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# To infinity and beyond: Efficient computation of ARCH(\infty) models 

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# To infinity and beyond: Efficient computation of $\operatorname{ARCH}(\infty)$ models* 

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#### Abstract

This paper provides an exact algorithm for efficient computation of the time series of conditional variances, and hence the likelihood function, of models that have an $\operatorname{ARCH}(\infty)$ representation. This class of models includes, e.g., the fractionally integrated generalized autoregressive conditional heteroskedasticity (FIGARCH) model. Our algorithm is a variation of the fast fractional difference algorithm of Jensen and Nielsen (2014). It takes advantage of the fast Fourier transform (FFT) to achieve an order of magnitude improvement in computational speed. The efficiency of the algorithm allows estimation (and simulation/bootstrapping) of $\mathrm{ARCH}(\infty)$ models, even with very large data sets and without the truncation of the filter commonly applied in the literature. In Monte Carlo simulations, we show that the elimination of the truncation of the filter reduces the bias of the quasi-maximum-likelihood estimators and improves out-of-sample forecasting. Our results are illustrated in two empirical examples.


JEL codes: C22, C58, C63, C87.
Keywords: Circular convolution theorem, conditional heteroskedasticity, fast Fourier transform, FIGARCH, truncation.

## 1 Introduction

Many autoregressive conditional heteroskedasticity (ARCH) models have a so-called $\mathrm{ARCH}(\infty)$ representation, which is similar to the linear representation for time series models for the conditional mean. The $\mathrm{ARCH}(\infty)$ representation, or model, appears to be introduced by Bollerslev (1986). An example is the very popular fractionally integrated GARCH (FIGARCH) model of Baillie, Bollerslev, and Mikkelsen (1996); see below for additional examples.

[^0]In $\operatorname{ARCH}(\infty)$ models, the sequence of conditional variances is a linear convolution of all previous squared innovations, where the weights are simple functions of the parameters of the model. By standard methods, the calculation of the sequence of conditional variances, and hence of the likelihood function, requires $O\left(T^{2}\right)$ arithmetic operations. Evaluating the likelihood function of an $\operatorname{ARCH}(\infty)$ model requires calculation of the sequence of conditional variances, and optimizing the likelihood typically requires many iterations and hence many calculations of the sequence of conditional variances with different parameter values. Thus, for large sample sizes, the $O\left(T^{2}\right)$ computational cost can be prohibitive for estimation of the model, and even more so for multivariate models. Even if it is not prohibitive for estimation with a given sample, it may render bootstrap inference or simulation methods infeasible.

Sample sizes commonly applied in empirical work have grown tremendously in recent years, especially in finance. Time series of daily observations over many years are very common; for example, our daily exchange rate data in Section 5.2 has 12,300 observations. However, much longer data series are sometimes analyzed. For example, in recent empirical work, models of the $\operatorname{ARCH}(\infty)$ class have been fitted to samples of size 100,032 (Han, 2008), 509,472 (Chortareas, Jiang, and Nankervis, 2011), 55,860 (Conrad, Rittler, and Rotfuß, 2012), and 156,672 (Naeema, Shahbazb, Saleemc, and Mustafaa, 2019).

To speed up computation it has been standard in the literature, at least since Baillie et al. (1996), to truncate the convolution at a fixed truncation number such as 1,000 . That is, only a part of the time series of innovations is used to calculate the sequence of conditional variances. Of course, this is an approximation which implies that the calculation is not exact.

In this paper, we discuss efficient computation of the sequence of conditional variances, and hence the likelihood function, for any model that has an $\operatorname{ARCH}(\infty)$ representation. We make three separate contributions.

First, we show how to apply the fast Fourier transform (FFT) of Cooley and Tukey (1965) and the circular convolution theorem to reduce the required number of arithmetic operations from $O\left(T^{2}\right)$ to $O(T \log T)$. Our proposed algorithm is a variation of an algorithm recently proposed by Jensen and Nielsen (2014) for the calculation of fractional differences. The variation is threefold. We (i) apply the FFT-based method to a transformation of a time series instead of the time series itself, (ii) account for possible truncation of the convolution, and (iii) apply arbitrary coefficients instead of the fractional difference coefficients. The algorithm is exact and does not rely on any approximation device. The increase in computational speed relative to linear convolution can easily be a factor of 10 or even much more for sample sizes that can be encountered in practical applications, and is sufficient to make bootstrap and simulation methods for $\operatorname{ARCH}(\infty)$ models feasible, even with very large sample sizes.

Second, in a small Monte Carlo simulation, we quantify the estimation bias introduced by using a fixed truncation number. We show that, for the truncation number typically applied in the literature, 1,000 , the bias can be substantial. However, because our algorithm is exact and sufficiently fast to eliminate the need for truncation, it does not suffer from this bias.

Third, in the Monte Carlo simulation, we also compare out-of-sample forecast performance with and without truncation. We find that the bias from truncation makes forecast less precise compared with no truncation. Moreover, the difference in forecast errors with and without truncation in fact increases with the sample size.

In two empirical applications we apply our algorithm to data sets of exchange rates. These highlight the differences obtained with and without the truncation number, and in
particular emphasize the very different computational time achieved with the new FFTbased algorithm compared with standard linear convolution.

Recently, Klein and Walther (2017) also proposed an application of the FFT and circular convolution theorem, based on Jensen and Nielsen (2014), to calculate the sequence of conditional variances for FIGARCH models. Their method and algorithm is closely related to ours. However, Klein and Walther (2017) maintain the fixed truncation number throughout and do not consider exact algorithms. They also do not consider the bias associated with the application of a fixed truncation number. We include some comparisons with their algorithm, which show that, in the absence of the fixed truncation number, its computational time is of the same order of magnitude as the standard linear convolution algorithm.

Many conditional heteroskedasticity models proposed in the literature have an $\mathrm{ARCH}(\infty)$ representation, going back to the GARCH model of Bollerslev (1986). However, the conditional variance of the (finite-order) GARCH model is a sum of only a fixed number of terms (in the same way that an ARMA model is a sum of a fixed number of terms, but it has an $\mathrm{MA}(\infty)$ representation). Because the GARCH model is a sum of only a fixed number of terms, it will not benefit noticeably from our method. In contrast, our method will be of great benefit to models where the conditional variance is not a sum of a fixed number of terms. In particular, this covers all the long-memory-type conditional variance models in the $\operatorname{ARCH}(\infty)$ class, e.g., the FIGARCH model of Baillie et al. (1996), the long-memory GARCH model of Karanasos, Psaradakis, and Sola (2004), and the hyperbolic GARCH model of Davidson (2004). Finally, our method also applies to some models that are not even in the $\mathrm{ARCH}(\infty)$ class. For example, it applies to the fractionally integrated asymmetric power ARCH model of Tse (1998) and the linear ARCH model of Robinson (1991) and Giraitis, Robinson, and Surgailis (2000), and it partly applies to the fractionally integrated exponential GARCH model of Bollerslev and Mikkelsen (1996); see Section 2.2 for details.

The next section defines the $\mathrm{ARCH}(\infty)$ class and explains our proposed FFT-based algorithm for calculation of the sequence of conditional variances. Section 3 presents results on the computational time for the new algorithm and compares with the standard linear convolution algorithm and the Klein and Walther (2017) algorithm. An analysis of the effects on computational time of using a fixed truncation number is also given. In Section 4 we present the results of a small simulation study to investigate the estimation bias that results from truncation and compare out-of-sample forecast performance. Section 5 illustrates our results with two empirical applications. Finally, in Section 6 we give some concluding remarks.

## 2 FFT algorithm for $\mathrm{ARCH}(\infty)$ models

We consider the following generic conditional variance model for the series $\left(\epsilon_{t}\right)_{t=1}^{T}$, which is either an observable series (such as asset returns) or the error term from a (regression) model,

$$
\begin{equation*}
\epsilon_{t}=\sigma_{t} z_{t}, \quad \mathbb{E}_{t-1}\left(\epsilon_{t}^{2}\right)=\sigma_{t}^{2}, \quad \text { and } \quad z_{t} \sim \text { i.i.d. }(0,1) \tag{1}
\end{equation*}
$$

Here, $\left(z_{t}\right)_{t=1}^{T}$ is an independently and identically distributed (i.i.d.) innovation series with mean zero and variance normalized to one, $\left(\sigma_{t}^{2}\right)_{t=1}^{T}$ is the conditional variance series, and $\mathbb{E}_{t-1}$ denotes the expectation conditional on the information available at time $t-1$.

The $\operatorname{ARCH}(\infty)$ model (or representation) for the conditional variance is

$$
\begin{equation*}
\sigma_{t}^{2}=c+\sum_{j=0}^{\infty} \lambda_{j} \epsilon_{t-j}^{2} \tag{2}
\end{equation*}
$$

where $c$ and $\lambda_{j}$ are the corresponding constant and coefficients, respectively. Of course, in practical applications, the series $\left(\epsilon_{t}\right)$ will be available only for $t=1, \ldots, T$, where $T$ denotes the sample size. In practice, therefore, the summation in (2) will need to be truncated at $j=t-1$. This results in the following $\operatorname{ARCH}(\infty)$ model, which can be viewed, for example, as a device to approximate the likelihood function,

$$
\begin{equation*}
\sigma_{t}^{2}=c+\sum_{j=0}^{t-1} \lambda_{j} \epsilon_{t-j}^{2} \tag{3}
\end{equation*}
$$

The summation in (3) is a linear convolution. By a standard linear convolution algorithm, for given series $\left(\epsilon_{t}\right)_{t=1}^{T}$ and $\left(\lambda_{j}\right)_{j=0}^{T-1}$, and a given value of $t$, the calculation (3) requires $2 t$ arithmetic operations ( $t$ multiplications and $t$ additions). Thus, the calculation of the series $\left(\sigma_{t}^{2}\right)_{t=1}^{T}$ requires $\sum_{t=1}^{T} 2 t=T^{2}+T$ arithmetic operations.

When the sample size is large, the computational burden of $T^{2}+T$ arithmetic operations makes the calculation of the conditional variance series $\left(\sigma_{t}^{2}\right)_{t=1}^{T}$ very slow. This can render optimization of a likelihood function, and certainly simulations and bootstrapping, infeasible. To overcome the computational burden of the calculation in (3), Baillie et al. (1996) suggested truncating the summation in (3) at a fixed number, say $n$, which is typically chosen to be 1,000 . This truncation has become standard in the literature. The resulting model is thus

$$
\begin{equation*}
\sigma_{t}^{2}=c+\sum_{j=0}^{\min \{t-1, n\}} \lambda_{j} \epsilon_{t-j}^{2} \tag{4}
\end{equation*}
$$

The required number of arithmetic operations to calculate the series $\left(\sigma_{t}^{2}\right)_{t-1}^{T}$ in (4) is only of order $n T$. Importantly, for a fixed truncation number $n$, the number of operations grows only linearly with the sample size and hence avoids the squared growth required for the calculation in (3).

While the introduction of the truncation number, $n$, in (4) allows for much faster calculation of the conditional variance series $\left(\sigma_{t}^{2}\right)_{t=1}^{T}$, it also introduces an approximation. It is no longer an exact algorithm to calculate the desired series given in (3). In Section 4 we present the results of some Monte Carlo simulations to illustrate and quantify the bias in parameter estimation resulting from this approximation.

We now introduce an alternative method to calculate the conditional variance series $\left(\sigma_{t}^{2}\right)_{t=1}^{T}$ in (3) without truncation using frequency domain techniques. Specifically, we will apply the fast Fourier transform (FFT) combined with the so-called circular convolution theorem. As discussed in Jensen and Nielsen (2014) in the context of calculating fractional differences, this method reduces the number of arithmetic operations to order $T \log T$. To describe this method, we will need the following two definitions.

Definition 1. The discrete Fourier transform (DFT), $f=\left(f_{j}\right)_{j=1}^{T}$, of a series $a=\left(a_{t}\right)_{t=1}^{T}$ is the solution to

$$
a=T^{-1} F f
$$

where $F$ is the Fourier matrix given by $F_{j k}=w_{T}^{(j-1)(k-1)}$ with $w_{T}=e^{2 \pi \mathrm{i} / T}$ and $\mathrm{i}=\sqrt{-1}$.
Definition 2. Let $a_{j}$ and $b_{j}$ be two periodic sequences, meaning that $a_{j+N T}=a_{j}$ and $b_{j+N T}=b_{j}$ for $N= \pm 0, \pm 1, \pm 2, \ldots$. Then the circular convolution of $\left(a_{j}\right)_{j=1}^{T}$ and $\left(b_{j}\right)_{j=1}^{T}$ is defined as

$$
(a \circledast b)_{t}=\sum_{j=1}^{T} a_{j} b_{t-j+1}=\sum_{j=1}^{t} a_{j} b_{t-j+1}+\sum_{j=t+1}^{T} a_{j} b_{T+t-j+1}, \quad t=1, \ldots, T .
$$

We next present two theorems. The first is a finite version of the circular convolution theorem, which shows how the circular convolution in Definition 2 can be calculated using the DFT in Definition 1. For periodic integrable functions, this result can be found in Zygmund (2003, Thm. 1.5, p. 36). The finite version has appeared in the early engineering literature as an application of the FFT; e.g. Stockham (1966, p. 230) and Cooley, Lewis, and Welch (1969, p. 32). Our version, presented in Theorem 1, is taken from Jensen and Nielsen (2014) and the proof can be found there. ${ }^{1}$

Theorem 1. Let $a=\left(a_{t}\right)_{t=1}^{T}$ and $b=\left(b_{t}\right)_{t=1}^{T}$ be two sequences. Then

$$
\begin{equation*}
a \circledast b=T^{-1} F(\bar{F} a \circ \bar{F} b), \tag{5}
\end{equation*}
$$

where $\circ$ denotes element-wise multiplication and $\bar{F}$ is the complex conjugate of $F$.
To apply the result in Theorem 1 in our context, we need to transform the linear convolutions in (3) and (4) into circular convolutions. To economize on notation, we consider only (4), but allow the possibility that $n=T-1$, in which case we obtain (3).

Our second theorem shows that the sequence of conditional variances in (3) or (4) (where the former is obtained by setting $n=T-1$ in the latter) can be calculated by extending the sequences with zeros and applying the result in Theorem 1. Specifically, we extend the vector of errors, $\left(\epsilon_{t}\right)_{t=1}^{T}$, and the vector of coefficients, $\left(\lambda_{j}\right)_{j=0}^{n}$, with zeros to length $2 T-1$. Thus, let the extended vectors be denoted $\tilde{\epsilon}=\left[\epsilon^{\prime}, O_{T-1}^{\prime}\right]$ and $\tilde{\lambda}=\left[\lambda^{\prime}, O_{2 T-1-(n+1)}^{\prime}\right]$, respectively, where $O_{m}$ denotes an $m \times 1$ vector of zeros. We then prove in Theorem 2 that the linear convolution of $\left(\epsilon_{t}^{2}\right)_{t=1}^{T}$ and $\left(\lambda_{j}\right)_{j=0}^{n}$ in (4) can be found as the first $T$ elements of the circular convolution of $\left(\tilde{\epsilon}_{t}^{2}\right)_{t=1}^{2 T-1}$ and $\left(\tilde{\lambda}_{j}\right)_{j=0}^{2 T-2}$.
Theorem 2. Let $\tilde{\epsilon}^{2}$ and $\tilde{\lambda}$ denote the $(2 T-1) \times 1$ extended vectors of squared errors and coefficients, respectively. In particular, the $t$ 'th element of $\tilde{\epsilon}^{2}$ is $\tilde{\epsilon}_{t}^{2}$. Then the vector of conditional variances, $\left(\sigma_{t}^{2}\right)_{t=1}^{T}, \underset{\sim}{i n}(3)$ or (4) can be calculated as the first $T$ elements of the $(2 T-1) \times 1$ vector $c+T^{-1} F\left(\bar{F} \tilde{\lambda} \circ \bar{F} \tilde{\epsilon}^{2}\right)$.

Because of the truncation number, $n$, the result in Theorem 2 may appear to be different from the corresponding result in Theorem 2 of Jensen and Nielsen (2014), which does not consider truncation. Although Theorem 2 does in fact follow from Theorem 2 of Jensen and Nielsen (2014), it seems intructive to give a short proof of Theorem 2 to explicitly demonstrate the role of the extended vectors $\tilde{\epsilon}$ and $\tilde{\lambda}$.
Proof. From Theorem 1 we know that $\tilde{\lambda} \circledast \tilde{\epsilon}^{2}=T^{-1} F\left(\bar{F} \tilde{\lambda} \circ \bar{F} \tilde{\epsilon}^{2}\right)$. Therefore, it only remains to be shown that $\left(\tilde{\lambda} \circledast \tilde{\epsilon}^{2}\right)_{t}=\sum_{j=0}^{\min \{t-1, n\}} \lambda_{j} \epsilon_{t-j}^{2}$ for $t=1, \ldots, T$. From Definition 2, noting that the extended sequences have $2 T-1$ elements and that the $\tilde{\lambda}_{j}$ sequence starts at index $j=0$, we find that

$$
\begin{equation*}
\left(\tilde{\lambda} \circledast \tilde{\epsilon}^{2}\right)_{t}=\sum_{j=1}^{t} \tilde{\lambda}_{j-1} \tilde{\epsilon}_{t-j+1}^{2}+\sum_{j=t+1}^{T+t-1} \tilde{\lambda}_{j-1} \tilde{\epsilon}_{2 T+t-j}^{2}+\sum_{j=T+t-1}^{2 T-1} \tilde{\lambda}_{j-1} \tilde{\epsilon}_{2 T+t-j}^{2} \tag{6}
\end{equation*}
$$

[^1]When $j=t+1, \ldots, T+t-1$ we have $2 T+t-j=T+1, \ldots, 2 T-1$, so that $\tilde{\epsilon}_{2 T+t-j}^{2}=0$ by definition. Similarly, when $j=T+t+1, \ldots, 2 T-1$ and $t=1, \ldots, T$ we have $\tilde{\lambda}_{j-1}=0$ by definition. This leaves only the first term on the right-hand side of (6). When $t-1 \leq n$, this term is equal to $\sum_{j=0}^{\min \{t-1, n\}} \tilde{\lambda}_{j} \tilde{\epsilon}_{t-j}^{2}$, which is what we needed to show. When $t-1>n$, we split the first term on the right-hand side of (6) as

$$
\sum_{j=1}^{t} \tilde{\lambda}_{j-1} \tilde{\epsilon}_{t-j+1}^{2}=\sum_{j=0}^{n} \tilde{\lambda}_{j} \tilde{\epsilon}_{t-j}^{2}+\sum_{j=n+1}^{t-1} \tilde{\lambda}_{j} \tilde{\epsilon}_{t-j}^{2}
$$

For $j=n+1, \ldots, t-1$ and $t=1, \ldots, T$ we have $\tilde{\lambda}_{j}=0$ by definition, so that the last summation is zero, which proves the desired result.

Theorem 2 shows that the linear convolution of $\left(\epsilon_{t}^{2}\right)_{t=1}^{T}$ and $\left(\lambda_{j}\right)_{j=0}^{n}$ in (4) can be found as the first $T$ elements of the circular convolution of $\left(\tilde{\epsilon}_{t}^{2}\right)_{t=1}^{2 T-1}$ and $\left(\tilde{\lambda}_{j}\right)_{j=0}^{2 T-2}$, which in turn can be calculated using the DFT. The power of this result lies in the fact that the required DFTs can be calculated extremely efficiently by the FFT. Indeed, the latter requires only an order $T \log T$ arithmetic operations, as shown by Cooley and Tukey (1965). This implies that, not only can the conditional variances be calculated an order of magnitude faster than using the standard linear convolution method, but also that the order of magnitude is the same with and without truncation. That is, there is no need to introduce a truncation number to speed up computation as in (4).

Most standard implementations of the FFT requires the length of the vectors to be a power of two. ${ }^{2}$ This is easily accommodated by further extending the vectors with zeros. Thus, to apply the FFT in the calculations, the vectors $\epsilon$ and $\lambda$ must be extended with zeros such that the number of elements in the vectors is equal to the smallest power of two that is at least $2 T-1$. This ensures that both Theorem 2 and the FFT can be applied.

The implementation described in the previous paragraph implies that conditional variance series for a range of sample sizes can be calculated in the same amount of time. For example, consider $T=1,025$ and $T=2,048$. In both cases, the smallest power of two that is at least $2 T-1$ is 11 , and consequently one must extend with 3,071 zeros when $T=1,025$ and 2,048 zeros when $T=2,048$. In both situations, the extended vectors have 4,096 elements, and thus it will take the same amount of time to compute the conditional variances using the FFT-based method in Theorem 2.

In Listings 1 and 2 we present our Matlab codes to implement the standard linear convolution algorithm and the FFT-based algorithm in Theorem 2, respectively. The former is a simple application of Matlab's filter function. Implementations of the FFT-based method in Theorem 2 for R and Ox can be found in Listings 3 and 4, respectively. All codes are downloadable from the authors' websites.

Listing 1: Matlab code to calculate $\operatorname{ARCH}(\infty)$ conditional variances by LC method

```
function [sigma2_arch] = arch_lc(cst, epsilon, lambda)
    sigma2_arch = cst + filter(lambda, 1, epsilon.^2);
end
```

[^2]Listing 2: Matlab code to calculate $\operatorname{ARCH}(\infty)$ conditional variances by FFT method

```
function [sigma2_arch] = arch_fft(cst, epsilon, lambda)
    T = size(epsilon, 1);
    np2 = 2.^nextpow2(2*T-1);
    sigma2_arch = ifft(fft(epsilon.^2, np2).*fft(lambda, np2));
    sigma2_arch = cst + sigma2_arch(1:T);
end
```

Listing 3: R code to calculate $\operatorname{ARCH}(\infty)$ conditional variances by FFT method

```
arch_fft <- function(cst, epsilon, lambda){
    iT <- length(epsilon)
    np2 <- nextn(2*iT-1, 2)
    sigma2_arch <- fft(fft(c(lambda, rep(0, np2-iT))) * fft(c(epsilon^2,
        rep(0, np2-iT))), inverse = T) / np2;
    sigma2_arch <- cst + sigma2_arch[1:iT]
    return(Re(sigma2_arch))
}
```

Listing 4: Ox code to calculate $\mathrm{ARCH}(\infty)$ conditional variances by FFT method

```
arch_fft(const cst, const epsilon, const lambda)
{
    decl T, sigma2_arch;
    T = rows(epsilon);
    sigma2_arch = fft(cmul(fft(lambda'~zeros(1,T)), fft((epsilon.^2)'~zeros
        (1,T)), 2);
    return cst + sigma2_arch'[0:T-1];
}
```


### 2.1 Example: FIGARCH model

We now consider an example that we will apply extensively in the remainder. The most well-known $\operatorname{ARCH}(\infty)$ model for the conditional variance is probably the $\operatorname{FIGARCH}(p, d, q)$ model of Baillie et al. (1996). For that model, the conditional variance specification is

$$
\begin{equation*}
\beta(L) \sigma_{t}^{2}=\omega+\left(\beta(L)-\phi(L)(1-L)^{d}\right) \epsilon_{t}^{2} \tag{7}
\end{equation*}
$$

where $\beta(L)=1-\sum_{i=1}^{p} \beta_{i} L^{i}$ and $\phi(L)=1-\sum_{i=1}^{\max \{p, q\}} \phi_{i} L^{i}$ are polynomials in the lag operator, $L$, whose roots all lie outside the complex unit circle. The fractional difference operator $(1-L)^{d}$ is defined in terms of the fractional coefficients $\pi_{j}(u)=u(u+1) \ldots(u+j-1) / j$ ! from the binomial expansion of $(1-z)^{-u}$, so that, for a generic series $\left(x_{t}\right)_{t=1}^{T}$, we have $(1-L)^{d} x_{t}=\sum_{j=0}^{t-1} \pi_{j}(-d) x_{t-j}$.

The specification (7) can be rearranged to get an $\operatorname{ARCH}(\infty)$ representation. For simplicity, suppose $p=q=1$, so that $\beta(L)=1-\beta L$ and $\phi(L)=1-\phi L$. This is the most commonly applied variant of the FIGARCH model. Then (7) can be written in the form (2)
with $c=\omega /(1-\beta)$ and

$$
\lambda_{j}= \begin{cases}0 & \text { for } j=0  \tag{8}\\ \phi-\beta+d & \text { for } j=1 \\ \beta \lambda_{j-1}+\phi \pi_{j-1}(-d)-\pi_{j}(-d) & \text { for } j \geq 2\end{cases}
$$

see Baillie et al. (1996) for details. Let $g_{1}=\phi-\beta+d$ and $g_{j}=\phi \pi_{j-1}(-d)-\pi_{j}(-d)$ for $j \geq 2$. Then the $\lambda_{j}$ coefficients in (8) can be rewritten as

$$
\lambda_{j}= \begin{cases}0 & \text { for } j=0  \tag{9}\\ \sum_{i=0}^{j-1} \beta^{i} g_{j-i} & \text { for } j \geq 1\end{cases}
$$

The last term in (9) is clearly a linear convolution. To speed up computation, the series of $\lambda_{j}$ coefficients should also be calculated using our FFT-based algorithm.

It follows that efficient calculation of the conditional variances in the $\operatorname{FIGARCH}(1, d, 1)$ model should use our FFT-based algorithm twice: once to calculate the conditional variances given the $\lambda_{j}$ coefficients, and once to calculate the $\lambda_{j}$ coefficients given the model parameters. The former calculation is described in detail in Theorem 2. Similarly, $\left(\lambda_{j}\right)_{j=1}^{n}$ can be found as the first $n$ elements of the $(2 n-1) \times 1$ vector $n^{-1} F(\bar{F} \tilde{\beta} \circ \tilde{g})$, where $\tilde{\beta}$ and $\tilde{g}$ denote the vectors $\left(\beta^{j}\right)_{j=0}^{n-1}$ and $\left(g_{j}\right)_{j=1}^{n}$ extended with zeros to length of the smallest power of two that is at least $2 n-1$. Matlab implementations of the calculation of $\left(\lambda_{j}\right)_{j=0}^{n}$ are shown in Listings 5 and 6 for the linear convolution algorithm and the FFT-based algorithm, respectively.

Listing 5: Matlab code to calculate $\lambda_{j}$ coefficients of $\operatorname{FIGARCH}(1, d, 1)$ by LC method

```
function [lambda] = lambda_lc(phi, beta, d, nTrunc)
    lambda = zeros(nTrunc+1,1);
    k = (1:nTrunc)';
    pij = cumprod((k(1:end)-d-1)./(k(1:end)));
    g = zeros(nTrunc,1);
    g(1) = phi - beta + d;
    g(2:end) = phi*pij(1:end-1) - pij(2:end);
    lambda(1) = 0;
    lambda(2:end) = filter(beta.^(0:nTrunc-1), 1, g);
end
```

Listing 6: Matlab code to calculate $\lambda_{j}$ coefficients of $\operatorname{FIGARCH}(1, d, 1)$ by FFT method

```
function [lambda] = lambda_fft(phi, beta, d, nTrunc)
    lambda = zeros(nTrunc+1,1);
    k = (1:nTrunc)';
    pij = cumprod((k(1:end)-d-1)./(k(1:end)));
    g = zeros(nTrunc,1);
    g(1) = phi - beta + d;
    g(2:end) = phi*pij(1:end-1) - pij(2:end);
    np2 = 2.^nextpow2(2*nTrunc-1);
    lambda(1) = 0;
    tmp = ifft(fft(beta.^(0:nTrunc-1)', np2).*fft(g, np2));
```

```
11 lambda(2:end) = tmp(1:nTrunc);
```

end

### 2.2 Application to non-ARCH ( $\infty$ ) models

Our FFT-based methodology can also be applied with great benefit to some models that are not even in the class of $\operatorname{ARCH}(\infty)$ models. In particular, the linear ARCH model of Robinson (1991) and Giraitis et al. (2000) is given by $\sigma_{t}=c+\sum_{j=1}^{\infty} \lambda_{j} \epsilon_{t-j}$, c.f. (2). Clearly, our FFTbased algorithms described above and presented in Listings 2 and 6 can easily be adopted to this model by simply replacing $\sigma_{t}^{2}$ and $\epsilon_{t-j}^{2}$ with $\sigma_{t}$ and $\epsilon_{t-j}$, respectively. Similarly, the fractionally integrated asymmetric power ARCH model of Tse (1998) is given as in (7), but with $\sigma_{t}^{2}$ and $\epsilon_{t}^{2}$ replaced by $\sigma_{t}^{\delta}$ and $h\left(\epsilon_{t}\right)$, respectively, where $h(\epsilon)=(|\epsilon|-\gamma \epsilon)^{\delta}$ and $\delta>0$. Again, our FFT-based algorithms apply with minimal changes to this model.

Finally, the popular fractionally integrated exponential GARCH (FIEGARCH) model of Bollerslev and Mikkelsen (1996) is given as in (7), but with $\sigma_{t}^{2}$ replaced by $\log \left(\sigma_{t}^{2}\right)$ and $\epsilon_{t}^{2}$ replaced by $h\left(\epsilon_{t} / \sigma_{t}\right)$, where $h(z)=\gamma_{1} z+\gamma_{2}(|z|-\mathbb{E}|z|)$. Thus, because $\sigma_{t}$ enters non-linearly on the right-hand side of the conditional variance equation, the sequence of conditional variances for the FIEGARCH model cannot be written as a convolution and our methodology does not apply to the calculation of the conditional variance sequence. However, the $\lambda_{j}$ coefficients for this model are identical to those for the FIGARCH model, and the calculation of these can therefore take advantage of our FFT-method as in Listing 6. At least this will reduce the computational time of calculating the $\lambda_{j}$ coefficients for this model from $O\left(T^{2}\right)$ to $O(T \log T)$ compared with standard linear convolution.

## 3 Computational time

In this section, we compare the computational time of our new algorithm in Theorem 2 and Listings 2,6 with the algorithm in Klein and Walther (2017) as well as the standard linear convolution implementation in Listings 1,5. All computations were performed in Matlab on a desktop with an Intel Core i5 (5250U) 1.6GHz processor running macOS Catalina 10.15.6.

In our first numerical experiment, we compare the difference in computational time for standard linear convolution and our FFT-based algorithm as a function of sample size. Specifically, for a given vector of coefficients, $[d, \phi, \beta, \omega]=[0.46,0.27,0.65,0.02]$, we calculate the conditional variances for the $\operatorname{FIGARCH}(1, d, 1)$ model using the method in Listings 1,5 (blue line) and the FFT-based algorithm in Listings 2,6 (red line). That is, we calculate the $\lambda_{j}$ coefficients in (9) followed by the conditional variances in (3), i.e. without truncation, with random numbers for $\epsilon_{t}$. This is done a large number of times ( 10,000 for $T<4,096$ and $1,000$ for $T \geq 4,096)$, and the median time in milliseconds across repetitions is reported in Figure 1 as a function of sample size. In Panel (a) the axes are linear and in Panel (b) the axes are logarithmic. In particular, the figure clearly shows the different orders of magnitude of the computational time for the two methods ( $T^{2}$ vs. $T \log T$ ). The figure also demonstrates how the computational time for the FFT-based algorithm is essentially a step function of the sample size, because of the padding to the smallest power of two that is at least $2 T-1$.

We also notice from Figure 1(b) that for $T=500$ to $T=1,000$, the two methods perform equally well, while linear convolution does better for smaller sample sizes and the FFTbased algorithm does better for larger sample sizes. Thus, to obtain the fastest possible

Figure 1: Computational time

algorithm across all sample sizes, we recommend implementing the linear convolution method for smaller sample sizes and the FFT-based algorithm for larger sample sizes. This can easily be achieved by combining the implementations in Listings 1 and 2 using an if statement, and similarly for Listings 5 and 6.

In our next experiment, we focus on the effect of truncation on computational speed, and compare linear convolution (LC), our FFT-based algorithm, and the algorithm of Klein and Walther (2017, henceforth KW). ${ }^{3}$ The setup is the same as in Figure 1, except for the introduction of the truncation number. To reflect samples sizes in modern empirical work (see references in the introduction), we consider $T=3,000$ to $T=500,000$ and truncation numbers from $n=1,000$ to $n=5,000$, as well as no truncation, $n=T-1$. For each $(T, n)$ pair, we repeated the calculation 10,000 times (for fixed $n$ ) or 1,000 times (for $n=T-1$ ), except for KW and LC with $T=500,000$ and $n=T-1$, where we only used 100 repetitions. In Table 1 we report the resulting median time across repetitions in milliseconds.

In Table 1 we first note that our FFT-based method is faster than the other methods for all sample sizes and truncation numbers considered. While the computational times of both the KW and LC algorithms are very sensitive to the choice of truncation number, the FFT-based method is much less so. For example, from $n=1,000$ to $n=T-1$, the computational time of the KW and LC algorithms increase by a factor of $7-8$ for $T=3,000$ and several hundred for $T=500,000$. On the other hand, the computational time of the FFT-based method only approximately doubles for either sample size. Hence, with our FFT-based algorithm, there is really no need for the truncation number and the associated approximation.

Using the exact filter to calculate the conditional variances, i.e. without truncation ( $n=$ $T-1$ ), the difference in computational time between the KW and LC algorithms on the one hand and our FFT-based algorithm on the other hand is very substantial. For $T=10,000$, which is a rather common sample size in finance, the FFT-based algorithm is $15-20$ times faster than the KW and LC algorithms. For the largest sample size considered, $T=500,000$,

[^3]Table 1: Computational time for different sample sizes and truncation numbers

|  | $n=1,000$ |  |  |  | $n=2,000$ |  |  |
| ---: | :---: | :---: | :---: | :---: | :---: | :---: | ---: |
| $T$ | KW | LC | FFT |  | KW | LC | FFT |
| 3,000 | 0.48 | 0.64 | $\mathbf{0 . 3 8}$ |  | 1.54 | 1.72 | $\mathbf{0 . 5 2}$ |
| 5,000 | 0.75 | 0.99 | $\mathbf{0 . 6 3}$ |  | 1.56 | 2.44 | $\mathbf{0 . 7 7}$ |
| 10,000 | 1.40 | 1.85 | $\mathbf{1 . 1 9}$ |  | 2.19 | 4.25 | $\mathbf{1 . 3 2}$ |
| 25,000 | 2.94 | 4.46 | $\mathbf{2 . 5 9}$ |  | 3.80 | 9.70 | $\mathbf{2 . 7 5}$ |
| 50,000 | 6.51 | 8.80 | $\mathbf{6 . 4 7}$ |  | 7.48 | 18.8 | $\mathbf{6 . 6 5}$ |
| 100,000 | 15.5 | 17.7 | $\mathbf{1 3 . 3}$ |  | 16.5 | 37.1 | $\mathbf{1 3 . 5}$ |
| 500,000 | 88.0 | 88.5 | $\mathbf{7 9 . 6}$ |  | 90.4 | 183 | $\mathbf{7 9 . 8}$ |
|  |  | $n=5,000$ |  |  |  | $n=T-1$ |  |
| $T$ | KW | LC | FFT |  | KW | LC | FFT |
| 3,000 | - | - | - |  | 4.05 | 4.70 | $\mathbf{0 . 6 8}$ |
| 5,000 | - | - | - |  | 12.1 | 14.2 | $\mathbf{1 . 3 0}$ |
| 10,000 | 11.1 | 23.9 | $\mathbf{1 . 9 2}$ |  | 37.6 | 51.4 | $\mathbf{2 . 6 1}$ |
| 25,000 | 12.6 | 53.4 | $\mathbf{3 . 3 3}$ | 205 | 352 | $\mathbf{6 . 2 8}$ |  |
| 50,000 | 16.2 | 102 | $\mathbf{7 . 1 7}$ |  | 868 | 1,611 | $\mathbf{1 4 . 2}$ |
| 100,000 | 25.3 | 201 | $\mathbf{1 4 . 1}$ | 3,440 | 6,610 | $\mathbf{3 4 . 0}$ |  |
| 500,000 | 99.8 | 988 | $\mathbf{8 0 . 0}$ | 147,506 | 293,788 | $\mathbf{1 9 2}$ |  |

Notes: Median computational time in milliseconds across 10,000 repetitions (fixed $n$ ) or 1,000 repetitions ( $n=T-1$ ) for different sample sizes and truncation numbers (only 100 repetitions for KW and LC with $T=500,000$ and $n=T-1$ ). KW is the algorithm of Klein and Walther (2017), LC is linear convolution (Listings 1,5), and FFT is our proposed algorithm (Listings 2,6). The fastest algorithm for each ( $T, n$ ) pair is highlighted in bold.
the FFT-based algorithm is over 750 times faster than the KW algorithm and 1500 times faster than the standard LC algorithm.

The numbers reported in Figure 1 and Table 1 are difficult to extrapolate directly to the time required for, e.g., estimation of a model or a simulation study. This is because, in estimation, each evaluation of the likelihood function and its derivatives involves several convolutions, and this is repeated for many iterations to obtain convergence. Practical examples of such timings and extrapolations to simulations are given at the end of Section 5.2.

## 4 Estimation bias

Since we are able to compute the sequence of conditional variances, and hence the likelihood function, of any model with an $\operatorname{ARCH}(\infty)$ representation very quickly using our FFT-based algorithm, we can explore the effects on the estimators from approximating the calculations. That is, we can simulate the estimation bias that results from using a fixed truncation number in the calculation of the conditional variances in (4). As an alternative to (4), Baillie et al. (1996) suggested terminating the summation in (4) at a fixed $n$ and replacing unobserved pre-sample values of $\epsilon^{2}$ by the unconditional sample variance, $T^{-1} \sum_{t=1}^{T} \epsilon_{t}^{2}$. We also consider

Table 2: Simulated bias and MCSE for $\operatorname{FIGARCH}(1, d, 1)$ model—DGP 1

| T | Par. | $n=1,000$ (P) |  | $n=1,000$ |  | $n=T-1$ |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Bias | MCSE | Bias | MCSE | Bias | MCSE |
| 5,000 | $d$ | 0.0808 | 0.0923 | 0.0036 | 0.0462 | -0.0042 | 0.0486 |
|  | $\phi$ | -0.0244 | 0.0434 | -0.0014 | 0.0349 | -0.0036 | 0.0356 |
|  | $\beta$ | 0.0511 | 0.0790 | 0.0016 | 0.0597 | $-0.0078$ | 0.0601 |
|  | $\omega$ | $-3.1 \times 10^{-5}$ | $3.7 \times 10^{-5}$ | $1.5 \times 10^{-5}$ | $3.7 \times 10^{-5}$ | $9.7 \times 10^{-6}$ | $3.4 \times 10^{-5}$ |
| 10,000 | $d$ | 0.0648 | 0.0505 | 0.0125 | 0.0310 | -0.0013 | 0.0331 |
|  | $\phi$ | -0.0139 | 0.0250 | 0.0010 | 0.0237 | -0.0017 | 0.0246 |
|  | $\beta$ | 0.0474 | 0.0420 | 0.0130 | 0.0338 | $-0.0028$ | 0.0361 |
|  | $\omega$ | $-1.6 \times 10^{-5}$ | $2.3 \times 10^{-5}$ | $2.0 \times 10^{-5}$ | $2.5 \times 10^{-5}$ | $5.2 \times 10^{-6}$ | $2.1 \times 10^{-5}$ |
| 25,000 | $d$ | 0.0504 | 0.0289 | 0.0203 | 0.0207 | -0.0006 | 0.0208 |
|  | $\phi$ | -0.0070 | 0.0146 | 0.0020 | 0.0143 | -0.0007 | 0.0151 |
|  | $\beta$ | 0.0407 | 0.0256 | 0.0212 | 0.0218 | $-0.0012$ | 0.0226 |
|  | $\omega$ | $1.8 \times 10^{-5}$ | $2.3 \times 10^{-5}$ | $4.1 \times 10^{-5}$ | $2.4 \times 10^{-5}$ | $3.3 \times 10^{-6}$ | $1.5 \times 10^{-5}$ |
| 50,000 | $d$ | 0.0436 | 0.0200 | 0.0247 | 0.0157 | -0.0001 | 0.0143 |
|  | $\phi$ | -0.0040 | 0.0104 | 0.0020 | 0.0102 | -0.0005 | 0.0107 |
|  | $\beta$ | 0.0372 | 0.0184 | 0.0254 | 0.0164 | $-0.0005$ | 0.0157 |
|  | $\omega$ | $5.3 \times 10^{-5}$ | $2.5 \times 10^{-5}$ | $6.9 \times 10^{-5}$ | $2.5 \times 10^{-5}$ | $2.4 \times 10^{-6}$ | $1.2 \times 10^{-5}$ |
| 100,000 | $d$ | 0.0393 | 0.0149 | 0.0279 | 0.0127 | $4.0 \times 10^{-5}$ | 0.0101 |
|  | $\phi$ | -0.0019 | 0.0074 | 0.0020 | 0.0074 | $-0.0003$ | 0.0077 |
|  | $\beta$ | 0.0352 | 0.0142 | 0.0283 | 0.0130 | $-0.0002$ | 0.0112 |
|  | $\omega$ | $9.8 \times 10^{-5}$ | $2.8 \times 10^{-5}$ | $1.1 \times 10^{-4}$ | $2.8 \times 10^{-5}$ | $1.8 \times 10^{-6}$ | $1.1 \times 10^{-5}$ |

Notes: Simulations are based on 10,000 replications. "(P)" denotes fixed truncation with pre-sample values equal to the unconditional sample variance. DGP 1 has $[d, \phi, \beta, \omega]=[0.4 ; 0.2 ; 0.6 ; 0.0001]$.
this possibility, and denote the results by " $(\mathrm{P})$ " (for pre-sample) in the following tables. ${ }^{4}$
Specifically, in this section we simulate the bias of the estimated coefficients in the baseline FIGARCH $(1, d, 1)$ model for a range of sample sizes, $T \in\{5,000 ; 10,000 ; 25,000 ; 50,000 ; 100,000\}$, and either fixed truncation, $n=1,000$, or no truncation, $n=T-1$. For each sample size, we simulate 10,000 samples from the $\operatorname{FIGARCH}(1, d, 1)$ model given by (1) and (7) with $z_{t} \sim$ i.i.d. $\mathcal{N}(0,1)$. We consider two different parameter values in the data generating process (DGP). First, DGP 1 has $\theta=[d, \phi, \beta, \omega]^{\prime}=[0.4 ; 0.2 ; 0.6 ; 0.0001]^{\prime}$. Second, we consider DGP 2 with $\theta=[0.4 ; 0.28 ; 0.68 ; 0.0001]^{\prime}$ to investigate the effect of a larger value of $\beta$ on the bias.

Given observations $\left(\epsilon_{t}\right)_{t=1}^{T}$ and truncation number $n$, the quasi-maximum-likelihood estimator is

$$
\begin{equation*}
\hat{\theta}=\underset{\theta}{\arg \max } \log L(\theta) \quad \text { with } \quad \log L(\theta)=-\frac{1}{2} \sum_{t=1}^{T}\left(\log \left(\sigma_{t}^{2}\right)+\frac{\epsilon_{t}^{2}}{\sigma_{t}^{2}}\right) \tag{10}
\end{equation*}
$$

where $\sigma_{t}^{2}$ is calculated from (4) and (9). The maximization is unconstrained, but inequality constraints on the parameters that guarantee non-negativity of the conditional variances are

[^4]Table 3: Simulated bias and MCSE for $\operatorname{FIGARCH}(1, d, 1)$ model—DGP 2

| $T$ | Par. | $n=1,000$ (P) |  | $n=1,000$ |  | $n=T-1$ |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | Bias | MCSE | Bias | MCSE | Bias | MCSE |
| 5,000 | $d$ | 0.1123 | 0.1098 | 0.0087 | 0.0465 | -0.0046 | 0.0497 |
|  | $\phi$ | -0.0502 | 0.0578 | -0.0029 | 0.0325 | -0.0009 | 0.0337 |
|  | $\beta$ | 0.0577 | 0.0622 | 0.0061 | 0.0397 | $-0.0047$ | 0.0423 |
|  | $\omega$ | $-4.2 \times 10^{-5}$ | $2.7 \times 10^{-5}$ | $1.1 \times 10^{-5}$ | $3.1 \times 10^{-5}$ | $8.8 \times 10^{-6}$ | $3.0 \times 10^{-5}$ |
| 10,000 | $d$ | 0.0886 | 0.0673 | 0.0204 | 0.0330 | -0.0019 | 0.0350 |
|  | $\phi$ | -0.0353 | 0.0347 | -0.0049 | 0.0224 | -0.0007 | 0.0235 |
|  | $\beta$ | 0.0493 | 0.0404 | 0.0148 | 0.0276 | $-0.0023$ | 0.0296 |
|  | $\omega$ | $-2.6 \times 10^{-5}$ | $2.4 \times 10^{-5}$ | $1.5 \times 10^{-5}$ | $2.5 \times 10^{-5}$ | $5.4 \times 10^{-6}$ | $2.2 \times 10^{-5}$ |
| 25,000 | $d$ | 0.0684 | 0.0347 | 0.0315 | 0.0228 | $-0.0009$ | 0.0223 |
|  | $\phi$ | -0.0239 | 0.0178 | -0.0074 | 0.0144 | -0.0003 | 0.0146 |
|  | $\beta$ | 0.0413 | 0.0230 | 0.0227 | 0.0183 | -0.0010 | 0.0186 |
|  | $\omega$ | $-6.1 \times 10^{-6}$ | $2.3 \times 10^{-5}$ | $3.3 \times 10^{-5}$ | $2.3 \times 10^{-5}$ | $3.4 \times 10^{-6}$ | $1.6 \times 10^{-5}$ |
| 50,000 | $d$ | 0.0597 | 0.0232 | 0.0373 | 0.0178 | $-0.0003$ | 0.0154 |
|  | $\phi$ | -0.0193 | 0.0123 | -0.0090 | 0.0106 | -0.0004 | 0.0104 |
|  | $\beta$ | 0.0376 | 0.0165 | 0.0266 | 0.0142 | $-0.0005$ | 0.0132 |
|  | $\omega$ | $-4.0 \times 10^{-5}$ | $2.4 \times 10^{-5}$ | $5.8 \times 10^{-5}$ | $2.4 \times 10^{-5}$ | $2.3 \times 10^{-6}$ | $1.3 \times 10^{-5}$ |
| 100,000 | $d$ | 0.0547 | 0.0176 | 0.0415 | 0.0149 | $-3.4 \times 10^{-5}$ | 0.0111 |
|  | $\phi$ | -0.0166 | 0.0100 | -0.0103 | 0.0095 | $-0.0003$ | 0.0079 |
|  | $\beta$ | 0.0354 | 0.0136 | 0.0291 | 0.0122 | $-0.0003$ | 0.0097 |
|  | $\omega$ | $-8.3 \times 10^{-5}$ | $2.8 \times 10^{-5}$ | $9.4 \times 10^{-5}$ | $2.8 \times 10^{-5}$ | $1.8 \times 10^{-6}$ | $1.1 \times 10^{-5}$ |

Notes: Simulations are based on 10,000 replications. "(P)" denotes fixed truncation with pre-sample values equal to the unconditional sample variance. DGP 2 has $[d, \phi, \beta, \omega]=[0.4 ; 0.28 ; 0.68 ; 0.0001]$.
checked post-estimation; see Baillie et al. (1996, footnote 19) and Conrad and Haag (2006).
The Monte Carlo simulation results can be found in Tables 2 and 3 for DGPs 1 and 2, respectively. For each sample size and truncation number, we report the bias and Monte Carlo standard error (MCSE) for the four parameters.

We first observe from Tables 2 and 3 that replacing unobserved pre-sample observations with unconditional variance estimates appears to induce a large bias. Indeed, the biases in the (P) column are substantially larger than in other columns, and particularly the estimates of $d$ and $\beta$ are heavily upwards biased. It would appear that adding the effect of many (up to $n-1$ ) constant observations near the start of the sample, but none later, induces a type of non-stationarity which biases estimates of $d$ and $\beta$ upwards. Johansen and Nielsen (2016) analyze bias in an ARFIMA model using higher-order asymptotic theory, and show that inclusion of a level parameter (in our model this is $\omega$ ) essentially eliminates all bias coming from pre-sample values, so there appears to be no need for plugging in arbitrary constant pre-sample values. A similar mechanism may apply in our setup.

Next, disregarding the $(\mathrm{P})$ column, we see that the truncation at $n=1,000$ induces a

Table 4: Simulated out-of-sample forecast RMSE $(\times 1,000)$ for $\operatorname{FIGARCH}(1, d, 1)$ model

| T | DGP 1 |  |  | DGP 2 |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | $n=1,000$ (P) | $n=1,000$ | $n=T-1$ | $n=1,000$ (P) | $n=1,000$ | $n=T-1$ |
| 5,000 | 1.9797 | 1.4328 | 1.2003 | 2.2205 | 1.5488 | 1.2619 |
| 10,000 | 1.8057 | 1.4575 | 0.9092 | 2.0225 | 1.5901 | 0.9440 |
| 25,000 | 1.8871 | 1.7448 | 0.7159 | 2.1068 | 1.9261 | 0.7488 |
| 50,000 | 2.2862 | 2.2152 | 0.5517 | 2.5440 | 2.4401 | 0.5827 |
| 100,000 | 2.5670 | 2.5736 | 0.4298 | 2.8311 | 2.8332 | 0.4461 |

Notes: We conducted 500 expanding window one-step-ahead out-of-sample forecasts and report the resulting forecast RMSE averaged across 100 simulation replications. " $(\mathrm{P})$ " denotes fixed truncation with pre-sample values equal to the unconditional sample variance. DGP 1 has $[d, \phi, \beta, \omega]=[0.4 ; 0.2 ; 0.6 ; 0.0001]$ and DGP 2 has $[d, \phi, \beta, \omega]=[0.4 ; 0.28 ; 0.68 ; 0.0001]$.
substantial bias that increases with the sample size, $T$. This is expected since a larger sample size implies that a fixed truncation affects a larger proportion of the sample. The parameters $d$ and $\beta$ seem particularly affected by the truncation. The biases in these estimators are likely several parameter standard errors in magnitude for the larger sample sizes.

Comparing across Tables 2 and 3 we note that biases are substantially larger for DGP 2 (in Table 3) compared with DGP 1 (in Table 2). It appears that, as the $\beta$ parameter increases, it is more difficult to identify a given $d$ from highly persistent autoregressive-type dynamics, thus increasing the bias of both $d$ and $\beta .{ }^{5}$

Finally, we conduct a small out-of-sample forecast comparison. For each sample, we estimate the model based on the first $T-k-1$ observations and calculate the one-step ahead forecast of observation $T-k$. We repeat this for $k=1, \ldots, 500$ and calculate the forecast root-mean-squared-error (RMSE) for these 500 one-step-ahead forecasts. We then average the RMSE across 100 simulation replications and report the results $(\times 1,000)$ in Table 4 for a range of sample sizes, truncation numbers, and the two DGPs.

In Table 4 we first note that truncation implies less precise forecasts. For each DGP and for each sample size, the smallest forecast RMSE is obtained by using the untruncated filter. The difference in forecast RMSE between truncation at $n=1,000$ and no truncation is always in favor of no truncation and varies from a reduction of about $16 \%$ (for DGP 1 and $T=5,000$ ) to over $80 \%$ (for DGP 2 and $T=100,000$ ). More generally, the results in Table 4 reflect those in Tables 2 and 3, in the sense that (i) setting pre-sample values equal to the unconditional variance estimate increases forecast RMSE, (ii) forecast RMSE is higher for DGP 2 compared with DGP 1, and (iii) the difference in forecast RMSE between truncation at $n=1,000$ and no truncation increases with the sample size.

In conclusion, the results of our Monte Carlo simulations in Tables 2-4 clearly illustrate the consequences of the approximation implied by truncating the convolution as in (4). The resulting bias in the estimators is substantial, and it is worse for larger sample sizes. Furthermore, the truncation and associated bias result in less precise out-of-sample forecasts. Importantly, however, these problems are easily avoidable by application of our proposed

[^5]Figure 2: Daily DEM-USD exchange rate 3/13/1979-12/31/1998


FFT-based algorithm, which is exact and does not rely on any approximation or truncation.

## 5 Empirical illustrations

The persistence in the volatility of nominal exchange rates has been well-documented in the literature, and exchange rates are often modeled using FIGARCH-type models. Indeed, a daily time series of Deutsche Mark-US dollar (DEM-USD) exchange rates was used as the empirical example in Baillie et al. (1996). An updated data set, covering until the end of the DEM era and start of the Euro, was subsequently analyzed in Baillie, Cecen, and Han (2000). In our first empirical illustration, we consider this data set. As a second empirical illustration, we consider a longer time series of daily US Dollar-British Pound (USD-GBP) exchange rates.

### 5.1 Deutsche Mark-US Dollar exchange rate from 1979 to 1998

In this subsection, we analyze the daily DEM-USD exchange rate from March 13, 1979 to December 31, 1998, for a total of 4,975 spot rate observations. ${ }^{6}$ In Figure 2(a) we plot the spot rate and in Figure 2(b) we plot the continuously compounded percentage returns. It is clear from the figure that the spot exchange rate is nonstationary. Indeed, standard arbitrage arguments would dictate that the spot rate should be a Martingale, which justifies modeling the returns as serially uncorrelated as in (1).

Table 5 reports the quasi-maximum-likelihood estimates for the $\operatorname{FIGARCH}(1, d, 1)$ and FIGARCH $(1, d, 0)$ models with either no truncation or with truncation at $n=1,000$ (with or without setting pre-sample value equal to unconditional variance estimates). Several interesting findings appear from these results. First, regardless of the truncation number, the additional parameter in the $(1, d, 1)$ specification compared with the $(1, d, 0)$ specification is highly significant as measured by a $t$-test on the additional $\phi$ parameter. Thus, we consider mainly the FIGARCH $(1, d, 1)$ model. Second, for this model and data set, we note that the parameter estimates for $d$, and to a lesser extent $\beta$ do vary with the truncation number. Third, with truncation at $n=1,000$ the computing times for our FFT-based method are

[^6]Table 5: FIGARCH models for DEM-USD exchange rate (1979-1998)

| $(p, d, q)$ | $n=1,000$ (P) |  | $n=1,000$ |  | $n=T-1$ |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | $(1, d, 1)$ | $(1, d, 0)$ | $(1, d, 1)$ | $(1, d, 0)$ | $(1, d, 1)$ | $(1, d, 0)$ |
| $\mu$ | 0.0033 | 0.0032 | 0.0038 | 0.0033 | 0.0041 | 0.0035 |
|  | (0.0087) | (0.0087) | (0.0086) | (0.0086) | (0.0086) | (0.0086) |
| $d$ | 0.4544 | 0.2860 | 0.4334 | 0.2741 | 0.4794 | 0.2854 |
|  | (0.0186) | (0.0180) | (0.0194) | (0.0176) | (0.0192) | (0.0168) |
| $\phi$ | 0.2929 | - | 0.2984 | - | 0.2801 | - |
|  | (0.0214) |  | (0.0222) |  | (0.0213) |  |
| $\beta$ | 0.6689 | 0.2143 | 0.6530 | 0.2026 | 0.6778 | 0.2130 |
|  | (0.0146) | (0.0234) | (0.0152) | (0.0236) | (0.0155) | (0.0232) |
| $\omega$ | 0.0131 | 0.0519 | 0.0179 | 0.0613 | 0.0147 | 0.0511 |
|  | (0.0034) | (0.0079) | (0.0036) | (0.0086) | (0.0042) | (0.0095) |
| Iterations | 34 | 16 | 34 | 16 | 37 | 17 |
| Fcn. evals. | 230 | 108 | 236 | 105 | 236 | 105 |
| Time FFT | 191 | 92 | 193 | 88 | 356 | 155 |
| Time LC | 311 | 174 | 277 | 150 | 3,356 | 1,225 |

Notes: Quasi-maximum-likelihood estimates for $\operatorname{FIGARCH}(p, d, q)$ models for the DEM-USD percentage returns from March 14, 1979 to December 31, 1998 for different truncation numbers and $T=4,974$ return observations. Standard errors are reported in parentheses, and " $(\mathrm{P})$ " denotes fixed truncation with presample values equal to the unconditional sample variance. Finally, the number of iterations and the total number of function evaluations are reported, along with the computing time in milliseconds using the FFTbased method and the linear convolution (LC) method.
about two-thirds of those for the LC method, but without truncation $(n=T-1)$ the computing time for our FFT-based method is almost ten times faster than for the LC method.

### 5.2 US Dollar-British Pound exchange rate from 1971 to 2020

In this subsection, we analyze the daily USD-GBP spot exchange rate from October 1, 1971 to October 2, 2020, for a total of 12,300 spot rate observations. The data series was downloaded from the FRED database and is plotted in Figure 3. As in the previous subsection, the spot exchange rate is nonstationary and we model the continuously compounded returns.

The estimation results for the USD-GBP data set are presented in Table 6, which is laid out precisely as Table 5. The results are qualitatively quite similar, except they are more pronounced in Table 6 because of the longer sample.

First of all, the $\operatorname{FIGARCH}(1, d, 1)$ specification is again statistically superior to the FIGARCH $(1, d, 0)$ specification, regardless of truncation number. Secondly, the estimates for $d$, and again to a lesser extent for $\beta$, are now quite different for the three different FIGARCH $(1, d, 1)$ estimates. The estimates in the " $(\mathrm{P})$ " column are larger than the others, as expected based the simulation findings in Section 4. Comparing truncation at $n=1,000$, which is standard in the literature, with no truncation, the estimates of $d$ differ by about one standard error. The parameter $\omega$, which is usually not of primary interest, differs even more.

Figure 3: Daily USD-GBP exchange rate 10/1/1971-10/2/2020


Table 6: FIGARCH models for USD-GBP exchange rate (1971-2020)

| $(p, d, q)$ | $n=1,000$ (P) |  | $n=1,000$ |  | $n=T-1$ |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | $(1, d, 1)$ | $(1, d, 0)$ | $(1, d, 1)$ | $(1, d, 0)$ | $(1, d, 1)$ | (1, $d, 0$ ) |
| $\mu$ | -0.0004 | 0.0007 | 0.0012 | 0.0020 | 0.0016 | 0.0025 |
|  | (0.0048) | (0.0045) | (0.0049) | (0.0046) | (0.0053) | (0.0050) |
| $d$ | 0.4948 | 0.3257 | 0.4230 | 0.3123 | 0.4459 | 0.3136 |
|  | (0.0182) | (0.0140) | (0.0177) | (0.0132) | (0.0194) | (0.0143) |
| $\phi$ | 0.2532 | - | 0.2633 | - | 0.2535 | - |
|  | (0.0177) |  | (0.0167) |  | (0.0170) |  |
| $\beta$ | 0.6543 | 0.2562 | 0.5970 | 0.2429 | 0.6036 | 0.2377 |
|  | (0.0165) | (0.0174) | (0.0163) | (0.0170) | (0.0172) | (0.0175) |
| $\omega$ | 0.0113 | 0.0311 | 0.0140 | 0.0327 | 0.0115 | 0.0250 |
|  | (0.0037) | (0.0070) | (0.0042) | (0.0073) | (0.0050) | (0.0091) |


| Iterations | 34 | 20 | 30 | 20 | 31 | 19 |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: |
| Fcn. evals. | 238 | 132 | 226 | 142 | 238 | 137 |
| Time FFT | 370 | 208 | 356 | 281 | 715 | 493 |
| Time LC | 664 | 422 | 654 | 466 | 16,581 | 9,272 |

Notes: Quasi-maximum-likelihood estimates for $\operatorname{FIGARCH}(p, d, q)$ models for the USD-GBP percentage returns from October 4, 1971 to October 2, 2020 for different truncation numbers and $T=12,299$ return observations. Standard errors are reported in parentheses, and " $(\mathrm{P})$ " denotes fixed truncation with presample values equal to the unconditional sample variance. Finally, the number of iterations and the total number of function evaluations are reported, along with the computing time in milliseconds using the FFTbased method and the linear convolution (LC) method.

The computing time using our FFT-based method is now much faster than the LC method. Without truncation, our FFT-based method is about 23 times faster than the LC method for the $\operatorname{FIGARCH}(1, d, 1)$ model. As an order of magnitude, a simulation study or a bootstrap with 10,000 replications for the $\operatorname{FIGARCH}(1, d, 1)$ model in Table 6 without
truncation would require 46 hours with the LC method but less than 2 hours with our FFTbased algorithm. This result confirms those found in Section 3, and could certainly be the difference between bootstrap inference or simulations being feasible or not.

## 6 Conclusion

In this paper we have considered the computation of the time series of conditional variances for models in the $\operatorname{ARCH}(\infty)$ class. This class is very large and contains many commonly applied models in empirical finance such as the FIGARCH model. For models in the $\operatorname{ARCH}(\infty)$ class, we have provided an exact algorithm for efficient computation of the conditional variances, and hence of the likelihood function. Our algorithm is based on the fast Fourier transform (FFT) and achieves an order of magnitude improvement in computational speed from $O\left(T^{2}\right)$ to $O(T \log T)$. The efficiency of the algorithm allows estimation, as well as simulation and/or bootstrapping of $\mathrm{ARCH}(\infty)$ models, even with very large data sets. We have focused on univariate models, but multivariate models will achieve a proportional increase in computational speed.

Furthermore, to speed up computation it has been completely standard in the literature to resort to a truncation of the $\mathrm{ARCH}(\infty)$ model at a fixed truncation lag. With our proposed algorithm, this is not needed. As additional contributions, we also showed that the elimination of the truncation substantially reduces both the bias of the quasi-maximumlikelihood estimators and the out-of-sample forecast errors.

We illustrated our results with two empirical examples that highlighted both the differences in the computational speed and in the parameter estimates for our FFT-based algorithm and the standard algorithm with and without truncation.

As sample sizes in economics and finance have become longer in recent years (see examples in the introduction), efficient computation has become increasingly more important. As we have shown, this is certainly true for daily time series, and when the models discussed in this paper are applied to high-frequency data this becomes crucially important. Furthermore, the gaining popularity of computationally intensive methods of inference, based on simulation, bootstrapping, or machine-learning techniques, that require estimation of (variations of) each model a large number of times, emphasizes these points even more.

## Data availability statement

The daily DEM-USD exchange rate data that support the findings of this study are openly available in the Federal Reserve H. 10 historical data website at https : //www. federalreserve. gov/releases/H10/hist/dat89_ge.htm and https://www.federalreserve.gov/releases/ H10/hist/dat96_ge.htm. The daily USD-GBP exchange rate data that support the findings of this study are openly available in the FRED database at https://fred.stlouisfed. org/series/DEXUSUK. The data files are included in the online supplementary materials together with the computer programs listed in the paper. The latter are also available on the authors' websites.

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[^1]:    ${ }^{1}$ In Jensen and Nielsen (2014) there is a small typo in the fifth and sixth lines of the proof, where $(t-1)(s-$ 1) should be $(t-1)(u-1)$ in the exponents. However, since the last equality sets $s=u$ the result is the same.

[^2]:    ${ }^{2}$ Some modern implementations require only that the length is a product of powers of small prime numbers, such as $2^{k} 3^{l} 5^{m}$, and the subsequent discussion is easily adapted to such cases.

[^3]:    ${ }^{3}$ The Klein and Walther (2017) Matlab code was downloaded from the publisher's website.

[^4]:    ${ }^{4}$ We are grateful to an anonymous referee for this suggestion.

[^5]:    ${ }^{5}$ We are grateful to an anonymous referee for suggesting a second DGP with a larger value of $\beta$ based on this motivation.

[^6]:    ${ }^{6}$ The Euro was introduced on January 1, 1999. Our data set was downloaded from the Federal Reserve H. 10 historical data website, and covers the same time period as that analyzed in Baillie et al. (2000), but has a slightly different number of observations.

