Robust parameter estimation from pulsar timing data

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ABSTRACT

Recently, global pulsar timing arrays have released results from searching for a nano-Hertz gravitational wave background signal. Although there has not been any definite evidence of the presence of such a signal in residuals of pulsar timing data yet, with more and improved data in future, a statistically significant detection is expected to be made. Stochastic algorithms are often used in a very large parameter space to infer results from data. In this paper, we attempt to rule out effects arising from the stochasticity of the sampler in the inference process. We compare different configurations of nested samplers and the more commonly used Markov chain Monte Carlo method to sample the pulsar timing array parameter space and account for times taken by the different samplers on the same data. Although we obtain consistent results on parameters from different sampling algorithms, we propose two different samplers for robustness checks on data in the future to account for cross-checks between sampling methods as well as realistic run-times.

Key words: gravitational waves – methods: data analysis – pulsars: general

1 INTRODUCTION

Pulsars Timing Arrays (PTAs) (Detweiler 1979; Hellings & Downs 1983; Jenet et al. 2009; Fordman et al. 2010; Hobbs et al. 2010; Manchester et al. 2013) aim to detect the stochastic Gravitational Wave Background (GWB). A GWB signal is likely created by the superposition of gravitational waves emitted by Super Massive Black Hole Binaries (SMBHBs) (Rosado, Sesana & Gair 2015), but there could be other sources such as a relic from inflation (Grishchuk 2005) or cosmic strings (Vilenkin 1981; Vilenkin & Shellard 2000). While increasingly constraining upper limits have been placed on the amplitude of the GWB, there has been no detection of this signature yet. However all operational PTAs are currently detecting a common but spatially uncorrelated red noise process (Arzoumanian et al. 2020; Chen et al. 2021; Goncharov et al. 2021; Antoniadis et al. 2022). This might indicate that the GWB signal will be detected with statistical significance in the near future. In this paper, we look at consistencies between a variety of stochastic samplers used to sample a large parameter space, where the latter is of paramount importance to inferring properties of the GWB signal.

The size of pulsar timing models necessitates the use of a hybrid frequentist and Bayesian analysis, where the pulsar timing model parameters are first obtained using iterative least square fitting with tools such as TEMPO2 (Edwards, Hobbs & Manchester 2006; Hobbs, Edwards & Manchester 2006; Hobbs et al. 2009) or PINT (Luo et al. 2021) to obtain a set of timing residuals. These timing residuals are then modelled to remove excess delays due to red noise processes as well as fluctuations from the variations in the ionized interstellar medium (ISM) codified as dispersion-measure models using Bayesian analysis, while analytically marginalizing over the timing model parameters. This is typically called single pulsar noise analysis (SPNA). Even with this simplification, the
estimation of the red noise and dispersion-measure model parameters remains computationally expensive. Further, in the final stage, when searching for the GWB, all pulsar models must be simultaneously fitted for, along with a model for the correlated signal from the GWB as well as any other correlated or uncorrelated common noise processes. Even when the search is optimized for the smallest number of pulsars, this can lead to final dimensions of the order of hundreds of parameters. Further, for the individual pulsar models as well as the final GWB search, it is desirable to carry out model selection (Raftery 1996; Jeffreys 1998) – a method which is particularly well suited for Bayesian analysis.

This mixed approach implies inherent uncertainties in the comparison of the algorithms themselves as well as any difference in the obtained results. We attempt to address this issue by adapting the most commonly used PTA analysis package, ENTERPRISE (Ellis et al. 2020), to utilize a number of nested sampling algorithms. We perform single pulsar noise analyses for a set of six pulsars, first utilized for the recent limits presented by the European Pulser Timing Array (EPTA) (Chen et al. 2021). Using the most performant nested sampling algorithm as determined from the SPNA analysis, we then search for the GWB using both the pulsar data set as well as the second International Pulser Timing Array (IPTA) second mock data challenge (MDC) (Hazboun, Mingarelli & Lee 2018).

In Section 2, we briefly summarize the data we have used for our inference process. Section 3 serves as an introduction to Bayesian inference with focus on noise models used in this paper and pulsar timing data in general. Some technical details to algorithms we use are also included. We give a summary of our results in Section 4 and conclude in Section 5.

2 DATA

We have used data recently utilized by the EPTA collaboration (Chen et al. 2021) and focused on six pulsars – PSRs J0613-0200, J1012+5307, J1600-3053, J1713 + 0747, J1744-1134, and J1909-3744. The times of arrival (TOAs) of these pulsars are fitted using the TEMPO2 software to pulsar timing models describing the pulsars astrometric, intrinsic, and environmental properties, along with simple polynomial models for the variations of IISM along the line of sight to these pulsars. The resulting ‘ timing residuals’ are shown in Fig. 1, where we highlight the large number of observing systems used for each pulsar data set by different colours. We refer the interested reader to (Chen et al. 2021) and forthcoming EPTA publications for more details on the individual observing systems but list the names here. The abbreviations correspond to the Pulsar Machine (P1, P2, and PuMal/II) instruments at the Westerbork Synthesis Radio Telescope (WSRT), the Reconfigurable Open Architecture Computing Hardware (ROACH) and the Digital Filter Bank (DFB) based devices at the Jodrell Bank Observatory (JBO), the Berkeley-Orleans- Nancy (BON) and the Nancay Ultimate Pulsar Sensitive Instrument (NUPPI) at the Nancay Radio Observatory, and the PSRIX instrument (labelled P217, P200, S110, and asterix) at the Effelsberg radio telescope. The residuals of Fig. 1 encode within them the signatures of contributions from pulsar specific low and high frequency processes as well as common astrophysical signals, such as perturbations due to Solar system bodies (Champion et al. 2010; Caballero et al. 2018), time-variable delays due to density fluctuations in the IISM along the line of sight to the pulsar or the spatially correlated GWB.

In addition to real data, to test different samplers, we have also used simulated timing data, generated by the IPTA collaboration (Verbiest et al. 2016) and used in the second MDC (Hazboun et al. 2018). From the MDC, we choose a data set containing a GWB signal. The data set consists of 33 pulsars, and in addition to the GWB, each individual pulsar is characterized by its own spin noise or red noise. The data also contains white noise characterizing the observing telescopes. The simulated data set spans a timeline of 15 yr and is observed at a central frequency of 1440 MHz. The TOAs are uniformly distributed with observations taken every 30 d.

The extraction of the GWB signal is a complicated process due to the need to transform radio pulsar observations into reference times at which a group of photons from each pulsar in the PTA arrive at Earth or Solar system Barycentre. While the observed data are the TOAs, the data analysis is done on timing residuals. For this the TOAs are first converted into residuals, obtained after subtracting the predicted timing model from the observed TOAs. If the predicted model fits the observations perfectly, the residuals will be identically 0. In addition to the presence of a GWB, additional non-gravitational-wave related noise sources may alter the TOAs, some of these noise models are described in Section 3.

3 BAYESIAN ANALYSIS AND PARAMETER ESTIMATION SETUP

We perform Bayesian inference on the pulsar timing data and sample over parameters corresponding to noise models describing the variations in the residuals as described below. We sample over single pulsars (henceforth, SPNA analysis) as well as the full pulsar timing array (henceforth, PTA analysis) and use different samplers to test the consistency of the inferred noise models. Since nested sampling provides direct access to the marginal likelihood (Buchner 2021a), hypothesis-testing may be done naturally from a nested sampling analysis, and PTA analysis could in turn quantify the support for the imprint of the quadrupolar versus non-quadrupolar correlations on data. However, analyses such as those from the Parkes PTA...
(PPTA) or NANOGRAV typically utilize the Parallel Tempering Markov Chain Monte Carlo (PTMCMC) (Ellis & van Haasteren 2017) method due to the lower computational cost. Since PTMCMC does provide direct access to the marginal likelihoods, methods such as the Savage–Dickey approximation and hypermodel sampling are employed for model comparison. Even though the EPTA and IPTA results have been presented in the literature, which utilize efficient multi-ellipsoidal nested sampling algorithms such as MULTINEST and POLYCHORD for SPNA, the final search for the GWB still utilizes PTMCMC or similar Markov Chain Monte Carlo (MCMC) based methods.

3.1 Bayesian inference

We provide a very brief summary of Bayesian inference to make our study self-contained. We point the reader to detailed resources like (Gregory 2005; Sivia & Skilling 2006) for further reading. Bayesian analysis estimates parameters from probability distribution functions (PDFs). The posterior PDF is obtained by providing the initial prior PDF and using that in combination with the likelihood, containing information about the data. The Bayes’ theorem can be written down as

\[
P(\hat{\theta}|d) = \frac{P(d|\hat{\theta}) P(\hat{\theta})}{P(d)},
\]

where \(\hat{\theta}\) refers to a multidimensional parameter set, \(d\) is the data, and the notation \(P(\hat{\theta}|d)\) refers to information on \(\theta\) given \(d\). Details of the likelihood calculation in case of analysis of pulsar timing data may be found in (Arzoumanian et al. 2015) and references therein. In addition to estimating parameters by using prior knowledge as well as knowledge from observed data, Bayesian analysis allows us to perform model selection. With the data remaining the same, this means performing an analysis each time with a different model. In that case, equation (1) may be rewritten as

\[
P(H|d) = \frac{P(d|H) P(H)}{P(d)},
\]

\(H\) represents a hypothesis, and in case of pulsar analysis, \(H\) may be assuming that the timing data contains a GW signal, \(H_{\text{GWB}}\) or only a common red noise signal \(H_{\text{CRN}}\), but not a GWB. From equation (2), if we then compute the ratio of probabilities \(P(H_{\text{GWB}})\) and \(P(H_{\text{CRN}})\), we get a quantitative measure of which model is more preferred by the data.

3.2 Noise models

When analysing single pulsar data, we focus only on individual pulsars’ intrinsic red noise (RN), the noise from dispersion measure arising from the interstellar medium (DM), and white noise (WN), inherent to the observing telescopes. In addition, when sampling over the parameter space of a PTA analysis, we include a CRN. When the common process includes spatial correlations, we search for a common GWB. We briefly describe each of the noise processes below:

3.2.1 White noise

The white noise itself can be divided into two parts: (i) a multiplicative factor of the estimated error bar on the observed TOAs, the EFAC, and (ii) an additional noise adding in quadrature to the error bars, the EQUIAD. Both these components vary across the different observing telescopes even if they all observe the same pulsar. The total error on a TOA, \(\sigma\) can be written as

\[
\sigma = \sqrt{(\sigma_{\text{EFAC}})^2 + \text{EQUIAD}^2}.
\]

EFAC represents possible uncertainty on the TOA error estimation during the cross-correlation of the pulsar profile with the standard template (Taylor 1992), and EQUIAD may arise from physical effects like pulsar jitter and give rise to additional scatter of the TOAs (EKERS & Moffet 1968).

3.2.2 Red noise

Red noise is intrinsic to each pulsar, and also commonly referred to as spin-noise. This arises primarily as a result of irregularities in pulsar-spin (Cordes & Downs 1985; D’Alessandro et al. 1995). The imprint on the pulsar residuals from the intrinsic noise is also a red process, like the GWB, and the power spectrum may be described as a power-law

\[
\phi_{\text{RN}} = \frac{A_{\text{RN}}^2}{12\pi^2} \left(\frac{1}{1 \text{ yr}}\right)^{-3} f^{-\gamma_{\text{RN}}} T,
\]

where \(A_{\text{RN}}\) and \(\gamma_{\text{RN}}\) are the amplitude and spectral index of the red noise process respectively, and \(T\) is the total timespan between latest and earliest TOA.

3.2.3 Dispersion measure noise

As the pulses from a pulsar travel through the interstellar medium, the imprint of the interstellar medium is also encoded on the TOAs. Dispersion measure is time-varying and defined as the integrated column density of free electrons in the pulsar’s line of sight (You et al. 2007). Unlike intrinsic red noise, this noise is frequency-dependent and follows a \(v^{-2}\) dependence, \(v\) being the radio frequency. This source may be further described by an additional power-law spectrum of the form

\[
\phi_{\text{DM}} = A_{\text{DM}}^2 \left(\frac{1}{1 \text{ yr}}\right)^{-3} f^{-\gamma_{\text{DM}}} T \left(\frac{1400 \text{ MHz}}{v}\right)^2,
\]

where \(A_{\text{DM}}\) and \(\gamma_{\text{DM}}\) are the amplitude and spectral index of the dispersion noise, respectively.

3.3 Samplers

As described above, the parameter space of even a single pulsar is multidimensional, and we use techniques of stochastic sampling to infer the noise properties of pulsars. To compare among different samplers, we use nested sampling (Skilling 2006) as well as MCMC methods, where the latter is also conventionally used in inference from pulsar timing data (Ellis & van Haasteren 2017). We have made use of the modular nature of the analysis code ENTERPRISE and incorporated different nested samplers to be used with the Likelihood function available within the code. We also use the native PTMCMC sampler, both with and without message-passing-interface (MPI) (Message Passing Interface Forum 2021), making a thorough study of performances from different kinds of samplers. We briefly describe the individual samplers used in this paper below.

3.3.1 PTMCMC

MCMC (Raftery 1996; Gamerman & Lopes 2006) is one of the commonest methods to stochastically sample a parameter space.
Furthermore, Parallel Tempering (Swendsen & Wang 1986; Geyer 1991) is incorporated to explore the parameter space at different temperatures, thereby enabling a denser sampling. PTMCMC is natively used in the pulsar timing software ENTERPRISE. MCMC directly samples the posterior distribution and after the initial stage, called burn-in, gathers samples which are the representative posterior samples. In this paper, we have used in addition to PTMCMC, also its MPI-enabled version (henceforth, PTMCMC-MPI), and we notice a speedup of around a factor of two in most cases when run using the same machine. Details of the number of cores used are given at the end of Section 4.1.

3.3.2 PyMultiNest

Conventionally, the nested sampling method samples the prior by distributing live points and exploring the parameter space by finding higher regions of likelihood. Each live point forms a contour on the likelihood surface which gets updated as live points corresponding to lower likelihood values get replaced by ones associated with higher likelihood values. Ref. (Feroz, Hobson & Bridges 2009) updated this method by forming regions on the likelihood surface and associating them to multiple multidimensional ellipsoids. Furthermore, this has been made more user-friendly by introducing a Python interface in (Buchner et al. 2014) called PYMULTINEST. In this paper, we use the parallelized version of the same by interfacing it with the MPI protocol.

3.3.3 Dynasty

The nested sampling method described above is known as Static Nested Sampling. In addition, the Dynasty sampler (Speagle 2020) also includes Dynamic Nested Sampling. Throughout our paper, we have however used the Static sampler from within DYNASTY. The configuration we have when using DYNASTY relies on constructing the ellipsoids as implemented in MULTINEST and as such differs only in the use of the parallelization through MPI as we now parallelize sampling the prior. In addition, the decision of when to construct multiple bounds differs in Dynasty as opposed to MULTINEST. Further information may be found also in the documentation.1 Our implementation follows the call to DYNASTY as in (Smith et al. 2020), and we have adopted the approach of parallelization as in the publicly available PBILBY code.

3.3.4 UltraNest

ULTRANEST (Buchner 2021b) is a newly introduced nested sampling algorithm. It is designed to ensure accurate sampling of the parameter space, especially in the cases of widely separated minima or tightly correlated parameter density distribution for which multi-ellipsoidal algorithms such as MULTINEST have been shown to fail. ULTRANEST utilizes the Radfriends (Buchner 2016) algorithm along with flexible penalization schemes which are dynamically reconfigured to allow resampling of previously sampled regions. We have utilized the Reactive nested sampling algorithm of ULTRANEST for our test, as we found no discernible benefits from using the static version in our initial testing. It should be noted that the hybrid frequentist and Bayesian approach of standard PTA analysis means this article presents a restrictive comparison for

| Parameter | Prior range |
|-----------|-------------|
| log(AKN)  | [−18, −10]  |
| γRN       | [0, 0.7]     |
| log(ADM)  | [−18, −10]  |
| γDM       | [0, 0.7]     |
| log(AGWB) | [−18, −10]  |
| γGWB      | [0, 0.7]     |

ULTRANEST as this algorithm is expected to perform better with very large numbers of model parameters.

4 RESULTS

We present the results from the SPNA analyses on the 6 pulsars from the EPTA and present results from the full PTA analyses from the simulated data set. For the SPNA analyses, we have recorded the time taken by each sampler. For the PTA analysis, we focus only on the fastest of the nested samplers and use PTMCMC for comparison between two types of samplers. The intrinsic parameters being sampled over and their respective prior ranges are given in Table 1.

4.1 SPNA

We present results only on RN and DM as these directly affect the TOAs and are intrinsic to the pulsars. Fig. 2 shows the amplitude and spectral index of the RN noise models inferred from the respective 6 EPTA pulsars. The models are presented in the form of posterior PDFs. Fig. 3 shows the same for the DM noise models. In each case, we show results obtained from different samplers; we show results from PTMCMC, PTMCMC-MPI, PYMULTINEST, DYNASTY, and ULTRANEST. In case of the nested samplers, we have used 4096 live points and have used $2 \times 10^9$ posterior samples for the PTMCMC-based runs for each pulsar. It is, however, to be noted that the final number of posterior samples for the nested samplers is much larger than the number of live points because of the scaling from parallelization and is comparable to the MCMC runs. Since we have used the samplers each time in combination with the generic package ENTERPRISE, the likelihood model therefore remains the same and the results show the robustness of the sampling as the PDFs are very consistent with each other. In Tables 2 and 3, we show the medians of the RN and DM parameters, respectively, by quoting the median values and the 5 per cent and 95 per cent quantiles obtained from the PDFs in Figs 2 and 3. The values, along with the widths of the credible intervals show the consistency of the results obtained from each sampler.

We quantify the differences in PDFs by giving the values of Kolmogorov–Smirnov (KS) statistic (Kolmogorov 1933; Smirnov 1948) in Table 4. If the cumulative distributions corresponding to two posterior distributions $P_1(x)$ and $P_2(x)$ are $P_1(x)$ and $P_2(x)$, respectively, the KS statistic is the largest difference:

$$KS = \sup_x |P_1(x) - P_2(x)|,$$

where $\sup_x$ is the supremum function, defined as the smallest element greater than or equal to all numbers of both distributions. From the above definition, the KS value always lies between 0 and 1. If we find the values to be closer to 0, we may consider the underlying PDFs $P_1(x)$ and $P_2(x)$ to be very close to each other. A source of differences

1https://dynesty.readthedocs.io/en/latest/faq.html
in PDFs is however the stochasticity of the algorithm itself; this is in
addition to inherent differences between the PDFs being compared.

To quantify for this and establish a threshold from the stochasticity
itself, for each parameter of each pulsar and for each sampler, we
generated 20 sets of resampled posterior samples and computed the
KS statistic values between all combinations of these 20 data sets.
The maximum KS statistic arising from this study for all parameters
and sampler for each pulsar is always $\sim 10^{-3}$, the highest values
overall being for J1744-1134, and the log $\Lambda_{\text{SS}}$ parameter, $\sim 0.007$.
So, from Table 4, we take values $> 10^{-3}$ to signify a difference in
the PDF arising inherently. From Table 4, the highest KS value is $\sim 0.3$
between PTMCMC and PYMULTINEST for J0613-0200 as well as
J1744-1134.

Based only on these values, the PDFs are not quantifiably close
to one another. The results of the PDFs quantified in Tables 2 and
3, however do confirm that the PDFs give consistent results. This
apparent discrepancy is due to the nature of the KS statistic and
its inherent dependence on the number of final posterior samples,
$\sim 10^5$ in our case. As the number of samples increase, the statistic
becomes more and more sensitive to inherent differences in sampling.

Having obtained consistent results, however, we may conclude that
the visual differences in the PDFs are likely due to a combination
of the stochasticity of the algorithms and the difference in
the individual samplers. We notice PTMCMC combined with MPI
gives a speedup in all cases, and while that is a significant gain
in runtimes, we note that the algorithm, when coupled with MPI,
is different from the native PTMCMC. PTMCMC, when used in
a single core, does not do parallel tempering (the name is a misnomer
in this case). It is only when coupled with MPI, that there is
a single temperature per thread and the parallel tempering kicks
in. Moreover, from Figs 2 and 3, we note that PTMCMC-MPI results
are closer to those obtained with parallel nested samplers. This
is likely because the multiple chains running with different
temperatures in case of PTMCMC-MPI, allow a more exhaustive
exploration of the parameter space, making the final posterior PDFs
closer to those obtained using nested samplers. We also note that,
among nested samplers, ULTRANEST and DYNesty show excellent agreement, whereas PYMULTINEST distributions tend
to be slightly different. While all the nested samplers that we have
used in this work rely on the underlying algorithm MULTINEST,
there are subtle differences among PYMULTINEST, DYNesty,
and ULTRANEST. ULTRANEST and DYNesty have slight
improvements over MULTINEST (and therefore PYMULTINEST)
and our results suggest that MULTINEST itself is probably not
good enough to sample the complex and high dimensional parameter
space of the pulsars except in the simplest cases. It may be worth
trying to do PYMULTINEST analyses with finer settings, however
that would not be a one-to-one comparison among samplers as is
our goal here. In this analysis, we have only changed the sampler. A
robust check of the likelihood function would be to keep the sampler
the same and change the likelihood definition. The existing software
TEMPOSEST (Lentati et al. 2014) is independent of ENTERPRISE
and defines the likelihood function independently. It inherently uses
the MULTINEST sampler, a check of the likelihood function may be
to repeat an analysis with TEMPOSEST and ENTERPRISE coupled
with PYMULTINEST. This will be studied in a future publication.

Finally, in Table 5 we note the walltime in hours taken by
each sampler on the same data. We also note the number of
dimensions, $\text{N}_{\text{dim}}$, and number of TOAs, $\text{N}_{\text{TOA}}$, for each pulsar.

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2We thank Michael Keith for pointing this out to us.
Table 2. Red noise parameters obtained from PDFs shown in Fig. 2. The values shown are the medians, the subscript, and superscript values indicate the 5% and 95% quantiles obtained from the distributions of the individual PDFs.

| Pulsar      | Parameter | PTMCMMC | PTMCMMC-MPI | PyMultiNest | Dynasty | UltraNest |
|-------------|-----------|---------|-------------|-------------|---------|-----------|
| J0613-0200  | γRN       | 5.2±6.2 | 4.90±7.2    | 4.87±7.3    | 4.98±6.7 | 4.99±6.7  |
|             | log AγRN  | −15.02  | −14.81      | −14.79      | −14.86   | −14.86    |
| J1909-3744  | γRN       | 3.26±7.4| 3.29±8.1    | 3.67±4.3    | 3.32±2.2 | 3.26±7.2  |
|             | log AγRN  | −14.06  | −14.08      | −14.11      | −14.10   | −14.06    |
| J1600-3053  | γRN       | 2.48±7.3| 2.64±1.5    | 2.92±0.9    | 2.60±1.1 | 2.63±4.0  |
|             | log AγRN  | −13.66  | −13.70      | −13.75      | −13.69   | −13.71    |
| J1012 + 5307| γRN      | 1.40±7.3| 1.53±1.0    | −          | 1.53±1.1 | −         |
|             | log AγRN  | −13.09  | −13.09      | −13.23      | −13.09   | −13.22    |
| J1713 + 0747| γRN      | 3.48±7.3| 3.55±2.0    | −          | 3.54±2.0 | 3.54±6.3  |
|             | log AγRN  | −14.25  | −14.27      | −14.71      | −14.27   | −14.71    |
| J1744-1134  | γRN       | 0.84±4.2| 1.04±1.1    | 1.65±2.2    | 1.12±1.3 | −         |
|             | log AγRN  | −13.41  | −13.44      | −13.84      | −13.45   | −13.77    |

Table 3. Dispersion measure parameters obtained from PDFs shown in Fig. 3. The values shown are the medians, the subscript, and superscript values indicate the 5% and 95% quantiles obtained from the distributions of the individual PDFs.

| Pulsar      | Parameter | PTMCMMC | PTMCMMC-MPI | PyMultiNest | Dynasty | UltraNest |
|-------------|-----------|---------|-------------|-------------|---------|-----------|
| J0613-0200  | γDM      | 2.62±6.6| 2.80±4.2    | 3.06±0.9    | 2.80±0.8 | 2.81±3.5  |
|             | log AγDM  | −13.66  | −13.73      | −13.83      | −13.73   | −13.73    |
| J1909-3744  | γDM      | 2.79±6.1| 2.79±6.6    | 3.55±6.9    | 2.76±6.6 | 2.83±6.9  |
|             | log AγDM  | −13.89  | −13.89      | −14.21      | −13.87   | −13.91    |
| J1600-3053  | γDM      | 2.88±6.2| 2.87±7.9    | 3.00±2.2    | 2.87±2.4 | 2.87±2.7  |
|             | log AγDM  | −13.30  | −13.30      | −13.31      | −13.31   | −13.30    |
| J1012 + 5307| γDM      | 2.00±6.4| 2.01±8.1    | −          | 1.96±6.3 | −         |
|             | log AγDM  | −13.64  | −13.64      | −13.64      | −13.62   | −13.45    |
| J1713 + 0747| γDM      | 1.10±6.4| 1.04±6.7    | −          | 1.05±6.4 | 1.04±6.8  |
|             | log AγDM  | −13.37  | −13.36      | −13.36      | −13.36   | −13.36    |
| J1744-1134  | γDM      | 4.14±8.3| 3.97±0.5    | 3.84±1.15   | 3.89±9.5 | −         |
|             | log AγDM  | −14.41  | −14.33      | −14.24      | −14.29   | −14.28    |

Table 4. Results on KS statistics on 6 pulsars from SPNA results from different samplers. The runs which ended up being unfinished after months do not have KS statistics’ values associated with them and are given as ‘−’. The values are shown differently for the red noise and dispersion measure models, and for the two parameters, the amplitude, and spectral index separately. The p-value in all cases is however, <10^-5, this is likely due to the large number of posterior samples we compare, ~10^7 for each PDF.

| Pulsar      | Parameter | PTMCMMC versus PTMCMMC-MPI | PTMCMMC versus PyMultiNest | PTMCMMC versus Dynasty | PTMCMMC versus UltraNest |
|-------------|-----------|---------------------------|---------------------------|------------------------|-------------------------|
|             |           | Red Noise | DM Noise | Red Noise | DM Noise | Red Noise | DM Noise | Red Noise | DM Noise | Red Noise | DM Noise |
| J0613-0200  | γ         | 0.154     | 0.141    | 0.172     | 0.291     | 0.119     | 0.13     | 0.132     | 0.145    |
|             | log A     | 0.151     | 0.137    | 0.173     | 0.291     | 0.116     | 0.127    | 0.129     | 0.137    |
| J1909-3744  | γ         | 0.01      | 0.009    | 0.094     | 0.117     | 0.023     | 0.041    | 0.012     | 0.016    |
|             | log A     | 0.009     | 0.006    | 0.074     | 0.120     | 0.020     | 0.043    | 0.012     | 0.018    |
| J1600-3053  | γ         | 0.079     | 0.038    | 0.196     | 0.185     | 0.062     | 0.035    | 0.086     | 0.037    |
|             | log A     | 0.074     | 0.03    | 0.251     | 0.119     | 0.055     | 0.017    | 0.08     | 0.026    |
| J1012 + 5307| γ         | 0.042     | 0.042    | −         | −         | 0.042     | 0.05    | −         | −        |
|             | log A     | 0.017     | 0.037    | −         | −         | 0.034     | 0.042    | −         | −        |
| J1713 + 0747| γ         | 0.073     | 0.095    | −         | −         | 0.062     | 0.077    | 0.072     | 0.088    |
|             | log A     | 0.073     | 0.057    | −         | −         | 0.064     | 0.036    | 0.072     | 0.049    |
| J1744-1134  | γ         | 0.129     | 0.114    | 0.291     | 0.153     | 0.154     | 0.154    | −         | −        |
|             | log A     | 0.113     | 0.112    | 0.295     | 0.166     | 0.136     | 0.150    | −         | −        |
Table 5. Walltime in hours for the SPNA runs with each sampler, for each pulsar the number of dimensions and number of TOAs are also given as Ndim and NTOAs, respectively. The unfinished runs’ times are shown as ‘-‘. From this table, we note that only PTMCMC and DYNESTY are expected to finish within a feasible time-scale. Furthermore, when used with MPI, runtimes with PTMCMC can be scaled up. For some pulsars, the speedup obtained is up to a factor of 2.

| Pulsar Name | Ndim | NTOAs | PTMCMC | PTMCMC-MPI | Sampler        | Dynasty | UltraNest |
|-------------|------|-------|--------|------------|---------------|---------|-----------|
| J0613-0200  | 50   | 3022  | 9.91   | 8.82       | 745.69        | 6.83    | 42.25     |
| J1909-3744  | 18   | 2817  | 13.61  | 5.60       | 3.40          | 1.68    | 2.11      |
| J1600-3053  | 30   | 3345  | 19.06  | 7.16       | 16.27         | 3.5     | 220       |
| J1012 + 5307| 56   | 5837  | 28.80  | 13.94      | -             | 11.92   | -         |
| J1713 + 0747| 58   | 5052  | 29.11  | 14.82      | -             | 15.95   | 141.32    |
| J1744-1134  | 38   | 1980  | 19.66  | 6.49       | 321           | 4.5     | -         |

We have used the same machine for all SPNA runs to have a fair comparison of walltimes. A reason for DYNESTY’s speedup is also the parallelization of the prior-sampling, as mentioned in (Smith et al. 2020). Specifically, for the pulsars J1713+0747 and J1012 + 5307, we were unable to get the sampler to converge after these runs took at least 80 d and we do not present their results. From Table 5, we note that PYMULTINEST becomes unusable for most pulsars; while this is mostly likely due to the increased dimensionality; indeed the missing pulsars for PyMultiNest have Ndim = 56 and Ndim = 58, this is likely also a combination of the high Ndim and the large NTOAs.

In Figs 2 and 3, all nested samplers and PTMCMC-MPI were run in parallel using 47 cores of a single CPU, where each CPU has the specification of Intel(R) Xeon(R) Gold 6252N CPU @ 2.30GHz 35.75 MB with 192 GB memory and 48 cores in total. PTMCMC in itself was run using a single core.

4.2 PTA

In this section, we choose the two fastest samplers from Table 5. In addition to being the fastest, we also use one nested sampler (DYNESTY) and one MCMC sampler (PTMCMC) for consistency checks between two different methods of sampling. Since we do an analysis on all 33 pulsars together which form the pulsar timing array in the IPTA-MDC2, we fix the WN parameters to their TEMP02 fit values to make the analysis computationally feasible. We analyse simulated data where a GWB has been injected as mentioned in Section 2. The injected value of the GWB amplitude is picked up by the resulting PDFs of the GWB amplitude by both samplers as shown in Fig. 4. The figure shows the results when the analysis is done by keeping the spectral index of the GWB power spectrum fixed to 4.33. In addition, we repeated the analysis by varying both the spectral index and the GWB amplitude. This is shown in Fig. 5. The upper panel shows the results of the GWB amplitude when γ is varied and kept fixed. In the lower panel, the amplitude for the varying γ case is plotted by choosing the amplitude values corresponding to those of γ lying between 4.3 and 4.4 and the resulting PDF looks very similar to the case when γ is kept fixed to 4.33.

In addition, we perform model selection between a GWB and a CRN, using both samplers. With PTMCMC, we use the hypermodel approach, available within ENTERPRISE to extract a Bayes’ factor in favour of one of the two models. The model selection remains inconclusive from the values of Bayes’ factors obtained with either sampler. Recently, (Chalumeau et al. 2021) also used these two samplers to get model selection results. The values obtained from both samplers are given in Table 6. This shows that we are unable to assign a model to the data even when the data is ideally simulated, contains no DM noise, and contains a GWB signal of considerable amplitude ADGBW = 1.3 × 10^{-15}. This problem of model selection will therefore become even more important in real data which will additionally contain unmodelled noise. Further, we note the uncertainties in Table 6 and the slightly higher uncertainty values associated with the DYNESTY runs. While for hypermodel sampling, one run suffices to assign a value of Bayes’ factor to a model, with the nested sampling approach, we have to resort to separate runs for each model to get a Bayes’ factor. The error is therefore added in quadrature and adds up in the case of the runs done with DYNESTY. In Section 5, we suggest a method to be able to assign a threshold value of Bayes’ factor to claim a detection from real data.

5 CONCLUSIONS

We have compared different algorithms to sample the parameter space of the pulsar likelihood. We have used samplers to infer models from six single pulsars as well as a PTA comprised of thirty three pulsars. In each case, we note generally good qualitative agreement between different sampling algorithms and from estimates of runtime as well as to maintain a balance between different ways of
Figure 5. Comparison between varying and fixing the spectral index $\gamma$. The top panel shows the 1D posterior PDFs of the log of the GWB amplitude, $\log A_{\text{GWB}}$. The three posteriors correspond to when the spectral index, $\gamma$ is varying in the sampling (black), when the spectral index is fixed to 1/3 in the run (dashed, red) and when the $\log A_{\text{GWB}}$ is restricted to the indices of $\gamma$ corresponding to a narrow range of $[4.3, 4.4]$ (blue) in the varying-$\gamma$ run. The injected value of $\log A_{\text{GWB}} = -14.89$ is shown as a green vertical line. The bottom panel shows the two-dimensional plot of $\gamma$ versus $\log A_{\text{GWB}}$ when $\gamma$ is being varied in the inference run; the $\gamma$ range of $[4.3, 4.4]$ corresponding to the blue PDF in the upper panel is shown in blue horizontal lines. We show only the results from DYNESTY here as Fig. 4 already shows good agreement between PTMCMC and DYNESTY.

Table 6. Model selection results: Bayes’ factors between models GWB and CRN compared with samplers PTMCMC and DYNESTY.

| Model      | $B^\text{GWBB}_{\text{CRN}}$ | $B^\text{GWBB}_{\text{Fix } \gamma}$ |
|------------|-------------------------------|--------------------------------------|
| Dynasty    | 0.534 ± 1.148                  | 0.945 ± 1.147                        |
| PTMCMC     | 1.279 ± 0.018                  | 0.955 ± 0.011                        |

sampling, propose the use of PTMCMC and DYNESTY as preferred methods for future inference from pulsar timing data.

In future, we will generate data with randomized sky positions for the pulsars, commonly referred to as sky-scrambling (Taylor et al. 2017), to resemble a realistic realization of measured data and will construct a distribution of Bayes’ factors to build a ‘background’. Furthermore, we will inject GWB signals and infer their properties and compare the resulting Bayes’ factors distribution. This will likely give us an idea of the threshold Bayes’ factor to claim a detection of a GWB, if present in data for those number of pulsars that we search over and hence their specific locations in the sky. In addition, this will also be a function of the overall timing precision and stability, meaning we have to repeat this experiment when adding new pulsars or for new data sets with upgraded instruments. This work is in progress and will be published separately.

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DATA AVAILABILITY

The paper has made use of data on 6 single pulsars, results from whose analyses have been published by the EPTA in (Chen et al. 2021) and MDC data simulated by the IPTA, available in https://github.com/ipta/mdc2/.

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