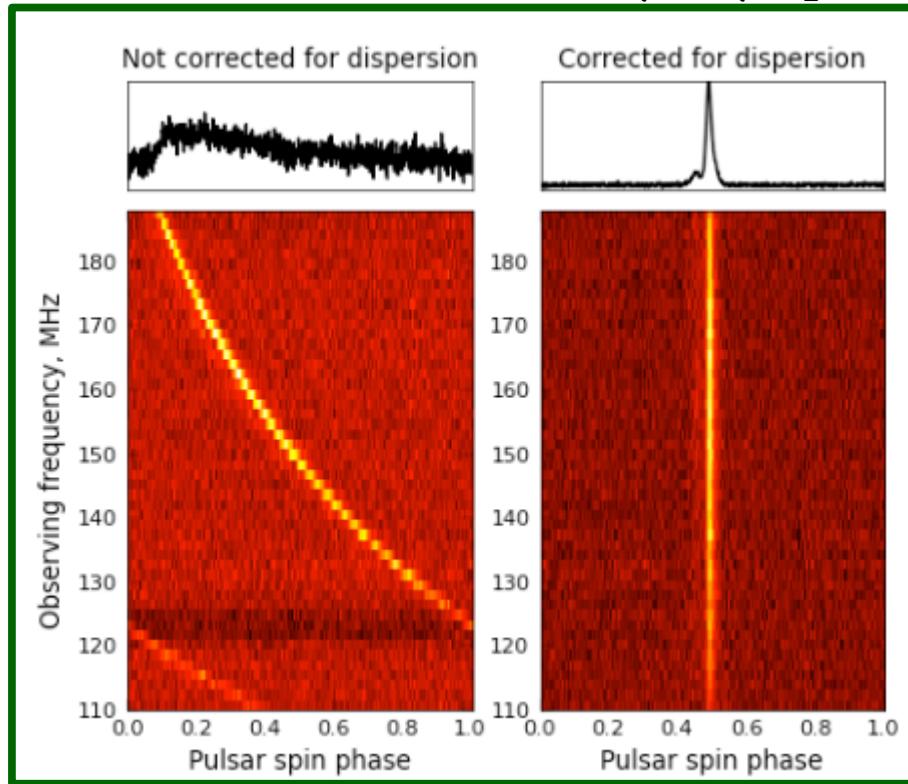
Credit: Anya Bilous

LOFAR HBA

Dispersion

PSR B2021+51

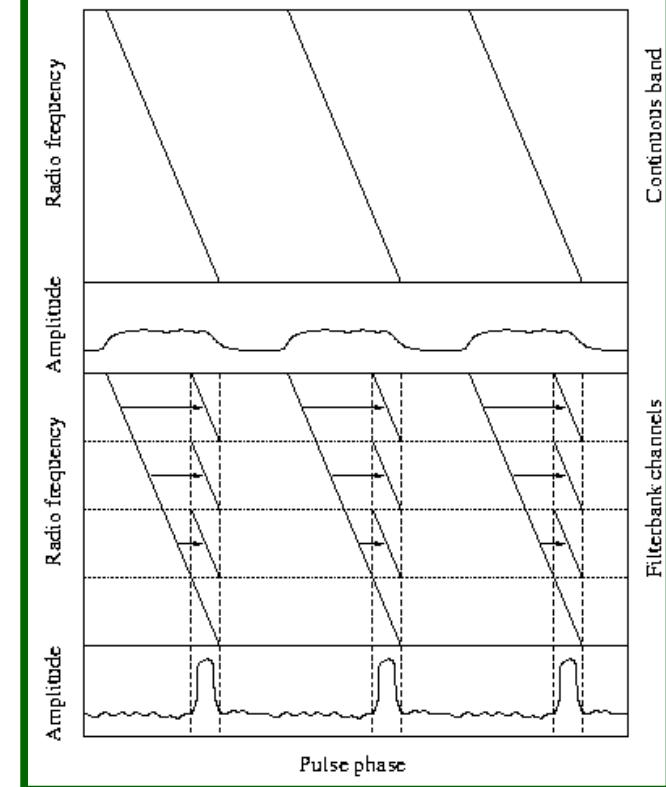
DM is off by only 3 pc/cc!



Credit: Anya Bilous

- DM [pc cm⁻³] measures the integrated column density of free electrons along the line of sight
- Can be corrected using (in)coherent dedispersion

Taken from "Handbook of Pulsar Astronomy" by Lorimer & Kramer

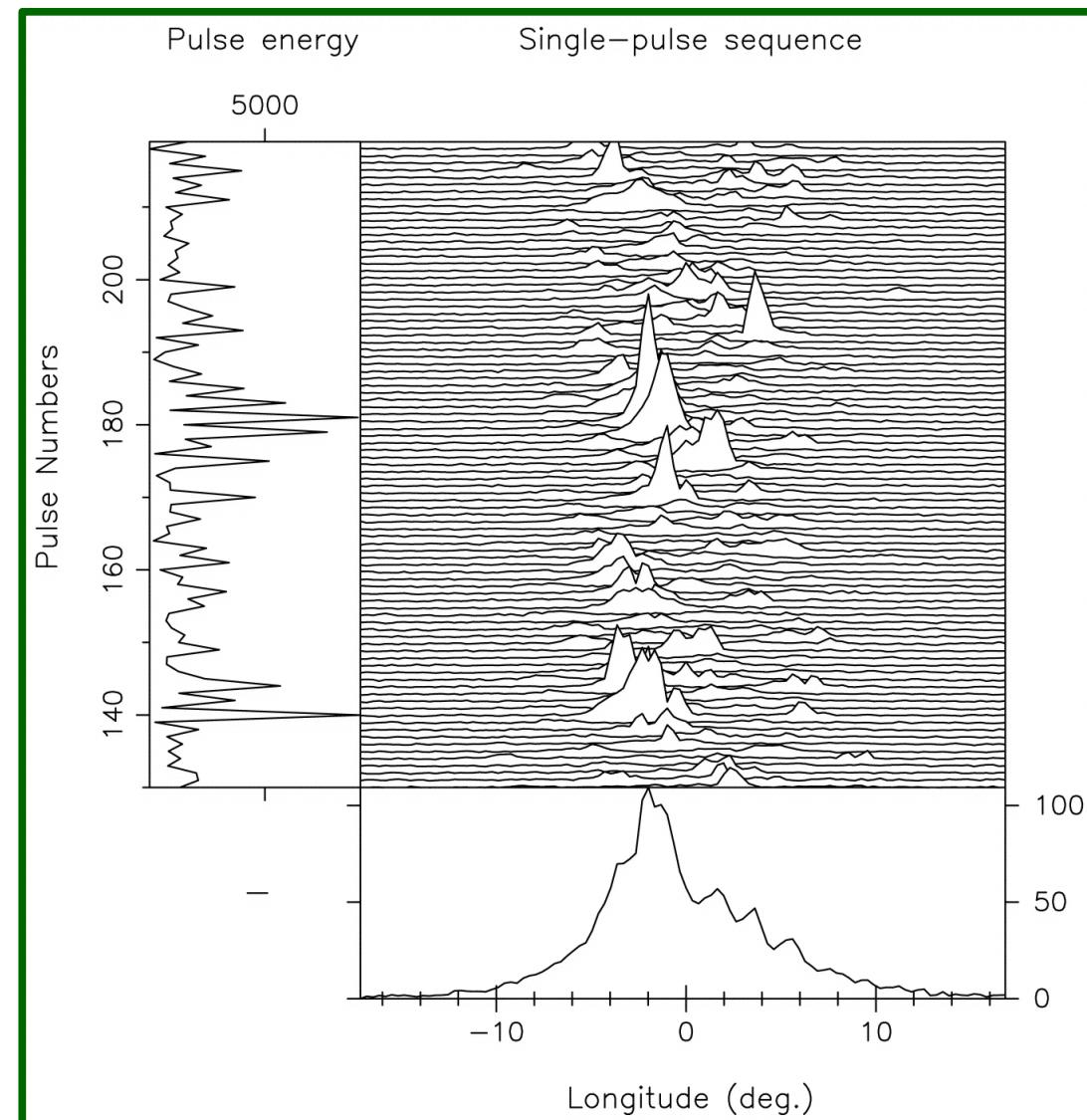
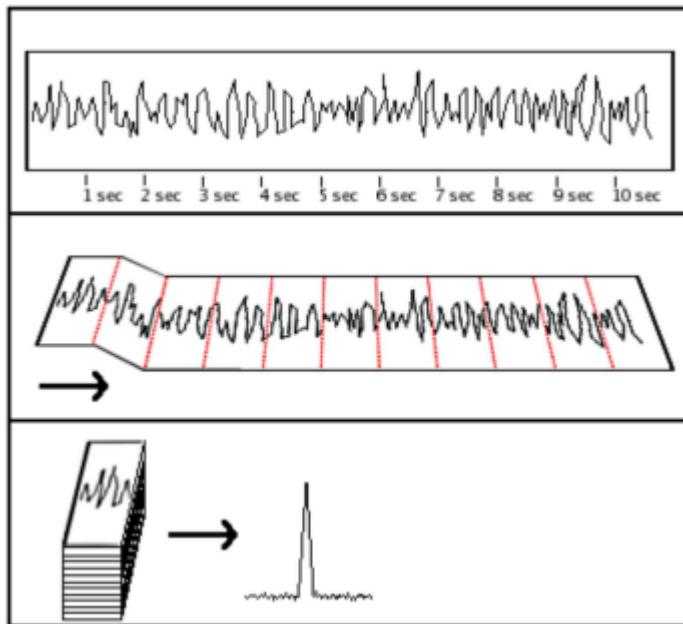


- Incoherent dedispersion – shifting channels in time
- Coherent dedispersion – requires complex-voltage data and is more computationally expensive

Folding

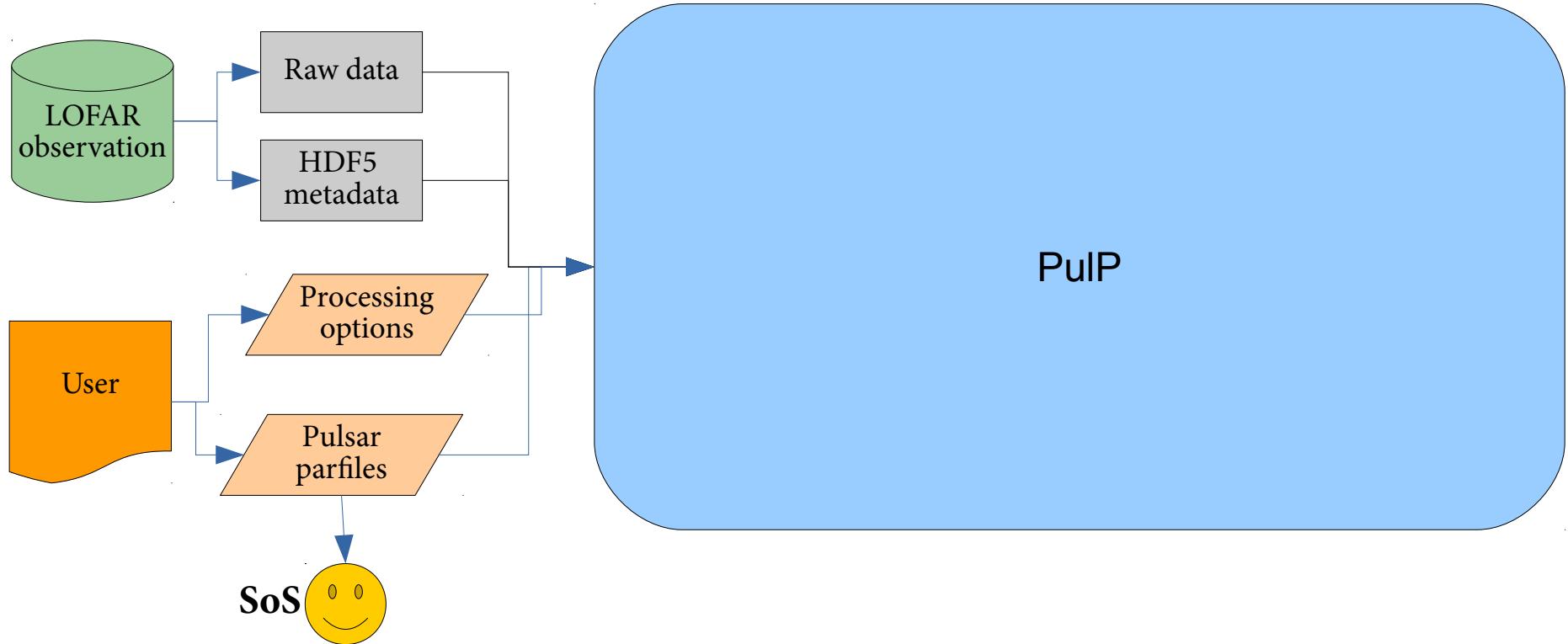
PSR B0943+10

in a nutshell



Deshpande & Rankin (1999)

PulP flowchart (1)



Parfiles: if parfiles are not given toPulP, then based on the target name it will try to find the corresponding pulsar in the ATNF catalog. If no pulsar is found in the catalog, PulP will look for the brightest pulsar in a given SAP and fold it.

Pulsar ephemeris (parfiles)

PSRJ	J0034-0534	
RAJ	00:34:21.8320019	1
DECJ	-05:34:36.81231	1
F0	532.7134297772821597	1
F1	-9.33246370303865163e-16	1
PEPOCH	49550.037311801294202	
POSEPOCH	49550.037311801294202	
DMEPOCH	49550	
DM	13.764894275846959064	1
DM1	0	
PMRA	8.0823671616462304792	0.13169130726699274092
PMDEC	-9.5157740417196312044	0.30750778304651565920
BINARY	ELL1	
PB	1.5892817926966151351	1
A1	1.4377774324431148653	1
TASC	49550.704855759820283	1
EPS1	6.4089927497823510916e-05	1
EPS2	-3.03855316458051531e-05	1
START	55959.632675467299123	1
FINISH	56448.301487586140865	1
TZRMJD	56190.012138005647902	
TZRFRQ	137.15199999999998681	
TZRSITE	t	
TRES	31.090	
EPHVER	5	
CLK	TT(TAI)	
MODE 1		
EPHEM	DE421	
NITS	1	
NTOA	218	
CHI2R	132.9672 207	

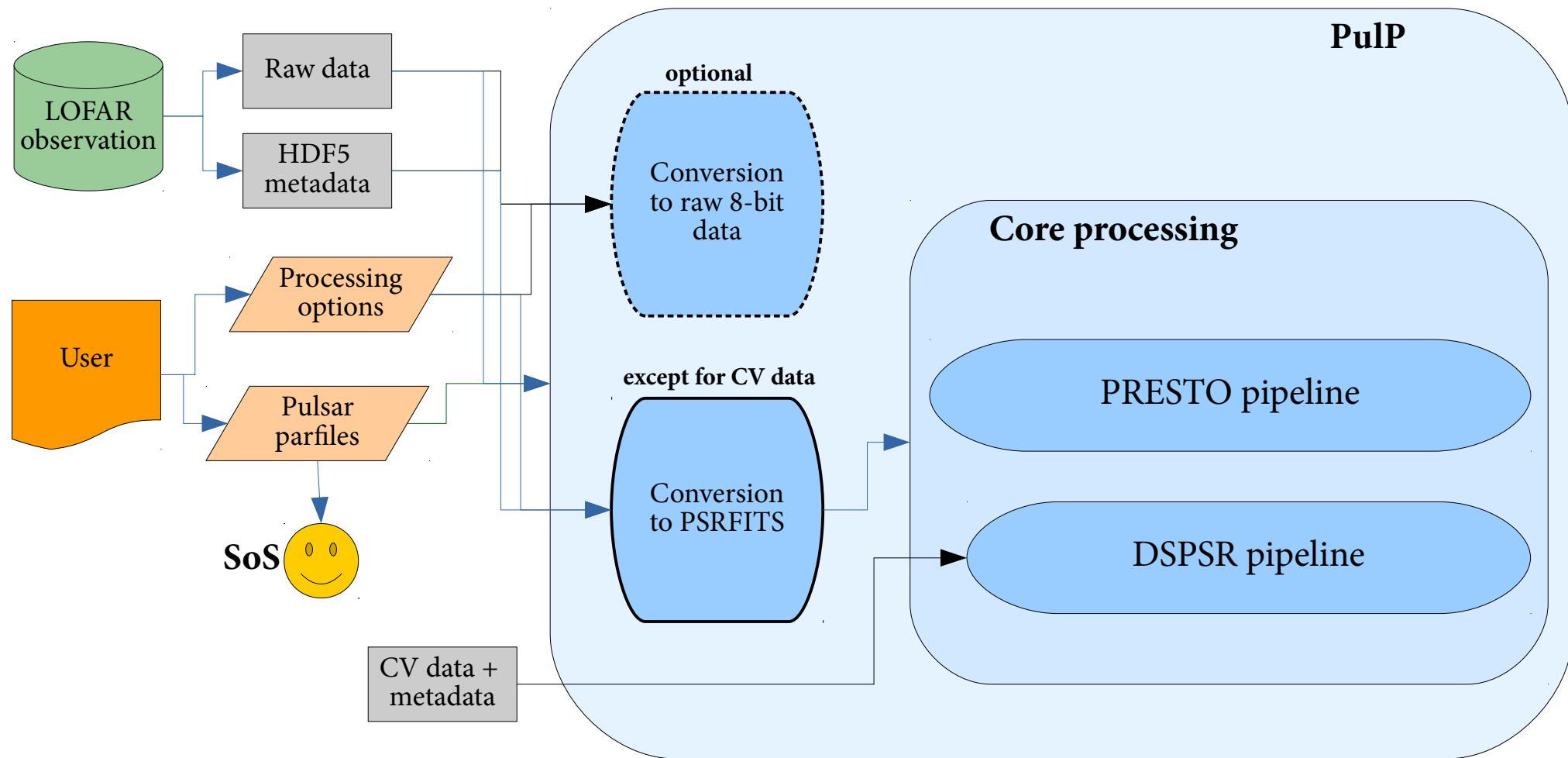
can be as simple as this:

PSR	J1706+35
RAJ	17:07:03.61
DECJ	+35:55:54.5
P0	0.159764851
P1	0.0
PEPOCH	58244.04308936660
DM	19.240
EPHEM	DE405
CLK	UNCORR

Input raw data

- HDF5 format
- Header information (metadata) is stored in `*_bf.h5` file
- The raw data itself is stored in `*_bf.raw` file
- This `.raw` file is linked from within `.h5` file and can be accessed directly via opened `.h5` file
- Filename structure:
→ Lnnnnnn_SAPxxx_Byyy_Sz_Pmmm_bf.h5
 - Lnnnnnn – LOFAR observation ID (ObsID)
 - xxx – Sub-array pointing (SAP) number
 - yyy – Tied-array beam (TAB) number
 - z – Stokes parameter, can only take values 0,1,2,3
 - Stokes I observation – have only S0 files
 - Stokes IQUV observation: S0 – I, S1 – Q, S2 – U, S3 – V
 - Complex-voltage data: S0 – Xreal, S1 – Ximag, S2 – Yreal, S3 – Yimag
 - mmm – Frequency part, i.e. when every file has only fraction of subbands

PulP flowchart (2)



Data conversion

- Conversion to raw 8-bit data (optional)
 - *digitize.py*
 - written by Marten van Kerkwijk
 - available at:
<https://github.com/mhvkv/scintellometry/blob/master/scintellometry/lofar/digitize.py>
 - *digitize.py -s 5 -o <output dir> <input .h5>*
- Conversion from raw 32-bit data to PSRFITS (for non-CV data)
 - custom-made program *2bf2fits*
 - written by Tom Hassall, Patrick Weltevrede, with contribution from Vlad Kondratiev
 - currently available at LOFAR Users Software Repository
 - will make it available at Github as well
 - does not save scales/offsets in PSRFITS
 - needs major revisiting...
 - Command example (very detailed input):
 - *2bf2fits -CS -H -append -nbits 8 -A 100 -sigma 3 -nsubs 400 -sap 0 -tab 0 -stokes 0 -o L667444_SAP0_BEAM0 -nsamples 24 -nchans 16 -ra 2.15980858832 -dec 1.30000703891 -psr B0809+74 -clock 200 -band HBA_110_190 -startdate 2018-09-12 -starttime 20:17:00.000000000 -samptime 0.0104858 -duration 299.977 -subs 54..453 -obsid L667444 -observer Pizzo /data/projects/PipelineTests/L667444/cs/L667444_SAP000_B000_S0_P000_bf.raw*

DSPSR Pipeline (1)

for every TAB, PSR, and frequency part

dedispersion/
/folding

```
dspsr -O <outputname> -b <nbins> -A -L <tsubint> -q -E <parfile> -t 2 <dspsr extra user options>  
OR: dspsr -O <outputname> -b <nbins> -A -q -E <parfile> -t 2 <dspsr extra user options: -s + other opts>
```

if Single Pulse Analysis = TRUE

creating
filterbank file

```
digifil -q -B 512 -b 8 -F <nchan> -D <dm> -o <outputname> <digifil extra user options>
```

Input data for dspsr:

- **CV data:** any one .h5 file for a given frequency part;
- **Stokes I/IQUV data:** PSRFITS file from the previous conversion step.

combining
frequency
parts

for every TAB
and PSR

Summary
plots

Tarball for
a given TAB and
frequency part

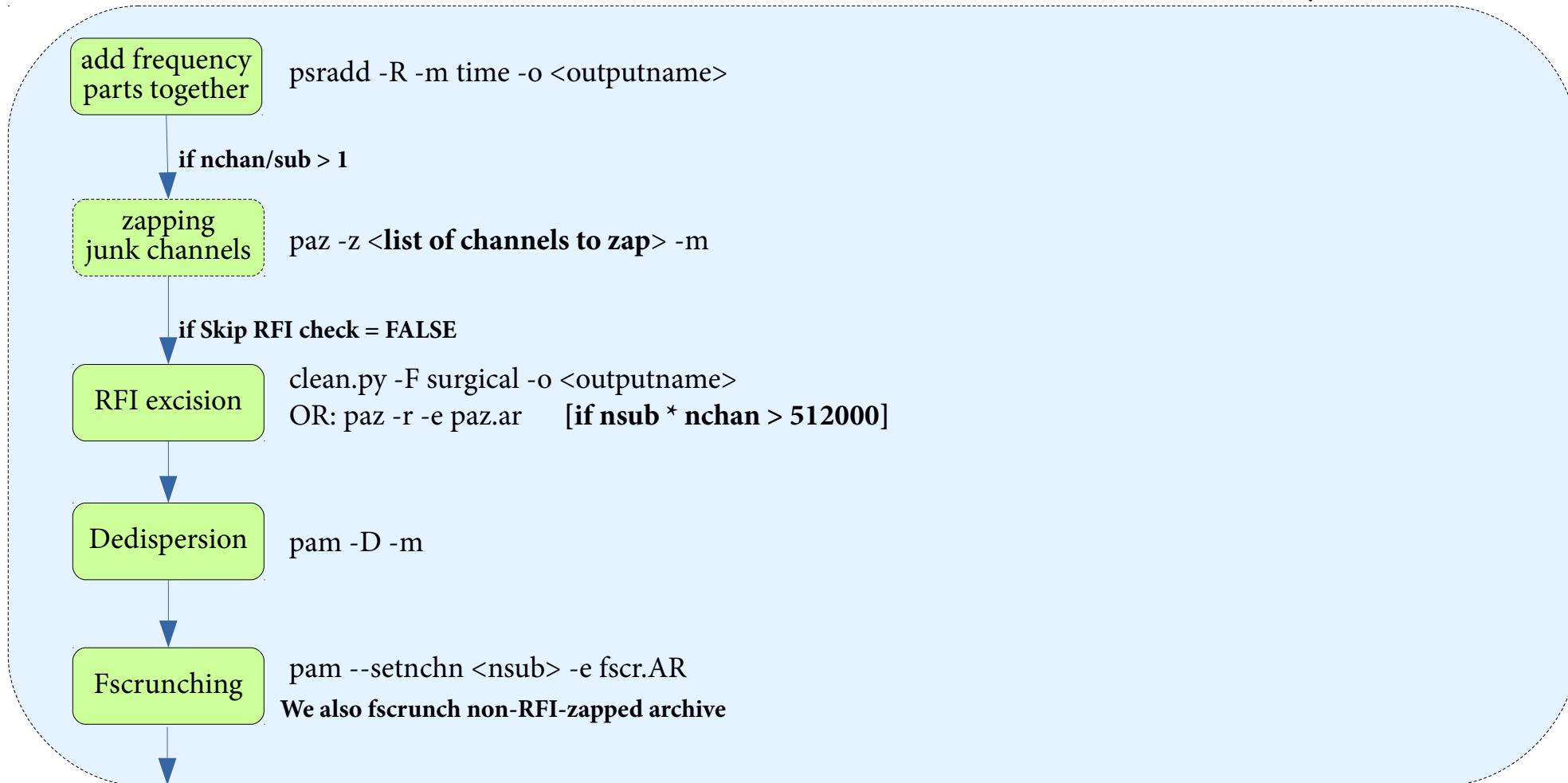
<nbins> - calculated automatically based on the sampling time and F0/P0 from the parfile.

Maximum possible <nbins>=1024

<nchan> - number of channels in a given frequency part. If number of channels = 1, then <nchan>=2

DSPSR Pipeline (2)

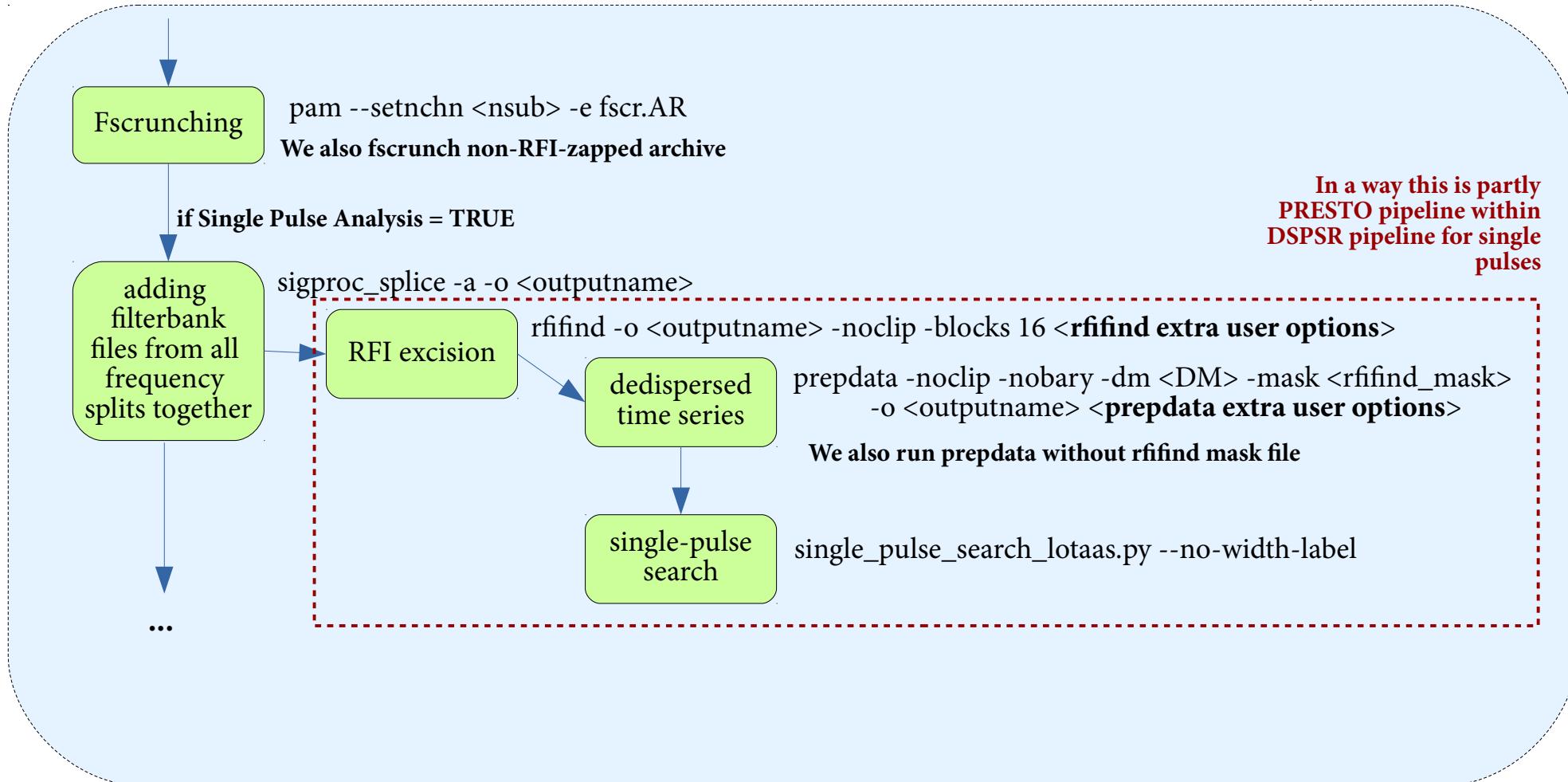
combining frequency parts;
for every TAB, PSR



<list of channels to zap> - if there are 16 chan/sub, we need to zap every 16th channel, then list becomes «0 15 31 47...»
This is necessary, as when 2nd PPF is used, the first channel in each subband gets corrupted

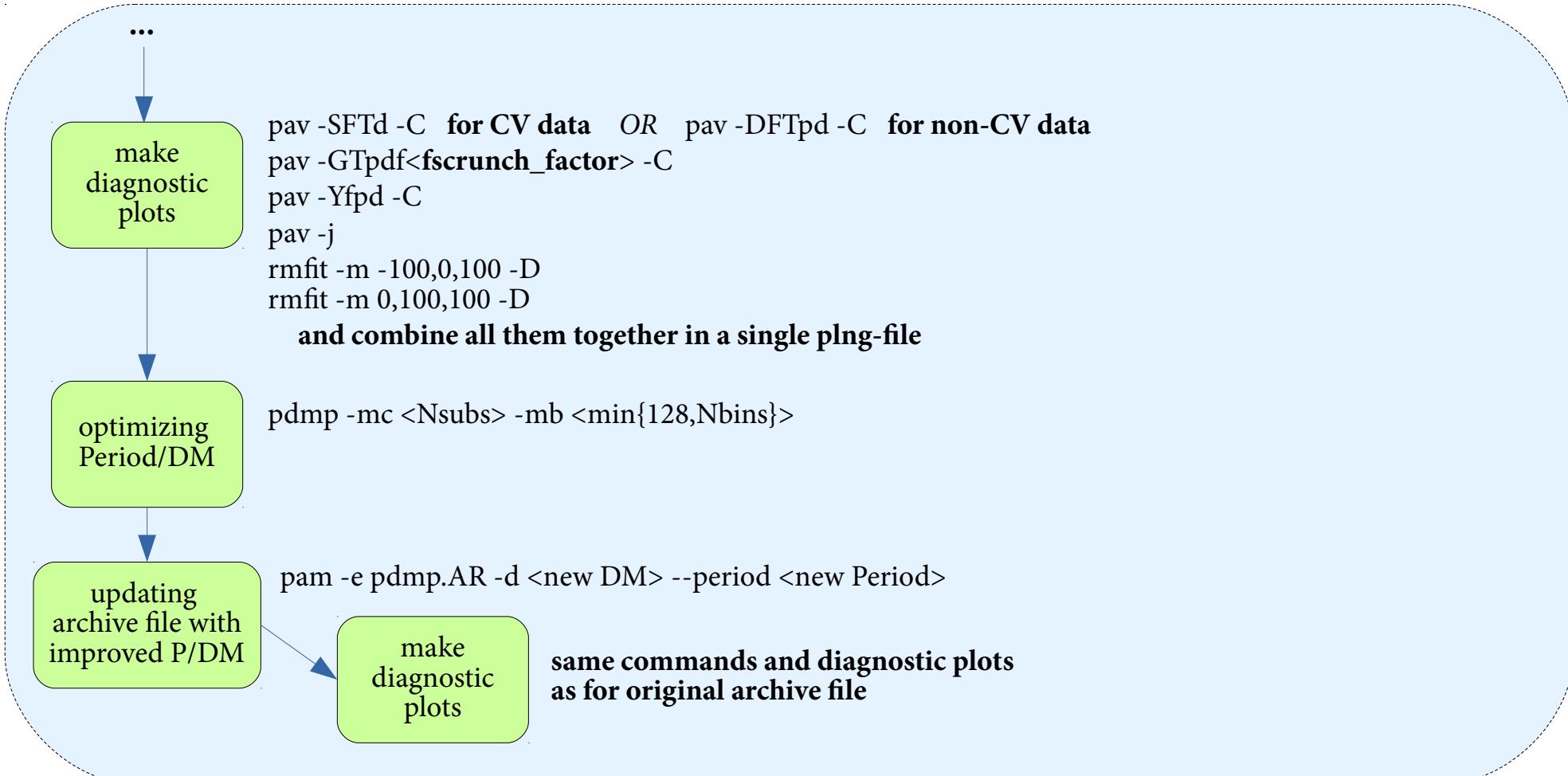
DSPSR Pipeline (2, cont.)

combining frequency parts;
for every TAB, PSR



DSPSR Pipeline (3, cont.)

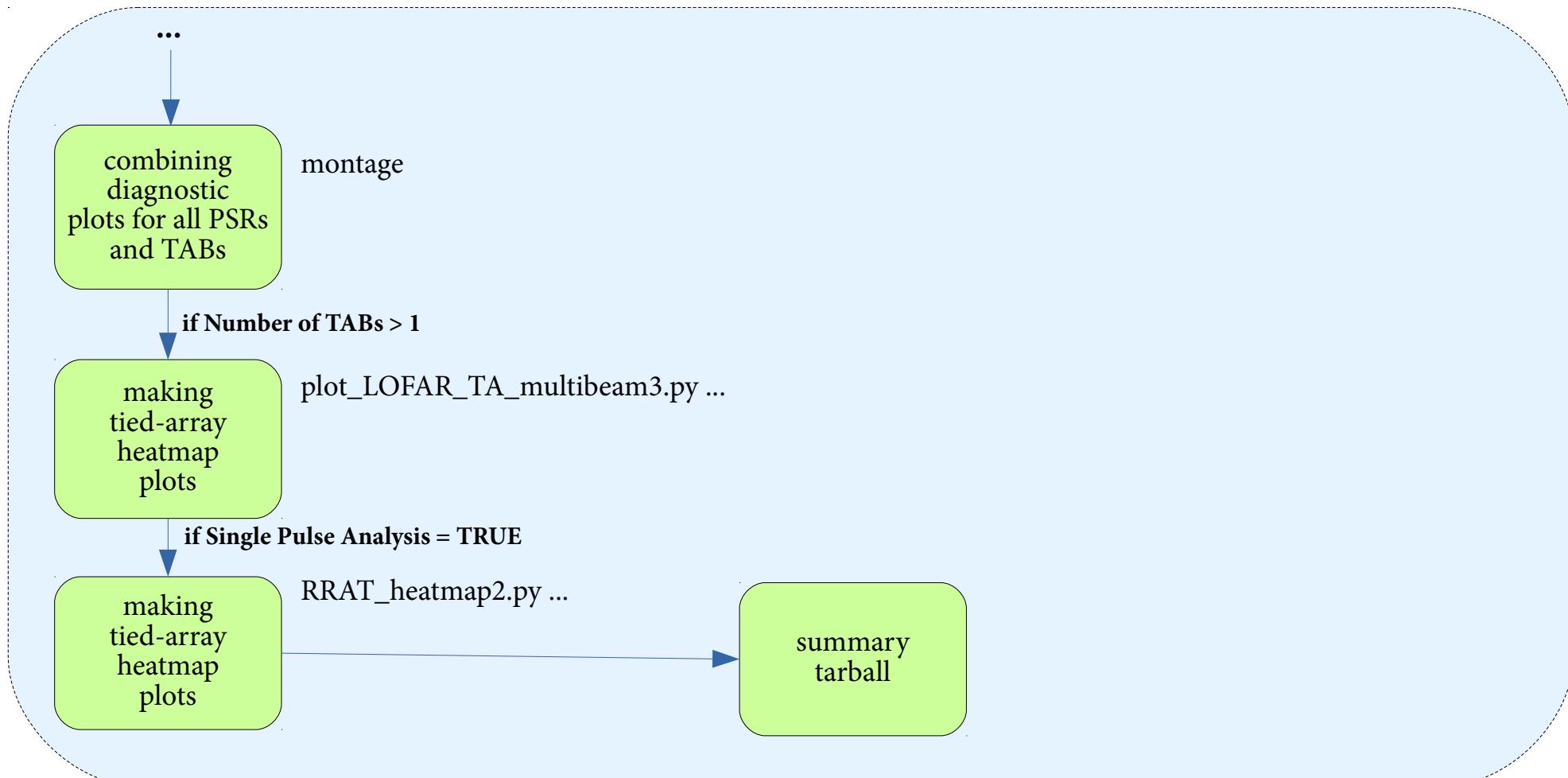
combining frequency parts;
for every TAB, PSR



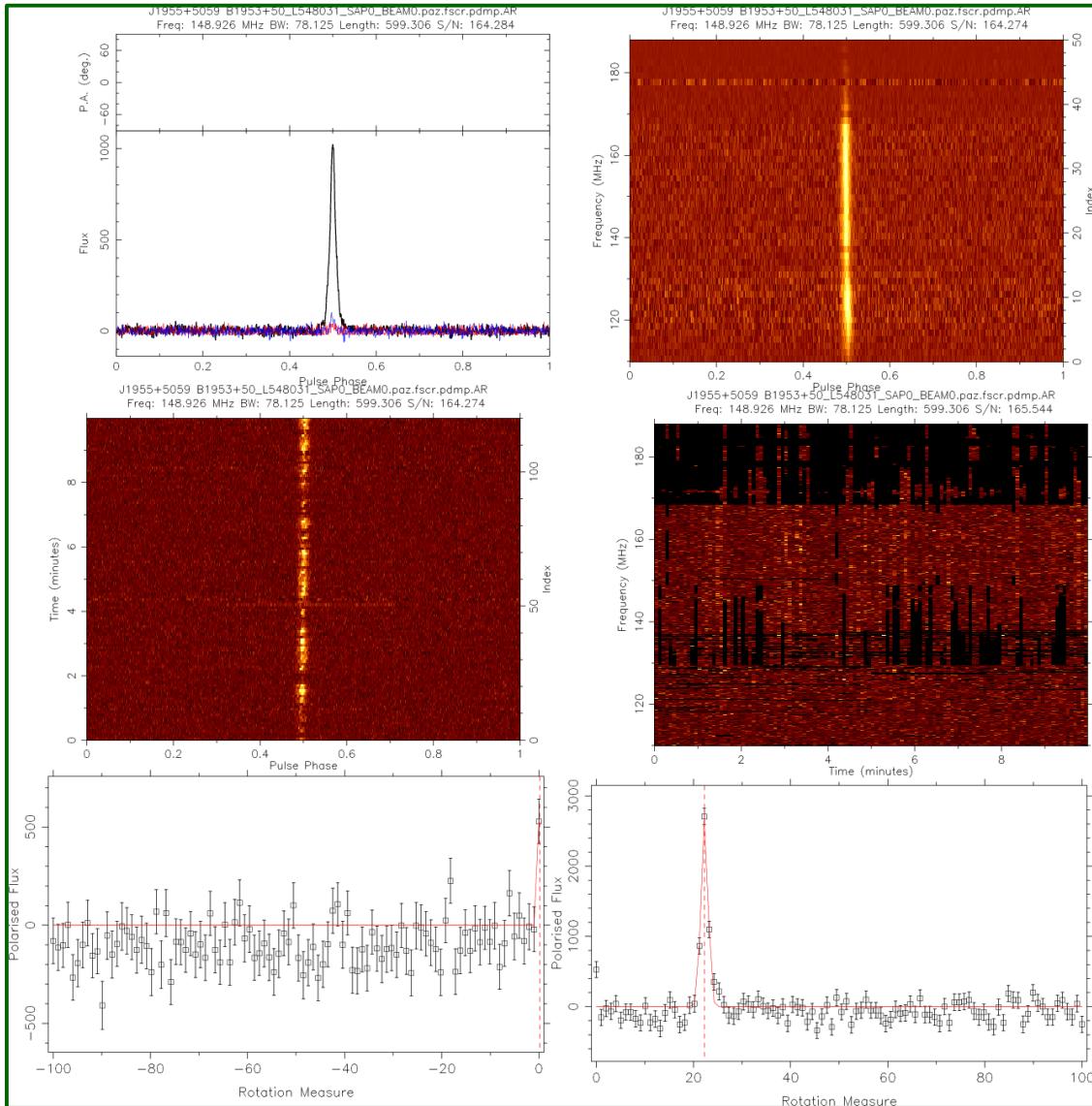
<fsrch_factor> = Nsubs / X, where X = highest common denominator of Nsubs between 1 and min{Nsubs, 63}

DSPSR Pipeline (4)

Summary plots



Diagnostic plots (1)

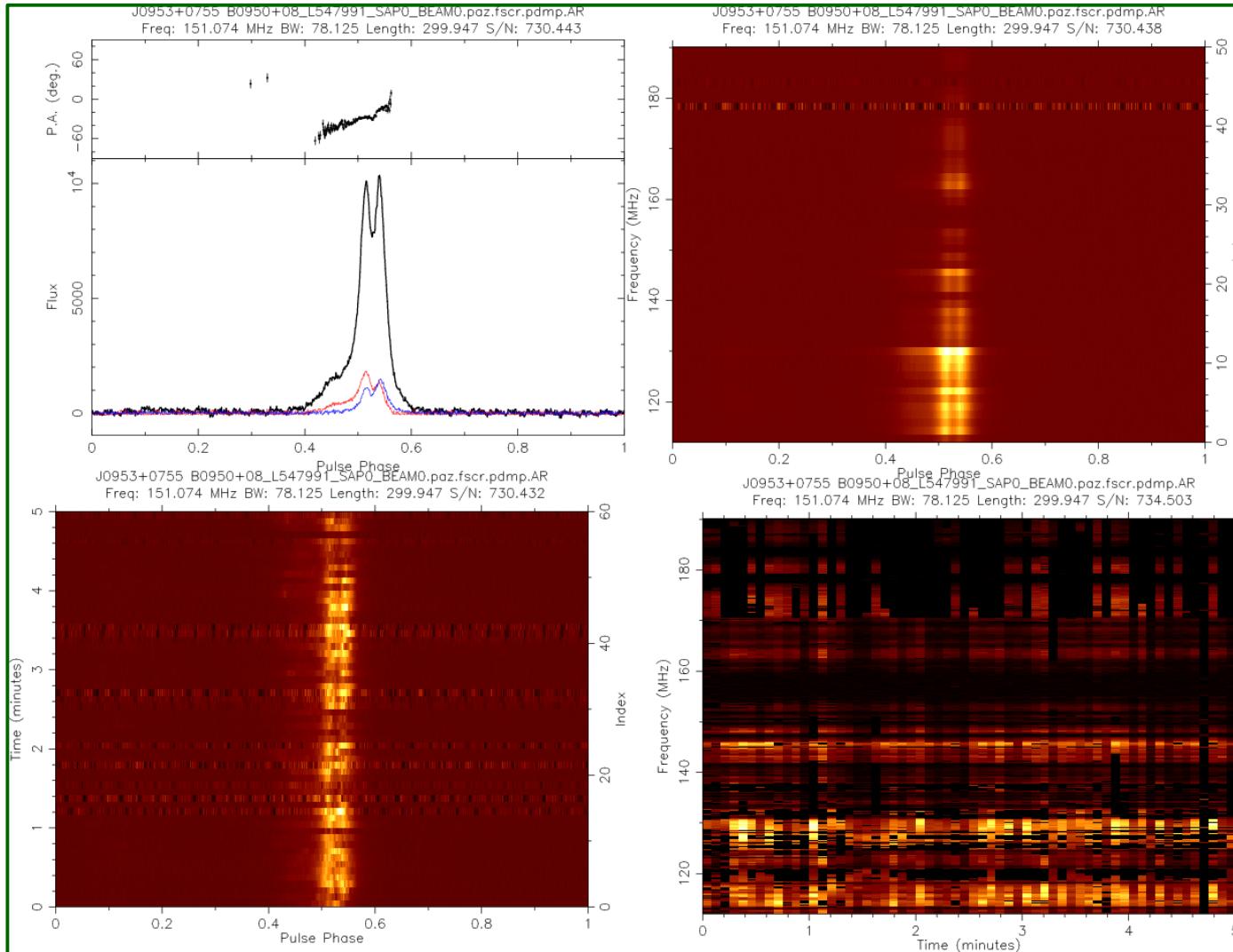


status.png

*_diag.png

*_diag_pdmp.png

Diagnostic plots (2)

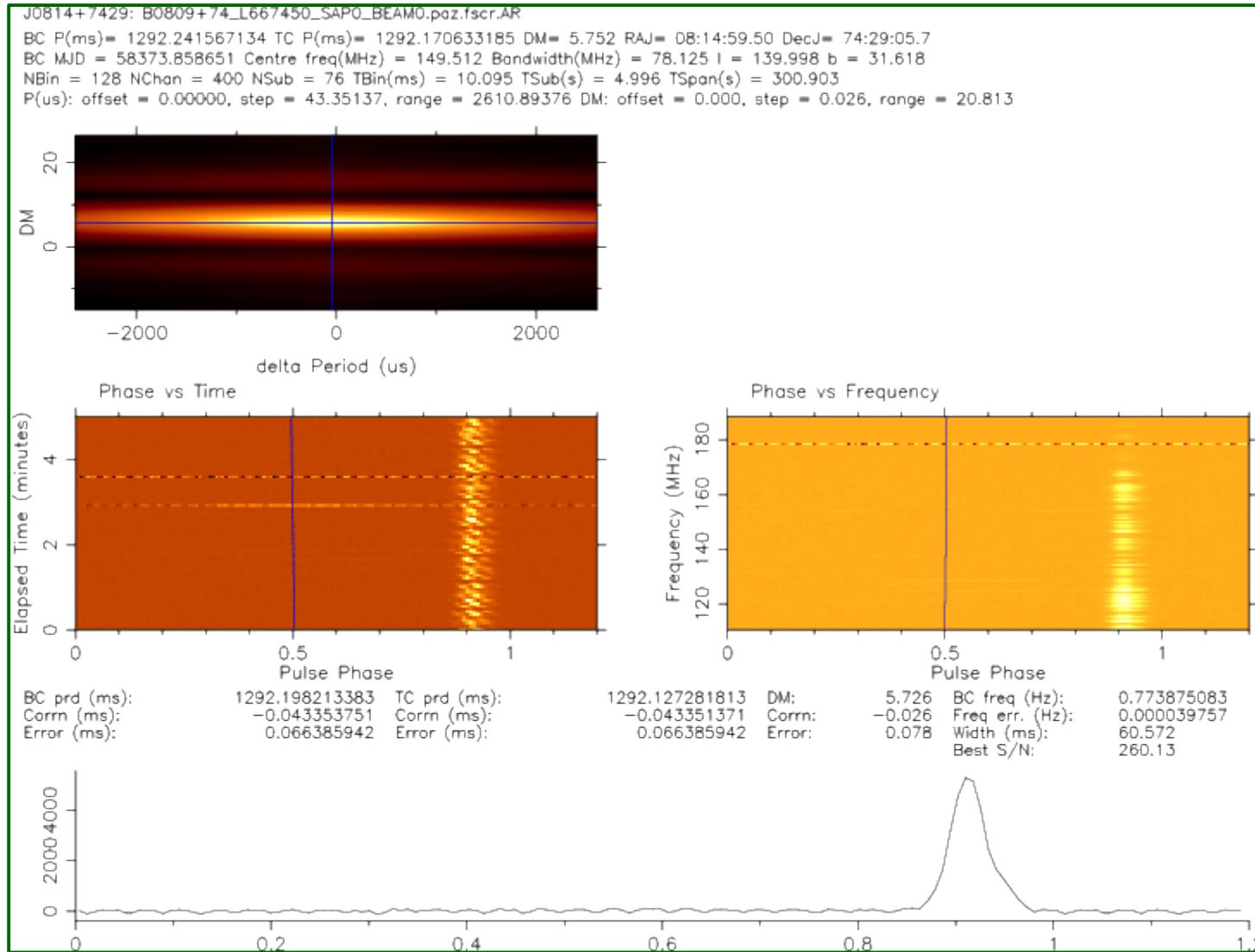


status.png

*_diag.png

*_diag_pdmp.png

Diagnostic plots (3)



*_pdmp.ps

PRESTO Pipeline (1)

for every TAB, and frequency part

RFI excision,
zapping
junk channels

```
rfifind -o <outputname> -psrfits -noclip -blocks 16 -zapchan 0:X:Y <rfifind extra user options>  
X = Nchan-1  
Y = Nchan/sub
```

if Skip dynamic average = FALSE

making
diagnostic
dynamic
spectrum

```
subdyn.py -psrfits -saveonly -n <samples_to_average> ...
```

for every PSR

Dedispersion/
/Folding

```
Prepfold -noscales -nooffsets -noxwin -psr <psrname> -par <parfile> -n <nbins> -nsub <nsubs>  
-fine -nopdsearch -mask <rfifind_mask> -o <outputname> <prepfold extra user options>
```

We also run prepfold without rfifind mask file

make
diagnostic
plots

convert montage → profile thumbnails

...

<**nbins**> - calculated automatically based on the sampling time and F0/P0 from the parfile.
Maximum possible <**nbins**>=1024

<**nsubs**> - if Nchan > 512, nsubs = 512. Otherwise, nsubs = Nchan

PRESTO Pipeline (2, cont.)

for every TAB, and frequency part

...

if Single Pulse Analysis = TRUE

dedispersed time series

for every PSR

```
prepdata -noscales -nooffsets -noclip -nobary -dm <DM> -mask <rfifind_mask>  
-o <outputname> <prepdata extra user options>
```

We also run prepdata without rfifind mask file

single-pulse search

```
single_pulse_search_lotaas.py --no-width-label
```

<nsub> – greatest common denominator of Nchan
between 1 and 1024
<lodm> – DM - 0.5*dmstep*numdms.
If lodm<=0, then lodm=0.01

dedispersed time series
for a range of DMs

```
prepdata -noscales -nooffsets -noclip -nobary -dm 0.0 -mask <rfifind_mask> -o <outputname>  
<prepdata extra user options>
```

We also run prepdata for DM=0.0 without rfifind mask file

```
prepsubband -noscales -nooffsets -noclip -nobary -nsub <nsub> -lodm <lodm> -dmstep 0.01  
-numdms 1000 -mask <rfifind_mask> -o <outputname> <prepsubband extra user options>
```

We also run prepsubband without rfifind mask file

single-pulse search

```
single_pulse_search_lotaas.py -p -g *.dat  
single_pulse_search_lotaas.py -t 5.5 —no-width-label -g *.singlepulse
```

Also running similar command but excluding DM=0

PRESTO Pipeline (3)

for every TAB, and frequency part

for every PSR

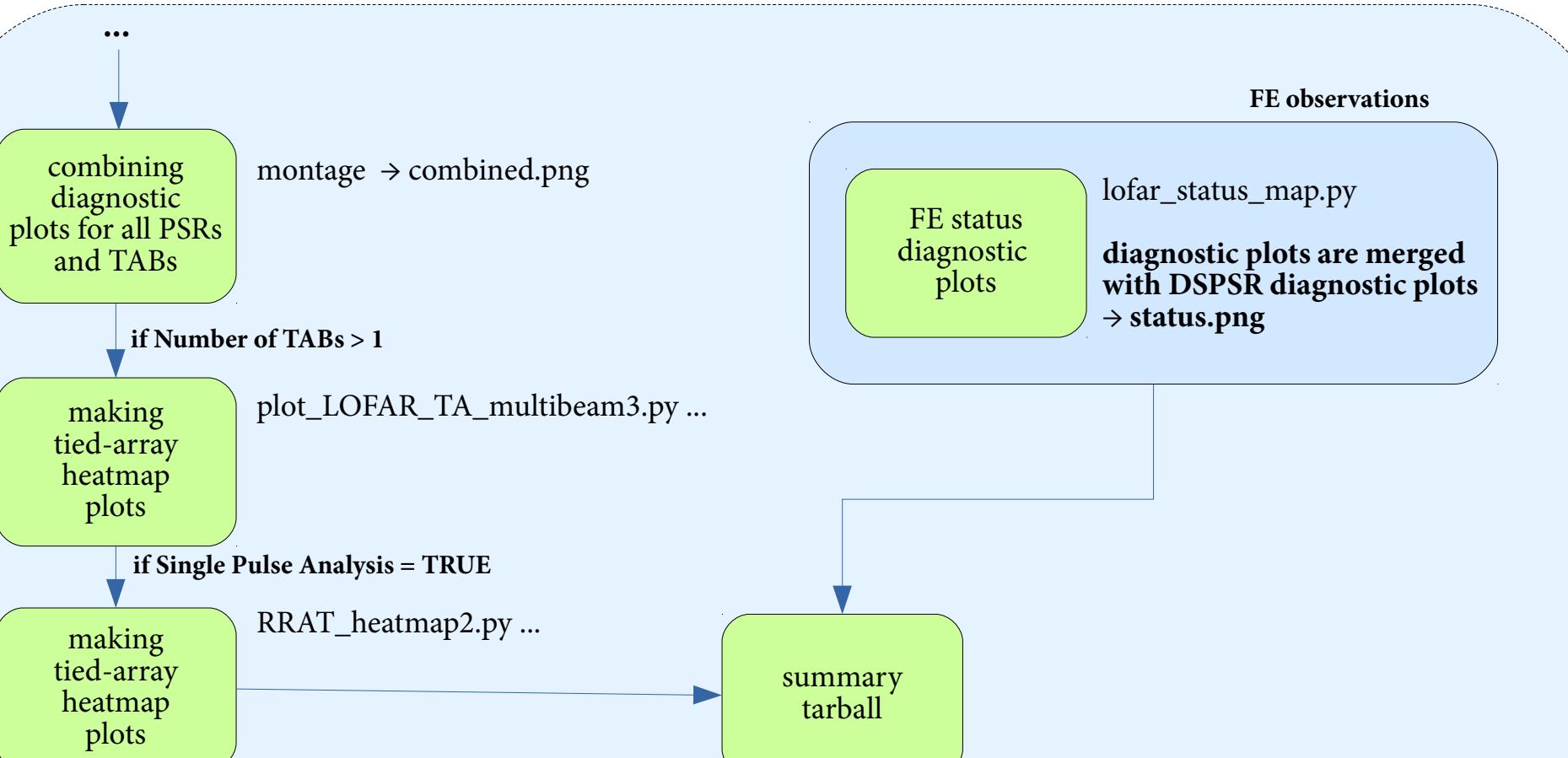
single-pulse
search

Tarball for
a given TAB and
frequency part

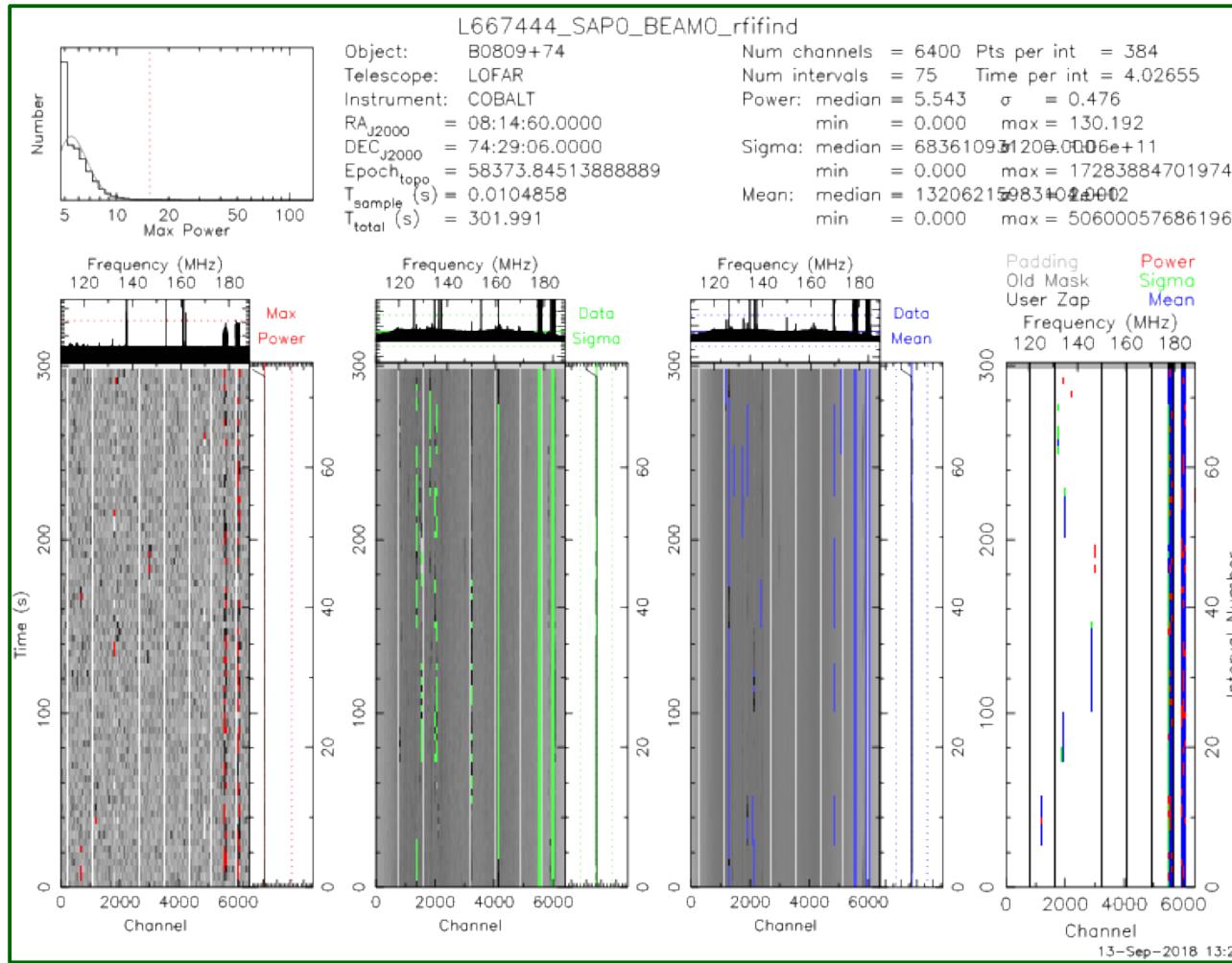
Summary
plots

PRESTO Pipeline (4)

Summary plots

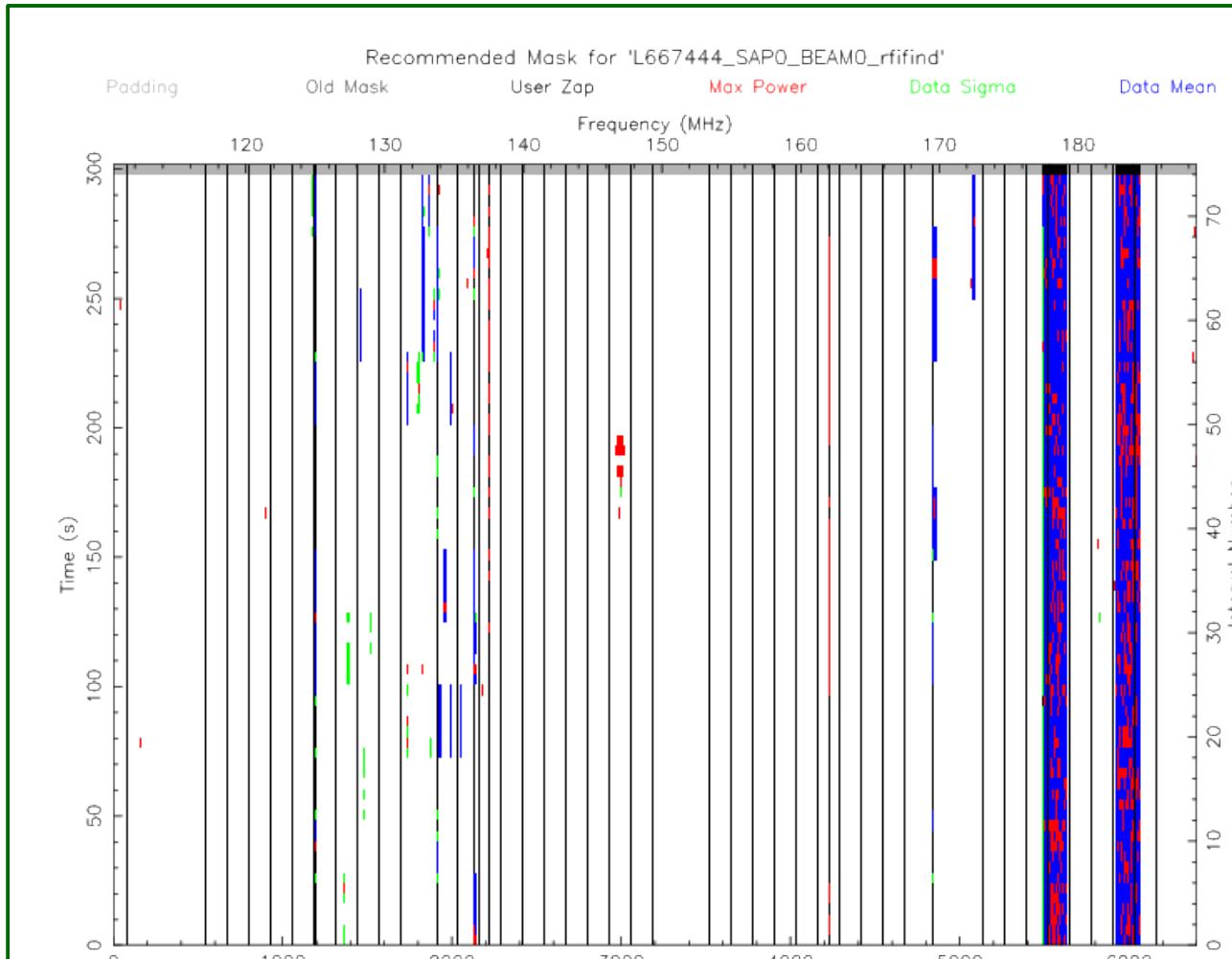


Diagnostic plots (1)



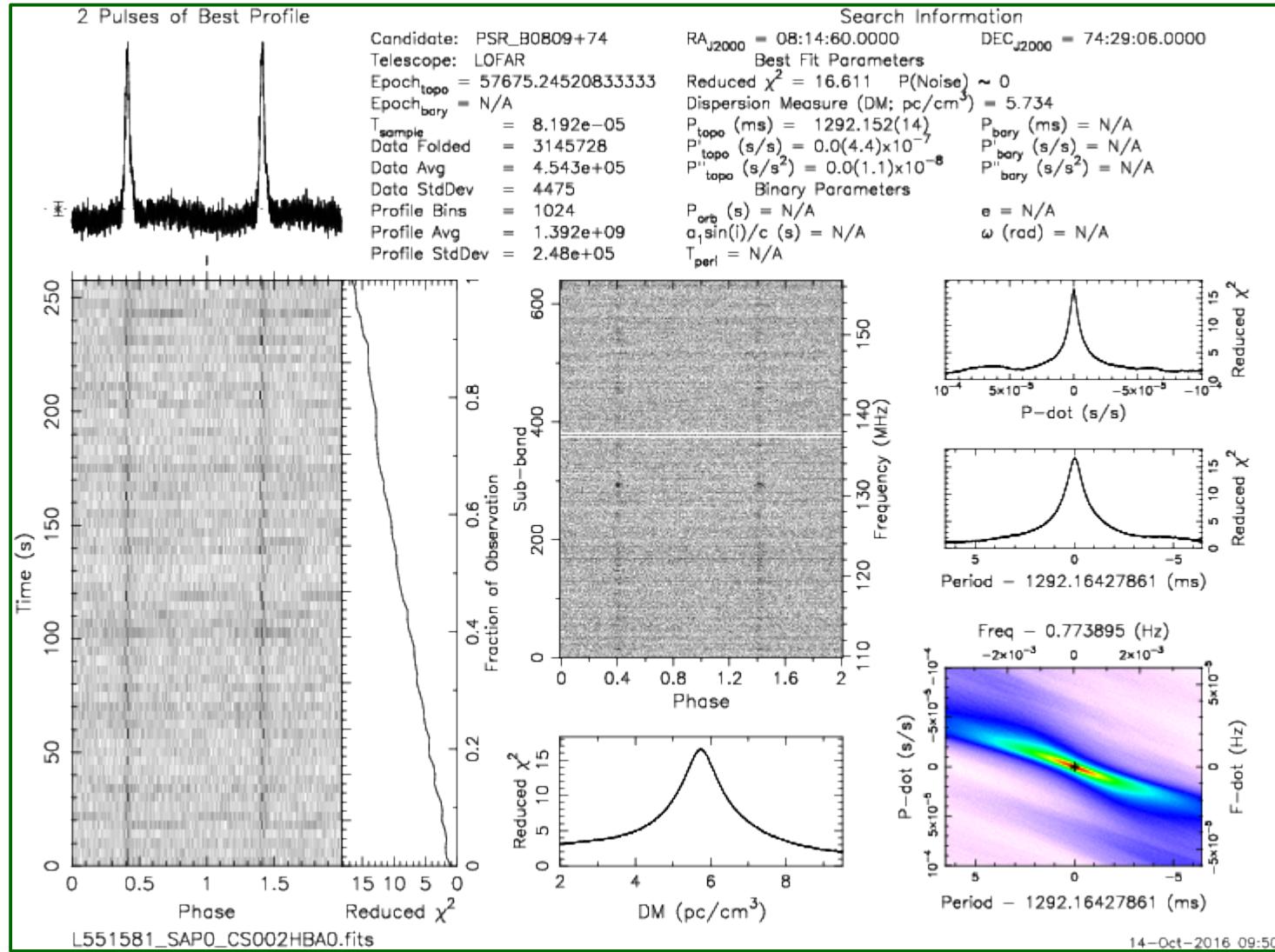
*_rfifind.ps

Diagnostic plots (1, cont.)



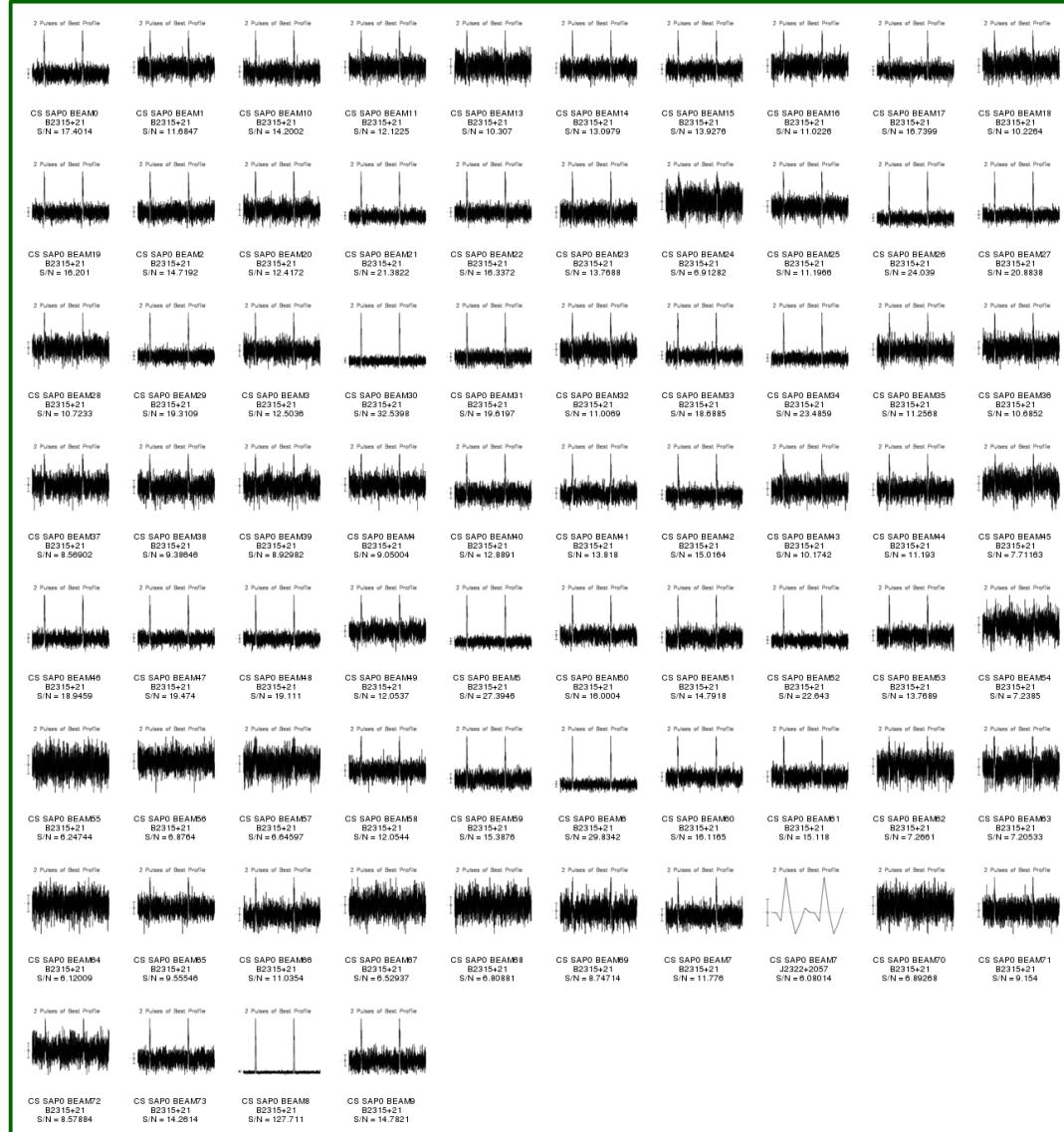
*_rfifind.ps

Diagnostic plots (2)



*_pdf.png

Diagnostic plots (3)

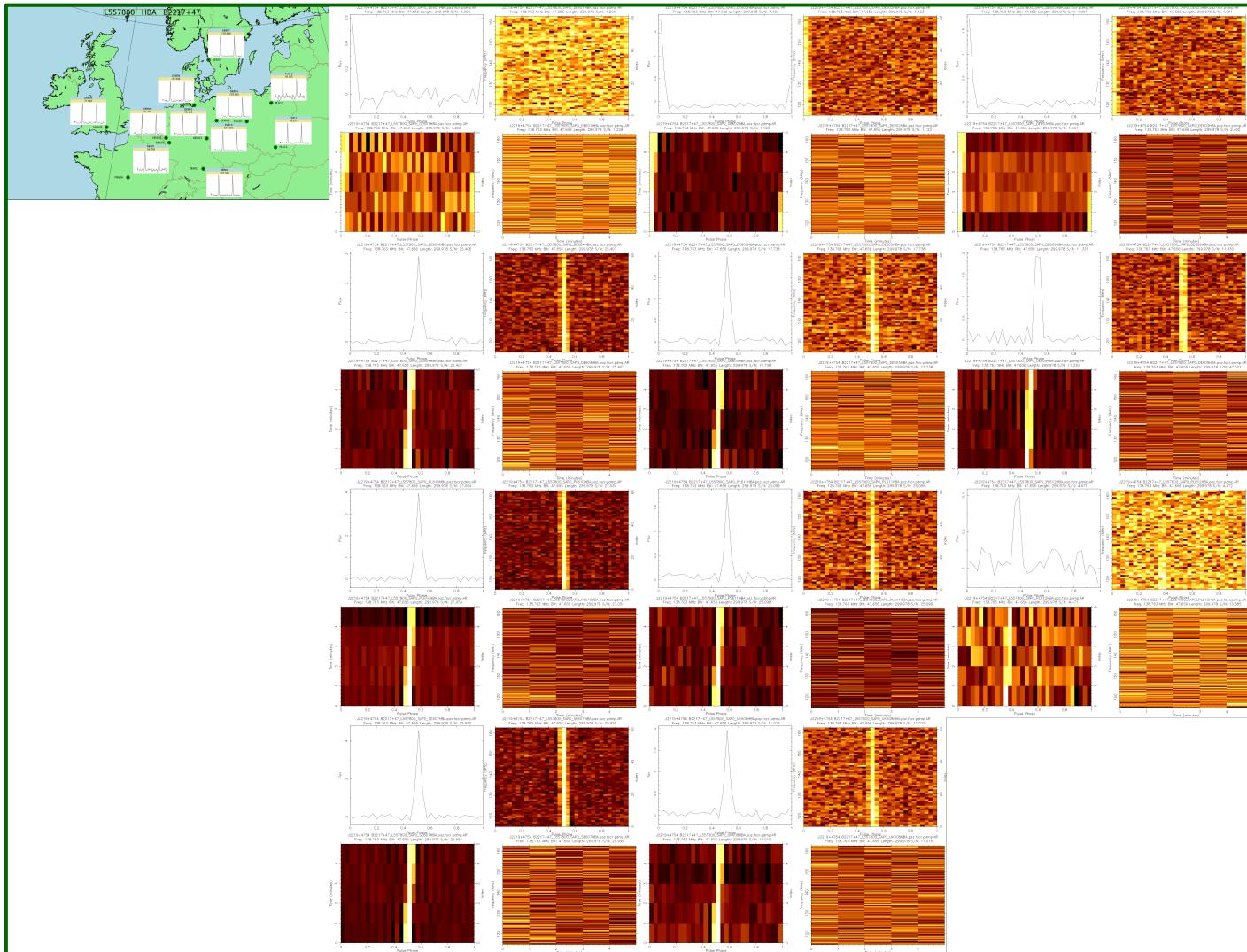


combined.png

multiple TABs
prefold

for FE observations

Diagnostic plots (4)

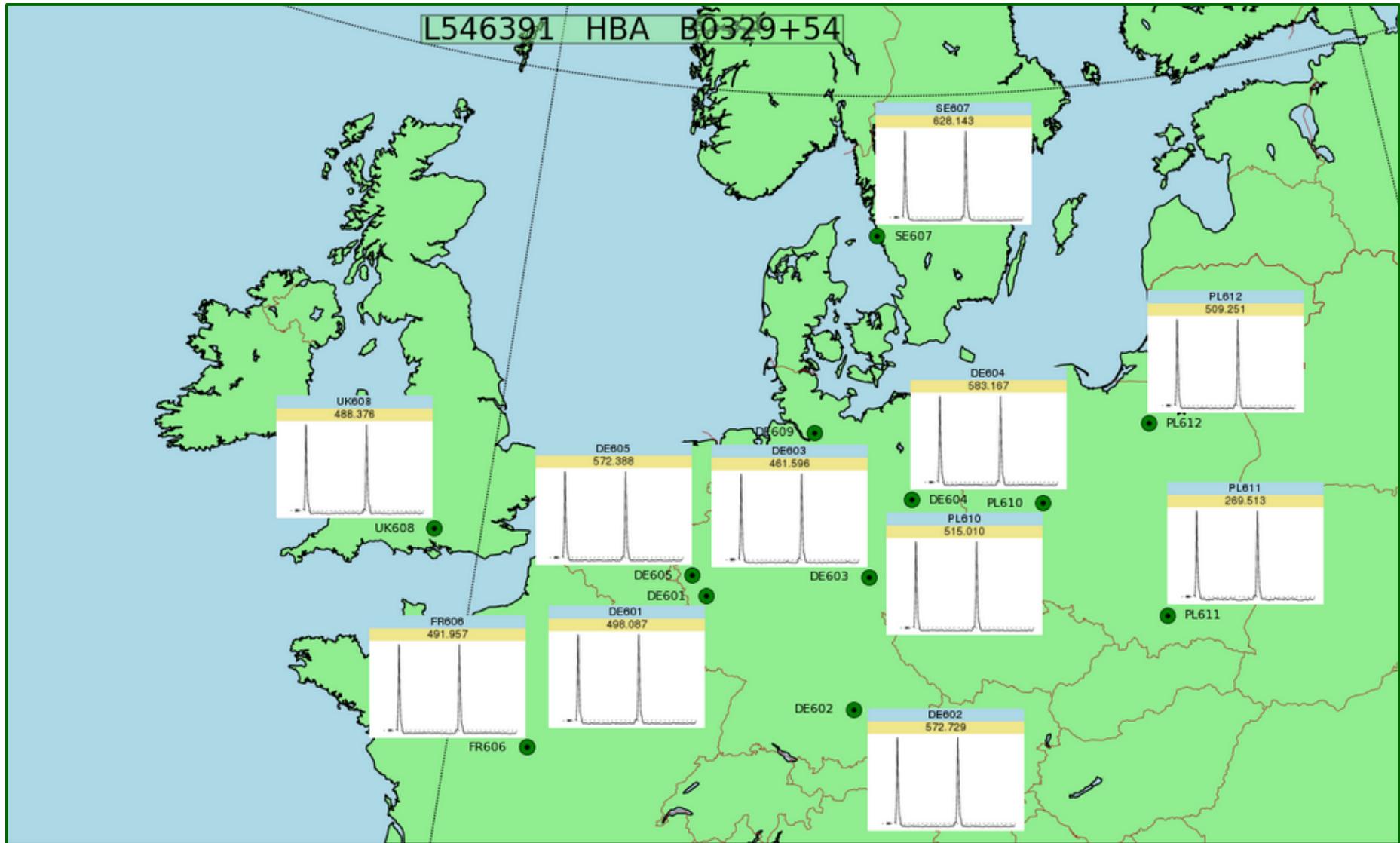


status.png

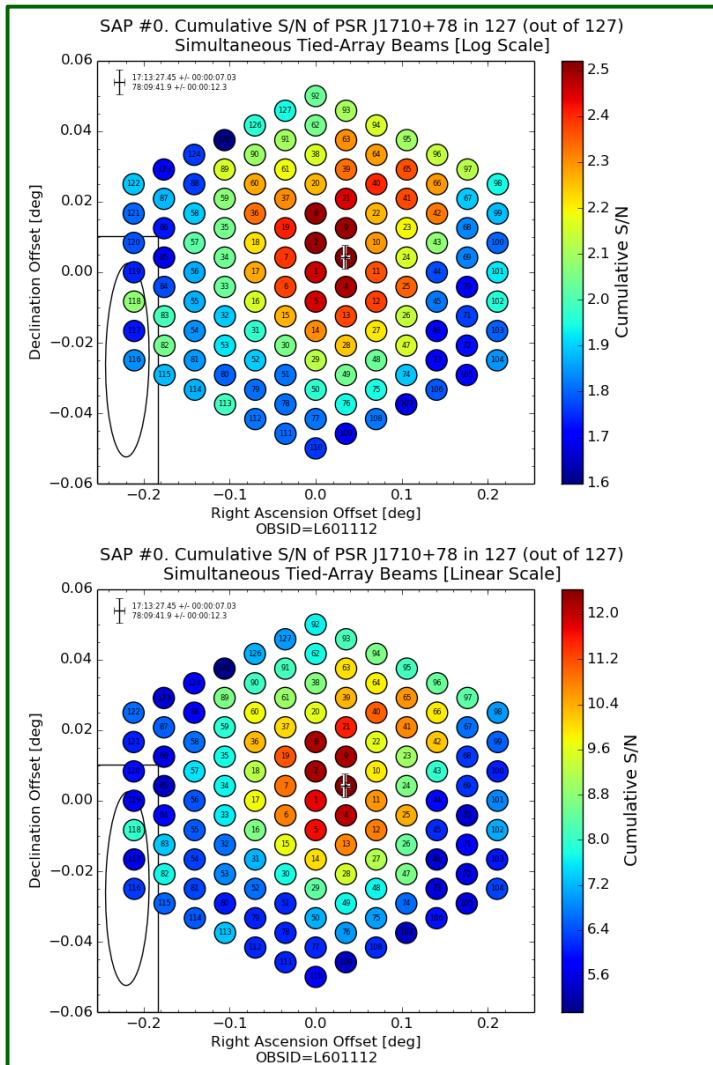
for FE observations

Diagnostic plots (4, cont.)

status.png



Diagnostic plots (5)



TAheatmap_*.png

PulP output data

- Raw data in 8-bit format (optional)
- Raw .h5 metadata files
- Pulsar data cubes (both from PRESTO and DSPSR pipelines)
- PRESTO pipeline:
 - rfifind mask
 - PSRFITS filterbank data
- DSPSR pipeline:
 - filterbank file(s) when SP analysis was done (optional)
- Single-pulse data (optional)
 - .singlepulse
 - Single-pulse plots
- Diagnostic plots
 - Plot with multiple profiles (multiple TABs, etc.) – *combined.png*
 - DSPSR diagnostic plots – *status.png*
 - Localization maps – *TAheatmap_*.png*

NorthStar: PulP (1)



Applicants Justification Observing Request Target List Additional information

 Specify a new observation

 Specify a new Pipeline :

NorthStar: PulP (1)



The screenshot shows the LOFAR Proposal interface. At the top, there is a banner with the text "LOFAR Proposal" in yellow and the "ASTRON LOFAR" logo. Below the banner, there is a navigation bar with five tabs: "Applicants", "Justification", "Observing Request", "Target List", and "Additional information". The "Observing Request" tab is highlighted with a red box and has the number "1" below it, indicating the first step. Below the tabs, there are two buttons: "Specify a new observation" and "Specify a new Pipeline :". The "Specify a new Pipeline :" button is also highlighted with a red box and has the number "2" below it, indicating the second step.

1

Specify a new observation

2

Specify a new Pipeline :

NorthStar: PulP (2)

Pipeline configuration

Pre processing parameters

Processing mode: **Pre processing only** *

Flagging strategy : **LBA** *

Averaging time factor: [steps] *

Averaging freq. factor: [steps] *

Demixing ? Yes No

Demixing sources : *

- CygA
- CasA
- TauA
- VirA
- HerA
- HydA

Imaging parameters

No imaging selected in processing mode

Subbands per image : [int] *

Field of view : [deg] *



Commit Pipeline

NorthStar: PulP (2)

Pipeline configuration

Pre processing parameters

Processing mode : **Pre processing only** * 3

Flagging strategy : **LBA**

Averaging time factor: [steps] *

Averaging freq. factor: [steps] *

Demixing ? Yes No

Demixing sources : *

- CygA
- CasA
- TauA
- VirA
- HerA
- HydA

Imaging parameters

No imaging selected in processing mode

Subbands per image : [int] *

Field of view : [deg] *



Commit Pipeline

NorthStar: PulP (3)

Pipeline configuration

Pre processing parameters

Processing mode: **Pre processing only** *

Flagging strategy : Pre processing only *

Averaging time factor: Calibration [ps] 1.0

Averaging freq. factor: Calibration + imaging

Demixing ? **Pulsar pipeline** 4

Long baseline calibration

User specified pipeline

CasA

TauA

VirA

HerA

HydA

Imaging parameters

No imaging selected in processing mode

Subbands per image : [int] *

Field of view : [deg] *



Commit Pipeline

NorthStar: Pulp (4)

Pipeline configuration

Pre processing parameters

Processing mode: **Pulsar pipeline** *

Skip RFI check : Yes No

Skip folding : Yes No

Skip pdmp : Yes No

Skip dsspr : Yes No

Skip prepfold : Yes No

Single pulse analysis : Yes No

RRATs analysis : Yes No

Skip dynamic average : Yes No

Length of subintegration (sec) : **0**

Convert HDF5 32-bit raw data to 8-bit : Yes No

Clip threshold (in units of sigma) : **5**

Sigma limit in conversion from raw HDF5 to PSRFITS : **3**

Number of blocks read at once in conversion to PSRFITS : **100**

Prepfold options :

Prepsubband options :

RFIfind options :

*Expert settings: not all options may result in a successful pipeline, therefore the Observatory may decline setting certain parameters.

Dsspr options :

Digifil options :

Prepdata options :

Extra options to convert from raw HDF5 to PSRFITS :

Pulsar :

Configuration comments :

Commit Pipeline

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NorthStar: PulP options (1)

- Skip RFI check
 - all RFI excision commands will be skipped (*rfifind*, *paz*, *clean.py*, *subdyn.py*)
- Skip folding
 - No folding will be done
 - For CV data, nothing will happen unless Single Pulse Analysis = TRUE, or conversion to 8-bit is requested
 - For non-CV data, *prepfold* will not run, only *2bf2fits* to create PSRFITS file and *rfifind*
- Skip pdmp
 - *pdmp* will not run in DSPSR pipeline
- Skip dspsr
 - DSPSR pipeline will not run
- Skip prepfold
 - *prepfold* will not run
- Single pulse analysis
 - Single pulse search part of both PRESTO and DSPSR pipelines will not be carried out
- RRATs analysis
 - RRATs part of the PRESTO pipeline will not be carried out

NorthStar: PulP options (2)

- Skip dynamic average
 - instead «average» should read «spectrum»!
 - *subdyn.py* will not run
- Length of subintegration (sec)
 - Specify sub-integration length for *dpsr* command (that goes with *-L* option)
 - By default it is 5 sec for CV data and 60 sec for non-CV data
 - If *-s* option is specified by a user in Dpsr extra options, then this option is ignored
- Convert HDF5 32-bit raw data to 8-bit
 - Self-explanatory, to run *digitize.py*
- Clip threshold (in units of sigma)
 - Clip threshold for *digitize.py*. Above this threshold raw data will be clipped. Default value is 5.0
- Sigma limit in conversion from raw HDF5 to PSRFITS
 - Same as for converting from raw 32-bit to raw HDF5 8-bit, but rather to PSRFITS data using *2bf2fits* command. Default value is 3
- Number of blocks read at once in conversion to PSRFITS
 - Determines the length of data where the running mean/rms is calculated when converting to PSRFITS using *2bf2fits*. The larger the number, the longer the time window for calculation. Default is 100

NorthStar: PulP options (3)

- Prepfold options
→ *prepfold* extra user options
- Prepsubband options
→ *prepsubband* extra user options
- RFIfind options
→ *rfifind* extra user options
- Dpsr options
→ *dpsr* extra user options
- Digifil options
→ *digifil* extra user options
- Prepdata options
→ *prepdata* extra user options
- Extra options to convert from raw HDF5 to PSRFITS
→ *2bf2fits* extra user options

NorthStar: PulP options (4)

- Pulsar

- Pulsar name for folding *without* «PSR» prefix
- Several (max 3) pulsar names can be specified separated by commas, no spaces
- Folding will be done for the first 3 pulsars
- If no pulsar name is given, then target name from the observation specification will be used.
- User must provide Science Support with the necessary ephemeris files (parfiles)
- If parfiles are not given, PulP will use ephemeris from the ATNF catalog
- If given pulsar is not known by ATNF and parfile is not provided, PulP will search for the brightest pulsar in the FoV of the SAP and fold the data for this brightest pulsar
- Instead of pulsar name, special words can be used, such as:
 - «meta» – use target name for each SAP separately in the HDF5 metadata
 - «parset» - use target name from the observational parset file for each SAP separately (obsolete)
 - «sapfind» – find brightest pulsar in the FoV for each SAP
 - «sapfind3» – find 3 brightest pulsars in the FoV for each SAP
 - «tabfind» – find brightest pulsar in the FoV for each TAB
 - «tabfind+» – look for pulsars first following «meta». If no legitimate pulsar is found, then look for pulsar following logic of «sapfind». If no pulsar found then look for it following logic of «tabfind»
- «NONE» as a pulsar name is ignored

PulP options (advice from the experience)

- For CV data:
 - Dpsr options: `-U minX1 -t 1`
 - Default subint time is 5 s, so for pulsars with P>5 s, consider using larger Length of subintegration, or make subintegrations equal to pulsar period (add option `-s` to Dpsr options)
- For non-CV data:
 - Extra options to convert from raw HDF5 to PSRFITS: `-nsamples 8192`
 - Number of blocks read at once in conversion to PSRFITS: 6
 - RFIfind options: `-blocks 1`
 - For short observations (<~ 3 min), change the default Length of subintegration of 60 s to a smaller number (5 — 10 s)

PSRCHIVE config file

- To process low frequency observations it is very important to have good parameters in your PSRCHIVE configuration file *psrrchive.cfg*
 - Possible locations of this file:
 - `~/.psrrchive.cfg`
 - `echo $PSRCHIVE_CONFIG`
 - If you do not have it, then you can run *psrrchive_config* which will print the contents of configuration to stdout. Redirect the input to the file, and inspect it to have proper values.
 - `psrrchive_config > ~/.psrrchive.cfg`
- What to pay attention to:
 - Dispersion::barycentric_correction. Should be ON
 - `Dispersion::barycentric_correction = 1`
 - FrequencyAppend::weight = none
 - WeightedFrequency::round_to_kHz. Should be OFF (we need maximum precision!)
 - `WeightedFrequency::round_to_kHz = 0`
 - Also possibly consider:
 - `FrequencyAppend::force_new_predictor = 1`
 - `Predictor::default = tempo2`
 - `Predictor::policy = default`



Pulsar visualisation credit: Alessandro Ridolfi