The NANOGrav TOA Generation Process

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http://nanograv.org/
The goal of this document is to describe the process of data reduction and TOA generation used by NANOGrav, including the generation of intermediate data files, in sufficient detail to allow it to be reproduced exactly.

1 Getting the software and data

The data reduction and TOA generation process makes use of several pieces of software. The full list of requirements is:

- **PSRCHIVE** ([http://psrchive.sourceforge.net/](http://psrchive.sourceforge.net/)). This is the main tool used to view and manipulate data in PSRFITS format. It consists primarily of command-line utilities, written in C, but also has a Python interface and a scripting language, `psrsh`. Traditionally, PSRCHIVE has been installed by building it from source (see [http://psrchive.sourceforge.net/installation.shtml](http://psrchive.sourceforge.net/installation.shtml) for full instructions). However, successfully configuring the build tools can be challenging, especially if its dependencies are installed in non-standard locations. A much easier way to install it is using `conda`. Once a conda environment has been set up, PSRCHIVE can be installed simply by running

  `conda install -c conda-forge psrchive`

  This should automatically handle downloading and installing dependencies.

- **psrtools** ([https://github.com/demorest/psrtools](https://github.com/demorest/psrtools)). This is a set of two command-line utilities (autotoa and normalize_rms) that depend on PSRCHIVE. Like PSRCHIVE, they are written in C, and can either be built from source or installed with conda:

  `conda install -c demorest psrtools`

- **nanopipe** ([https://github.com/demorest/nanpipe](https://github.com/demorest/nanpipe)). This is a collection of Python and `psrsh` scripts used to coordinate the various steps of the NANOGrav calibration and TOA generation pipeline. It can be installed by cloning the repository from GitHub:
To install nanopipe, run:

```bash
git clone https://github.com/demorest/nanopipe.git
cd nanopipe
pip install .
```

- **toagen** ([https://gitlab.nanograv.org/nano-time/toagen](https://gitlab.nanograv.org/nano-time/toagen)). This is a special-purpose repository which holds the full directory structure used for the calibration and TOA generation process, including configuration files which are not part of nanopipe. Currently access is limited to NANOGrav members. It is hosted on NANOGrav’s internal GitLab server. If you are a NANOGrav member but have not used GitLab previously, you will have to log in to gitlab.nanograv.org with your NANOGrav username and password and add an SSH key (user icon in upper right > Settings > SSH keys) before you can access the repository. You can then clone it by running:

```bash
git clone git@gitlab.nanograv.org:nano-time/toagen.git
```

toagen is not an installable program or library, but rather contains a set of configuration files and Makefiles that are used in the TOA generation process, along with the current template profiles and .tim files for all NANOGrav pulsars. It does not contain the actual profile .fits files, as these are much too large to be version-controlled (see below).

- Profile data. The working copy of NANOGrav’s profile data is stored on servers at WVU and is accessible from Bowser (bowser.phys.wvu.edu) or the notebook server (notebook.nanograv.org). VEGAS, GUPPI, and PUPPI data are stored under

  `/nanograv/timing/data/`

The data files are grouped into directories by pulsar name (without the “B” or “J” prefix), backend, and year, and are always in a directory called rawdata at the lowest level. For example, the first part of an Arecibo observation of PSR J1713+0747 taken on March 29, 2013 can be found at

```
/nanograv/timing/data/1713+0747/puppi/2013/
rawdata/puppi_56380_1713+0747_0542_0001.fits
```

Calibration scans are found in the same directories, and can easily be identified by the presence of cal in the filenames. Flux calibrator observations are found in directories corresponding to the name of the flux calibrator.¹

¹A quasar, usually B1442+101 = J1445+099, but at Arecibo sometimes J1413+151, and at GBT 3C190 is used for one 2011 observation. There are multiple directories with slight variations on the name, and some with suffixes, but the most recent calibrator observations all seem to go into 1442+101.
Some data from other telescopes used by NANOGrav are also available on the WVU servers, although not in the same place as the GBT and Arecibo data. YUPPI (VLA) data are stored under

/hyrule/data/users/pdemores/VLA/

and are grouped by semester and proposal number. Some CHIME data, grouped by pulsar, are also available, under

/hyrule/data/CHIME/NANOGrav/

As of September 29, 2021, only observations of PSR J1713+0747 from January to September of 2021 are available, but additional data should be copied over soon.

2 Setup and directory structure

The basic processing scripts are contained in nanopipe. As established in the instructions for nanopipe (https://github.com/demorest/nanopipe/blob/master/doc/basic_instructions.txt), the work takes place in a base directory which has subdirectories for each pulsar being processed. Within toagen, there is a separate base directory for each backend (ASP, GASP, PUPPI, GUPPI, YUPPI).

Flux calibrator observations are stored in a separate directory, called fluxcal, inside the base directory for each backend. Before processing the rest of the observations, one needs to create .fcal files from the flux calibrator observations. This is done using the PSRCHIVE tool fluxcal:

```
fluxcal -f -e fcal
```

The results can be checked using the PSRCHIVE calibrator viewer, pacv.

The base directory for each backend also contains several psrsh scripts (copied from nanopipe’s config directory): the basic list is

```
zap_minmax
zap_and_tscrunch
update_be_delay.psrsh
process_fluxcal
img_$be
```

Here $be is the backend name.

There should also be a configuration file called make_psr_make.config.py in the base directory for each backend, containing, at minimum, these lines:

```
ver_id = "$project_id"
tscrunch_arg = "-J ../ zap_and_tscrunch"
```

Here $project_id is a string identifying the particular version of the processing being carried out.
This basic setup is customized in a backend-dependent way in the real NANOGrav TOA generation process, in ways that are captured in the toagen repository. Once the scripts and configuration files are in place, a makefile can be generated for each pulsar by running (from within the corresponding subdirectory)

```make
make_psr_make $basedir > Make.psr
```

For $basedir one should substitute the full path of the base directory. The makefile can be used to produce TOAs by running

```make
make -f Make.psr toas
```

(again from within the pulsar subdirectory). This command can be altered to run several processes in parallel by adding the option `-j $n_procs`, where $n_procs is the number of processes, or by replacing `toas` with another makefile target from this list:

```
rf
calibration
zap
scrunch
templates
toas
```

to only run part of the analysis.

## 3 Per-pulsar makefiles

The main process is laid out in the per-pulsar makefiles (Make.psr), which are generated by the `make_psr_make` script in nanopipe. The processing carried out by `Make.psr` breaks down into the following seven steps, most of which correspond directly to makefile targets:

1. Target `rf`: Combine the raw .fits files into .rf files. The raw data files must be copied in from where they are stored on WVU’s servers. Several corrections are applied at this stage, namely:

   (a) Epoch error due to polyco REF_MJD precision in PSRFITS is corrected. This accounts for the effects of a bug in PSRCHIVE, discovered during the 9-year analysis, which meant that polyco reference MJDs were not stored using enough numerical precision.

   (b) Backend delays (stored in the BE_DELAY PSRFITS header item) are set appropriately. These measure extra signal latency within each backend, usually just digital filter latency, and are determined from the backend design (not measured experimentally). For GUPPI and PUPPI, the delay is a function of the number of channels and channel bandwidth in the observation.
(c) ADC ghost image corrections (Alam et al. 2021, §2.3.1; cf. Kurosawa et al. 2001) are applied.

(d) Par files are installed into the FITS header. This also means realigning the profiles according to the par file being installed. Usually the “predictive” par file from the previous data set is used here, if it is available.

(e) The DM value in the FITS header is set appropriately. This DM value is taken from the par file used above, and is used for de-dispersion prior to frequency averaging.

(f) Specified bands are zapped and time-frequency zapping (Offringa et al., 2010) is applied.

(g) The source and receiver names are fixed to make them uniform across the data set.

If this process fails, the file is noted in the list of cal failures. Finally, the database of calibrator archives is built up.

Commands:

```
psradd -T -j 'fix refmjd' -J
   ../update_be_delay.psrsh -J ../img_$be -j
   'install par $psr.basic.par' -j "e dm=$dm"
   -J ../zap_and_tscrunch_with_list_$tel -j "e
rcvr:name="fix_receiver_name $obsj.fits"
   -j "e name="get_proper_name $obsj.fits" -o
   $obs.rf $obs1.fits $obs2.fits
psradd -T -j 'fix refmjd' -J
   ../update_be_delay.psrsh -J ../img_$be -j
   ../zap_and_tscrunch_with_list_$tel -j "e
rcvr:name="fix_receiver_name $calj.fits"
   -j "e name="get_proper_name $calj.fits" -o
   $cal.cf $cal1.fits $cal2.fits
```

These commands (and the others quoted below) are taken from the Make.psr file, but reformatted to make them independently runnable and as generic as possible. The parsh scripts

```
update_be_delay.psrsh
img_puppi
img_guppi
zap_and_tscrunch_with_list_gb
zap_and_tscrunch_with_list_ao
```

can be found in nanopipe’s config directory (however, the ones found in nanopipe use zap_median, as was done for the 12.5-year and earlier data sets, rather than zap tfzap, used in toagen and the 15-year data set). The Python scripts
fix_receiver_name
get_proper_name

can be found in nanopipe's scripts directory.

2. Target calibration: Calibrate the data (.rf files) to produce .calib files. This performs a very basic flux/polarization calibration (cal gain and phase correction) using pac. Matching with the calibrator observations is done on a per-channel basis, and the pulse phase of the cal transition is set to 0.5 turns. Calibrator Stokes parameters are derived from the fluxcal data. If this process fails, the file is noted in the list of cal failures. Then the data goes through another round of RFI zapping, removing a specified fraction of the band edge, as well as more specified frequency bands, but these seem to mostly be duplicates. Any zapping options in the variable $XPAZ are applied. Finally, if the file $obs.calib does not exist, the file $obs.calibP is removed.

Commands:

```
pac -w
pac -a -j "config SquareWave::transition_phase=0.5" -x -e x.calib -d database.txt $obs.rf
paz -m -E2.0 $zapchannels $(XPAZ) $obs.x.calib
```

$zapchannels is a space-separated list of -F options specifying channels to zap on a per-receiver basis, configured in make_psr_make.

3. Target zap: Apply “min-max” zapping to produce .zap files.

Command:

```
psrsh -e zap ../zap_minmax $obs.x.calib
```

The psrsh script zap_minmax can be found in nanopipe's config directory. (But, just as for zap_and_tscrunch_with_list_$tel above, this is an older version of zap_minmax that uses zap median rather than zap tfzap. The zap tfzap version can be found in toagen.)

4. Target scrunch (1/2): Scale the weights so that the offpulse RMS of the weighted data is 1, producing .norm files.

Command:

```
normalize_rms -w $obs.x.zap
```

5. Target scrunch (2/2): Frequency average to 64 channels, and time average to a number of subintegrations set by the observation length (the length divided by 1800 s, rounded down, plus one), to produce .ff files.

Command:
pam -e ff -f$factor --setnsub `psredit -Q -c length $obs.x.norm | awk '{print int($2/1800.0) + 1}'' $obs.x.norm

$factor$ varies depending on the backend. It is 8 for receivers with 512 channels (GUPPI and PUPPI L-band, PUPPI S-band) and 2 for GUPPI 820 MHz, which has 128 channels.

6. Target templates: Iteratively determine the template and TOAs using autotoa (an implementation of the method outlined in Chapter 2 of Demorest 2007), with a list of all .ff files for this pulsar-frontend-backend combination as input. Use a Gaussian of width 0.1 as the initial guess for the template, and perform at most 3 iterations. Then rotate the final profile by half a turn, and apply UD8 wavelet smoothing to it.

Commands:

```bash
def f in {obs.ff}; do echo f >> rcvr.fflist autotoa -g0.1 -i3 -S rcvr.sum $rcvr.fflist pam -r0.5 -m rcvr.sum psrsmooth -W -t UD8 rcvr.sum
```

7. Target toas: Use pat to calculate the final narrowband TOAs from the .ff files and template and write them to a .tim file in tempo2/IPTA format, adding appropriate flags. Calculate errors using MCMC. Then use the same .ff files to calculate wideband TOAs based on the 12.5-year wideband templates, and write them to a .tim file with a similar format and flags.

Commands:

```bash
echo 'MODE 1' > rcvr.nb.tim pat -A FDM -e err=num -C chan -C subint -C snr -C wt -C flux -C fluxe -f "tempo2 IPTA" -X "-proc 15y -pta NANOGrav -ver TEMP $(XFLAG)" -s rcvr.sum.sm -M $rcvr.fflist >> rcvr.nb.tim pptoas.py --print-flux --quiet --flags=proc,15y,pta,NANOGrav,ver,TEMP$(XFLAG) -m $rcvr.12y.x.avg_port.spl -d $rcvr.fflist -o $rcvr.wb.tim
```

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References


