

ML Estimation of the Parameters of FBM Traffic with Geometrical Sampling

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Abstract

Traffic model based on the fractional Brownian motion (fBm) contains three parameters: the mean rate m , variance parameter a and the Hurst parameter H . The estimation of these parameters by the maximum likelihood (ML) method is studied. Explicit expressions for the ML estimates \hat{m} and \hat{a} in terms of H are given, as well as the expression for the log-likelihood function from which the estimate \hat{H} is obtained as the minimizing argument. A geometric sequence of sampling points, $t_i = \alpha^i$, is introduced in order to see the scaling behaviour of the traffic with fewer samples. It is shown that by a proper ‘descaling’ the traffic process is stationary on this grid leading to a Toeplitz-type covariance matrix. Approximations for the inverted covariance matrix and its determinant are introduced. The accuracy of the estimation algorithm is studied by simulations. Comparisons with corresponding estimates obtained with linear grid show that the geometrical sampling indeed improves the accuracy of the estimate \hat{H} with a given number of samples.

1 Introduction

One of the simplest and most studied models for aggregated data traffic is the fractional Brownian motion (fBm) model [7], which is a model for truly self-similar Gaussian traffic. Though the model has its limitations and, in particular, breaks down at small time scales, it has gained popularity because of its simplicity allowing e.g. the queueing behaviour to be studied analytically by making use of the scaling properties [4, 6]. Furthermore, an important feature of the fBm model is its parsimony [3]: in its basic form the model contains only three parameters, the mean rate m , the variance parameter a and the Hurst parameter H describing the scaling behaviour of the traffic. A small number of

traffic parameters is a very desirable feature from the point of view of the applicability of the model for traffic engineering purposes. The estimation of even a small number of parameters poses a problem for long range dependent traffic. Some early work [7] suggested that to obtain a reasonable accuracy a very large number of sample points may be required. The problem arises e.g. in the estimation of the Hurst parameter H . As H describes the scaling behaviour of the traffic variability, the sample points have to cover several time scales in order to determine H reliably, i.e., the total time range must be several orders of magnitude greater than the finest time resolution in the measurement.

In this paper we show that by an appropriate choice of the sampling instants, the number of sampling points can be considerably reduced. In particular, we will introduce a grid of geometrically distributed sampling points $t_i = \alpha^{i-1}$, $i = 1, \dots, n$ where α is some constant (< 1). The idea here is that such sampling grid covers several time scales with fewer points. The second point is that the geometrical grid, being “self-similar” fits well with the traffic process and gives rise to a simple structure in the covariance matrix.

Throughout this work we apply the maximum likelihood estimation (MLE) method [1]. MLE method has previously been applied to this problem by Deriche and Tewfik [2] and Ninness [5] using ordinary linear sampling. Explicit formulas for the estimators of m and a are given along with the log-likelihood function for determining the estimator for H . A major difficulty in this method is the calculation of the inverse and determinant of the covariance matrix appearing in the likelihood function. An approximate calculation is facilitated if the process is stationary whence the matrix is of Toeplitz type. For the original fBm process the increment process is stationary. We show that another stationary process is obtained from the fBm process by ‘descaling’ and changing the process’ index to logarithmic time, i.e., on the geometrical sampling grid the descaled process is stationary. It turns out that the elements of the inverse matrix far from the diagonal are small, enabling us to derive a simple approximation for the inverse matrix directly without using e.g. Whittle’s method [1] based on the spectral analysis.

We compare the effectiveness of the MLE estimator based on ordinary evenly spaced sampling grid with that obtained with a geometrical grid by simulations. In the comparison the total time range covered by the sample points was kept constant while the total range is always chosen to be $[0, 1]$ (the actual upper limit is immaterial because of the self-similarity). The simulations indicate that in the estimation of H the geometrical grid gives an advantage, whereas for the estimation of a alone it still gives satisfactory results although the linear sampling is slightly better. The estimation of mean rate m depends very little on the method and is always essentially given by the total traffic arrived between the first and last sample points.

The rest of this paper is organized as follows. In section 2 we review the fractional Brownian motion traffic model with its three parameters. The general problem of the estimation of these parameters by the maximum likelihood method is considered in section 3. The idea of geometrical sampling and the descaled process, along with an approximate form of the MLE, are introduced in section 4. For comparison, in section 5 we present the MLE method for the case of ordinary linear sampling. In section 6, we present results for estimating the fBm parameters with the described methods from simulated realizations of the process. Section 7 concludes the paper.

2 Fractional Brownian traffic

A normalized *fractional Brownian motion* with Hurst-parameter $H \in [0.5, 1)$, denoted by $Z(t)$, ($t \in \mathbb{R}$), is characterized by the following properties [6]:

1. $Z(t)$ has stationary increments;
2. $Z(0) = 0$, and $\mathbb{E}[Z(t)] = 0$ for all t ;
3. $\text{Var}[Z(t)] = \mathbb{E}[Z(t)^2] = |t|^{2H}$ for all t ;
4. $Z(t)$ has continuous paths;
5. $Z(t)$ is a Gaussian process, i.e., all its finite-dimensional marginal distributions are Gaussian.

In the special case $H = 0.5$, $Z(t)$ is the standard Brownian motion. It follows from the above properties that $Z(t)$ is a self-similar process whose scaling behaviour is defined by the Hurst-parameter H as follows

$$Z(\alpha t) \sim \alpha^H Z(t). \quad (1)$$

The covariance structure of the process is given by

$$\text{Cov}[Z(t_1), Z(t_2)] = \frac{1}{2} \{t_1^{2H} + t_2^{2H} - |t_2 - t_1|^{2H}\}. \quad (2)$$

Furthermore, in the case $H > 0.5$ the strongly correlated stationary sequence $Z(n+1) - Z(n)$, the increment process of $Z(t)$, (often called *fractional Gaussian noise*) is ergodic [6].

Fractional Brownian motion is a popular model for long-range dependent traffic. Norros [6] has suggested the following model

$$X(t) = mt + \sqrt{a}Z(t), \quad (3)$$

where X_t represents the amount of traffic arrived in $(0, t)$. The model has three parameters, m , a and H with the following interpretations and intervals for allowed values: $m > 0$ is the mean input rate, $a > 0$ is a variance parameter, and $H \in [0.5, 1)$ is the self-similarity parameter of $Z(t)$.

3 Exact Gaussian MLE

We use the notation of Beran [1]. Assume the traffic has been observed at n time instants forming the vector $\mathbf{t} = (t_1, \dots, t_n)^t$ where $(\cdot)^t$ denotes the transpose. And let $\mathbf{X} = (X(t_1), \dots, X(t_n))^t$ be the vector of observed traffic values at these instants. Since $X(t)$ is Gaussian, the joint probability density function of \mathbf{X} is

$$h(\mathbf{x}) = (2\pi)^{-\frac{n}{2}} |\mathbf{\Gamma}|^{-\frac{1}{2}} e^{-\frac{1}{2}(\mathbf{x}-\mathbf{m})^t \mathbf{\Gamma}^{-1}(\mathbf{x}-\mathbf{m})}, \quad (4)$$

where $\mathbf{x} = (x_1, \dots, x_n)^t \in \mathbb{R}^n$, $\mathbf{m} = m\mathbf{t}$, and $|\mathbf{\Gamma}|$ is the determinant of the covariance matrix

$$\mathbf{\Gamma} = \text{Cov}[\mathbf{X}, \mathbf{X}^t] = \text{E}[\mathbf{X}\mathbf{X}^t] - \text{E}[\mathbf{X}]\text{E}[\mathbf{X}^t]. \quad (5)$$

The MLE for m is obtained by maximizing $\log h(\mathbf{X}; m)$ with respect to m , resulting in the estimator

$$\hat{m} = \hat{m}(H) = \frac{\mathbf{t}^t \mathbf{\Gamma}^{-1} \mathbf{X}}{\mathbf{t}^t \mathbf{\Gamma}^{-1} \mathbf{t}}. \quad (6)$$

Note, that the estimate is unbiased, irrespective whether our estimate for H is correct or not. The variance of \hat{m} can also be calculated with the assumption that H is known exactly, $\hat{H} = H$. With straightforward calculations we get

$$\text{Var}[\hat{m}] = \frac{a}{\mathbf{t}^t \mathbf{\Gamma}^{-1} \mathbf{t}} \quad (7)$$

The variance of our estimator is smaller than the estimator based on the total sample mean, by the factor in the denominator (which is close to 1).

Next, consider the estimator for a . $\mathbf{\Gamma}$ is a simple linear function of a , $\mathbf{\Gamma} = a \mathbf{\Gamma}_H$, where $\mathbf{\Gamma}_H$ is independent of a and is given by

$$\mathbf{\Gamma}_H = \text{E}[\mathbf{Z}\mathbf{Z}^t] = [\text{Cov}[Z(t_i), Z(t_j)]]_{i,j=1,\dots,n}. \quad (8)$$

The MLE of a is obtained by maximizing the log-likelihood function $\log h(\mathbf{X}; a)$ with respect to a , and from that we get

$$\hat{a} = \hat{a}(m, H) = \frac{1}{n} (\mathbf{X} - \mathbf{m})^t \mathbf{\Gamma}_H^{-1} (\mathbf{X} - \mathbf{m}). \quad (9)$$

If we don't know the mean input rate m in advance, \mathbf{m} in Eq.(9) should be replaced by $\hat{m}\mathbf{t}$. Using Eq.(6) and Eq.(9) we get

$$\hat{a}(H) = \frac{1}{n} \frac{(\mathbf{X}^t \mathbf{\Gamma}_H^{-1} \mathbf{X})(\mathbf{t}^t \mathbf{\Gamma}_H^{-1} \mathbf{t}) - (\mathbf{t}^t \mathbf{\Gamma}_H^{-1} \mathbf{X})^2}{\mathbf{t}^t \mathbf{\Gamma}_H^{-1} \mathbf{t}}. \quad (10)$$

Again, assuming for the time being that H is known correctly the expectation and variance of \hat{a} can be calculated. Using $\text{E}[\mathbf{Z}^t \mathbf{\Gamma}_H^{-1} \mathbf{Z}] = \text{E}[\mathbf{N}^t \mathbf{N}] = n$ since $\mathbf{Z} \sim \mathbf{\Gamma}_H^{1/2} \mathbf{N}$ where \mathbf{N} is a vector of independent standard Gaussian variables, finally we have

$$\text{E}[\hat{a}] = \frac{n-1}{n} a, \quad \text{Var}\left[\frac{n}{n-1} \hat{a}\right] = \frac{2a^2(n-1)}{n^2}. \quad (11)$$

Thus \hat{a} has the "normal" $(n-1)/n$ bias.

Finally, we are left with the maximization of the H -dependent part of the log-likelihood function, i.e., essentially we have to minimize

$$\tilde{L}(\mathbf{X}; H) = \log |\mathbf{\Gamma}_H| + n \log \frac{(\mathbf{X}^t \mathbf{\Gamma}_H^{-1} \mathbf{X})(\mathbf{t}^t \mathbf{\Gamma}_H^{-1} \mathbf{t}) - (\mathbf{t}^t \mathbf{\Gamma}_H^{-1} \mathbf{X})^2}{\mathbf{t}^t \mathbf{\Gamma}_H^{-1} \mathbf{t}}. \quad (12)$$

The first term is a decreasing function of H , and the second term is an increasing function of H . The minimum is obtained for some value \hat{H} which is the MLE estimate; the corresponding MLE estimates for m and a are $\hat{m} = m(\hat{H})$ and $\hat{a} = a(\hat{H})$.

4 Geometrical sampling

The Hurst parameter H describes the scaling behaviour of the traffic. Therefore, in order to determine its value from measured traffic, the sample points have to cover several time scales, i.e. the total time range of the measurements has to be many orders of magnitude greater than the finest resolution (smallest interval between the sampling points). With the ordinary linear sampling, i.e. sampling points at constant intervals, this leads to the requirement of very large number of sampling points. Obviously, because of the correlations, there is a lot of redundancy in measured traffic values at these points. In order to use the measurements more efficiently we introduce a geometric sequence of sampling points, $t_i = \alpha^i$, $i = 1, \dots, n$, with some $0 < \alpha < 1$. The vector of observed traffic values at the sampling points is denoted by $\mathbf{X} = (X(t_1), X(t_2), \dots, X(t_n))^t$.

In addition to distributing the sampling points in a better way on different time scales, geometric sampling fits neatly with the self-similar behaviour of the fBm traffic. We show first that by a simple transformation we can obtain from the fBm process another process which is a stationary process of logarithmic time. As a geometric sequence corresponds to equidistant points in logarithmic time, it follows that the samples of the modified process constitute a stationary sequence. This leads to a simple Toeplitz-type structure of the covariance matrix and allows us to develop approximations to the inverse and determinant of the covariance matrix.

4.1 Descaled process

$Z(t)$ has the self-similar property $Z(\alpha t) \sim \alpha^H Z(t)$. Now consider the ‘descaled’ process $\check{Z}(t) \stackrel{d}{=} t^{-H} Z(t)$ which has the scaling property

$$\check{Z}(\alpha t) \sim (\alpha t)^{-H} Z(\alpha t) = t^{-H} Z(t) = \check{Z}(t). \quad (13)$$

Further let us take a new time variable $u = -\log t$ and denote $\tilde{Z}(u) \stackrel{d}{=} \check{Z}(e^{-u}) = \check{Z}(t)$. Now we have

$$\tilde{Z}(u - \log \alpha) = \check{Z}(e^{-u+\log \alpha}) = \check{Z}(\alpha e^{-u}) = \check{Z}(\alpha t) \sim \check{Z}(t) = \tilde{Z}(u). \quad (14)$$

Thus the process $\tilde{Z}(u)$ is stationary and has the following covariance structure:

$$\text{Cov} [\tilde{Z}(u_1), \tilde{Z}(u_2)] = \frac{1}{2} e^{H(u_2-u_1)} \left\{ 1 + e^{-2H(u_2-u_1)} - \left(1 - e^{-(u_2-u_1)} \right)^{2H} \right\}, \quad (15)$$

so the descaled process $\tilde{Z}(u)$ is short range dependent.

If we ‘descale’ the process $X(t)$ by the factor t^{-H} and use u as the process index, we finally have

$$\tilde{X}(u) \stackrel{d}{=} m e^{(H-1)u} + \sqrt{a} \tilde{Z}(u). \quad (16)$$

The covariance matrix $\tilde{\Gamma}$ of the descaled samples $\tilde{\mathbf{X}} = (\tilde{X}(u_1), \tilde{X}(u_2), \dots, \tilde{X}(u_n))^t$ with $u_i = -\log t_i = (1-i) \log \alpha$ can be written as

$$\tilde{\Gamma} = \text{E} [\tilde{\mathbf{X}} \tilde{\mathbf{X}}^t] = a \cdot \text{E} [\tilde{\mathbf{Z}} \tilde{\mathbf{Z}}^t]. \quad (17)$$

Note, that our geometrical grid is now equally spaced with regard to u . Thus, if we use the notation $\tilde{Z}_i = \tilde{Z}(u_i)$ the process $\tilde{\mathbf{Z}} = (\tilde{Z}_1, \tilde{Z}_2, \dots, \tilde{Z}_n)$ is a stationary process in discrete time with zero mean and unit variance and its auto-correlation function $\rho(k)$ can be defined as

$$\rho(i-j) = \frac{1}{2} \alpha^{-H|i-j|} \left\{ 1 + \alpha^{2H|i-j|} - (1 - \alpha^{|i-j|})^{2H} \right\}. \quad (18)$$

and thus

$$\tilde{\Gamma}_{ij} = a\rho(i-j), \quad i, j = 1, 2, \dots, n. \quad (19)$$

4.2 Descaled MLE

When doing the maximum likelihood estimation of the model parameters m , a and H , one can utilize the stationarity and short range dependent properties of the descaled process. Using the ‘descaling matrix’ $\mathbf{D} = \text{diag}(t_1^{-H}, \dots, t_n^{-H})$ we can easily derive $\tilde{\Gamma} = \mathbf{D} \Gamma \mathbf{D}$, and from this we get

$$\Gamma_H^{-1} = \mathbf{D} \tilde{\Gamma}_H^{-1} \mathbf{D}. \quad (20)$$

The determinant $|\Gamma_H|$ can be also calculated as

$$|\Gamma_H| = \alpha^{Hn(n-1)} |\tilde{\Gamma}_H|. \quad (21)$$

4.3 Approximate MLE

In practice, the exact MLE poses computational problems. And this is not just because of the computation time needed in case of large data sets, but because of the evaluation of the inverse and the determinant of the covariance matrix may be numerically unstable. To avoid these problems, one can use approximate methods to calculate the estimates. In [1], several possible approaches to approximating the Gaussian likelihood function are discussed, among them the well known Whittle’s approximate MLE.

In our case we focus on the properties of the covariance matrix Γ_H , trying to take advantage of its special structure and to find efficient approximations for its inverse and determinant.

Using the notations $\beta = \alpha^{-H}$ and

$$g(x) = \frac{1}{2} (1 + x^{2H} - (1-x)^{2H}), \quad (22)$$

the elements of the autocorrelation matrix $\tilde{\Gamma}_H$ can be written as $(\tilde{\Gamma}_H)_{i,j} = \beta^{|i-j|} g(\alpha^{|i-j|})$, $i, j = 1, 2, \dots, n$. It is interesting to note, that $g(x)$ is nearly completely linear for $x \in (0, 1)$. Figure 1 shows the difference of $g(x) - x$ for different values of H . It can be seen from the plot that the largest absolute difference is less than 0.02 for each value of H . This observation gives us the idea to use the approximation $g(x) \approx x$. So $\tilde{\Gamma}_H$ can be approximated as $\tilde{\Gamma}_H \approx \mathbf{R}$, where \mathbf{R} is a Toeplitz-type matrix of the form $[\mathbf{R}]_{ij} = \gamma^{|i-j|}$, $i, j = 1, 2, \dots, n$, with $\gamma = \alpha^{1-H}$.

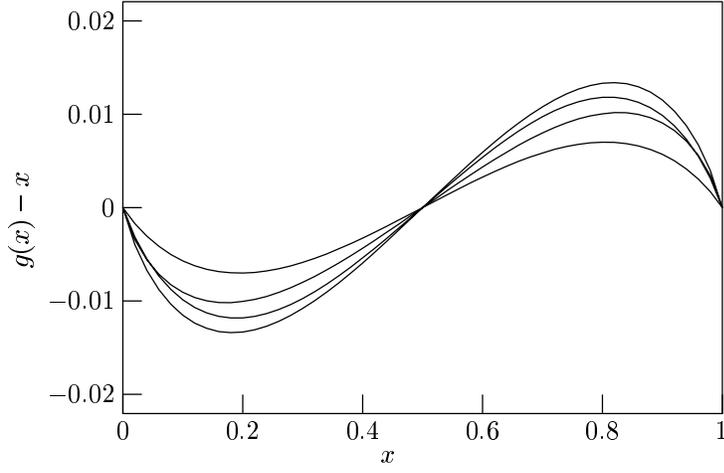


Figure 1: Error of approximation $g(x) \approx x$ for $H = 0.6, 0.7, 0.8$ and 0.9 .

The inverse of \mathbf{R} can be easily calculated as [8]

$$\mathbf{R}^{-1} = \frac{1}{\frac{1}{\gamma} - \gamma} \begin{pmatrix} \frac{1}{\gamma} & -1 & 0 & \cdots & 0 \\ -1 & \gamma + \frac{1}{\gamma} & -1 & \ddots & \vdots \\ 0 & -1 & \gamma + \frac{1}{\gamma} & \ddots & 0 \\ \vdots & \ddots & \ddots & \ddots & -1 \\ 0 & \cdots & 0 & -1 & \frac{1}{\gamma} \end{pmatrix}, \quad (23)$$

and the determinant of \mathbf{R} is given by [8]

$$|\mathbf{R}| = (-1)^{n-1} \prod_{i=1}^{n-1} \begin{vmatrix} \gamma^{1-i} & \gamma^{i-1} \\ \gamma^{-i} & \gamma^i \end{vmatrix} \cdot \gamma^{n-1} = (1 - \gamma^2)^{n-1}. \quad (24)$$

Using the fact that $\mathbf{t}^t \mathbf{D} \mathbf{R}^{-1} \mathbf{D} \mathbf{t} = 1$ and $\mathbf{t}^t \mathbf{D} \mathbf{R}^{-1} \mathbf{D} = (1, 0, \dots, 0)$, we get

$$\hat{m}(H) = X_1, \quad (25)$$

so using the above approximation the MLE estimate for m reduces simply to the sample mean. As for the estimate for a we get

$$\hat{a}(H) = \frac{1}{n} (\mathbf{X}^t \mathbf{D} \mathbf{R}^{-1} \mathbf{D} \mathbf{X} - X_1^2). \quad (26)$$

Finally, to get an estimate for H we have to minimize the function

$$L(\mathbf{X}; H) = \frac{n-1}{n} \log(\alpha^{nH} (1 - \alpha^{2-2H})) + \log(\mathbf{X}^t \mathbf{D} \mathbf{R}^{-1} \mathbf{D} \mathbf{X} - X_1^2). \quad (27)$$

It should be noted that though the linear approximation to $g(x)$ is rather accurate, the resulting inverse matrix \mathbf{R}^{-1} of Eq.(23) is rather poor an approximation to $\tilde{\mathbf{\Gamma}}^{-1}$ for large n . Nevertheless, the use of \mathbf{R}^{-1} in the log-likelihood function (27), as we will see, yields a good estimate for H , while the accuracy of the estimate $\hat{a}(H)$ suffers more from this approximation.

4.4 Improved approximation for $\mathbf{\Gamma}^{-1}$

Since the matrix $\tilde{\mathbf{\Gamma}}$ is a Toeplitz-type matrix with decreasing elements as we go farther from the diagonal, we expect that its inverse can be well approximated with a band matrix whose elements are zero starting from a given distance from the diagonal. Let $\tilde{\mathbf{\Gamma}}_H^{-1} \approx \mathbf{C}$ so that the approximate inverse has the following structure:

$$\mathbf{C} = \begin{pmatrix} c_1 & c_2 & \cdots & c_p & 0 & \cdots & 0 \\ c_2 & c_1 & c_2 & \cdots & c_p & \ddots & \vdots \\ \vdots & c_2 & c_1 & c_2 & \vdots & \ddots & 0 \\ c_p & \vdots & c_2 & c_1 & c_2 & \cdots & c_p \\ 0 & c_p & \cdots & c_2 & c_1 & \ddots & \vdots \\ \vdots & \ddots & \ddots & \vdots & \ddots & \ddots & c_2 \\ 0 & \cdots & 0 & c_p & \cdots & c_2 & c_1 \end{pmatrix}. \quad (28)$$

Our aim is to set the p parameters c_1, \dots, c_p to get $\mathbf{C}\tilde{\mathbf{\Gamma}}_H \approx \mathbf{E}$. For example, this can be achieved by solving the equation

$$(c_p, \dots, c_2, c_1, c_2, \dots, c_p) \cdot \mathbf{G} = (0, \dots, 0, 1, 0, \dots, 0), \quad (29)$$

where $\mathbf{G} = (\tilde{\mathbf{\Gamma}}_H)_{(2p-1) \times (2p-1)}$ and from this we have

$$c_i = \mathbf{G}_{p(p+i-1)}^{-1}, \quad i = 1, 2, \dots, p. \quad (30)$$

With this approximation we only need to calculate the inverse of a $(2p-1)$ -by- $(2p-1)$ matrix.¹ To improve the approximate inverse, its elements in the upper-left and lower-right corners can be corrected, for example, by solving

$$(\hat{c}_{11}, \hat{c}_{12}, \dots, \hat{c}_{1(p-1)}, c_{1p}) \cdot (\tilde{\mathbf{\Gamma}}_H)_{(p-1) \times p} = (1, 0, \dots, 0), \quad (31)$$

and

$$(\hat{c}_{21}, \hat{c}_{22}, c_{23}, \dots, c_{2p}) \cdot (\tilde{\mathbf{\Gamma}}_H)_{2 \times p} = (0, 1). \quad (32)$$

And because of the symmetric structure, $c_{ij} = c_{ji} = c_{(n-i+1)(n-j+1)}$, $i, j = 1, \dots, n$.

¹To be more exact, because of the symmetric structure we only need to calculate the inverse of a p -by- p matrix using slightly more complicated formulas.

5 Linear sampling

Let $\mathbf{X} = (X(t_1), X(t_2), \dots, X(t_n))^t$ be the vector of observed traffic values at instances

$$t_i = \frac{i}{n}, \quad i = 1, 2, \dots, n. \quad (33)$$

The increment sequence (Y_1, Y_2, \dots) with $Y_i = X(t_i) - X(t_{i-1})$ (substituting $X(t_0) \equiv X(0) = 0$) is a strongly correlated stationary sequence with

$$\text{Cov}[Y_i, Y_j] = \frac{1}{2} a n^{-2H} \left(|i - j + 1|^{2H} + |i - j - 1|^{2H} - 2|i - j|^{2H} \right), \quad i, j = 1, 2, \dots, n. \quad (34)$$

The formulas for the exact Gaussian MLE for this increment process are nearly the same as in Section 3, we only need to replace the covariance matrix $\mathbf{\Gamma}$ with $\mathbf{\Sigma} = [\text{Cov}[Y_i, Y_j]]_{i,j=1,2,\dots,n}$, and the vector \mathbf{t} with the vector $(1/n, 1/n, \dots, 1/n)^t$. After some minor simplifications we get an estimate for m

$$\hat{m} = \hat{m}(H) = \frac{\mathbf{1}^t \mathbf{\Sigma}^{-1} \mathbf{Y}}{\mathbf{1}^t \mathbf{\Sigma}^{-1} \mathbf{1}} \cdot n \quad (35)$$

where $\mathbf{1}$ is a vector of ones, and $\mathbf{\Sigma} = a \mathbf{\Sigma}_H$. For a we have the estimator

$$\hat{a}(H) = \frac{1}{n} \left(\mathbf{Y}^t \mathbf{\Sigma}_H^{-1} \mathbf{Y} - \frac{(\mathbf{1}^t \mathbf{\Sigma}_H^{-1} \mathbf{Y})^2}{\mathbf{1}^t \mathbf{\Sigma}_H^{-1} \mathbf{1}} \right). \quad (36)$$

Again, finally we have to minimize

$$\tilde{L}(\mathbf{Y}; H) = \log |\mathbf{\Sigma}_H| + n \log \left(\mathbf{Y}^t \mathbf{\Sigma}_H^{-1} \mathbf{Y} - \frac{(\mathbf{1}^t \mathbf{\Sigma}_H^{-1} \mathbf{Y})^2}{\mathbf{1}^t \mathbf{\Sigma}_H^{-1} \mathbf{1}} \right). \quad (37)$$

The minimum is obtained for some value \hat{H} which is the MLE estimate.

However, to calculate the inverse and the determinant of $\mathbf{\Sigma}_H$ the same problems arise as in the case of geometrical sampling with the covariant matrix $\tilde{\mathbf{\Gamma}}_H$. Since $\mathbf{\Sigma}_H$ is also a Toeplitz type matrix, the same method as described in section 4.4 can be used to approximate $\mathbf{\Sigma}_H^{-1}$ with \mathbf{C} of Eq.(28). The only difference is that we need the determinant $|\mathbf{\Sigma}_H|$, or at least an approximation of it, to calculate the estimate of H using Eq.(37). Fortunately, the determinant of \mathbf{C} can be calculated (see Appendix A), and we get $|\mathbf{\Sigma}_H| \approx |\mathbf{C}|^{-1}$.

6 Simulation results

The fBm samples were generated using the fact $\mathbf{Z} \sim \mathbf{\Gamma}_H^{1/2} \mathbf{N}$ (or, correspondingly, $\mathbf{Z} \sim \mathbf{\Sigma}_H^{1/2} \mathbf{N}$ for the linear sampling) where \mathbf{N} is a vector of independent standard Gaussian variables.

In all the simulation presented here the model parameters were set as $m = 1$, $a = 1$ and $H = 0.8$ as an example, but similar results were obtained using different values

of the parameters. The parameter α for the geometrical grid was chosen so that the difference between the nearest two measurement time instants (the ‘resolution’ of the measurement) was 10^{-6} . Figure 2 shows the results of H estimates as a function of the number of sample points using both geometrical and linear sampling. In the geometrical case Eq.(27) was minimized while for the linear sampling we used the formula Eq.(37) where the inverse of Σ_H was approximated with a band matrix of Eq.(28) with $p = 2$ and the determinant of the correlation matrix was approximated by Eq.(44). Assuming Gaussianity, the 95% confidence interval was obtained by repeating the simulations 100 times and calculating the sample variance of the estimates. The results show that the

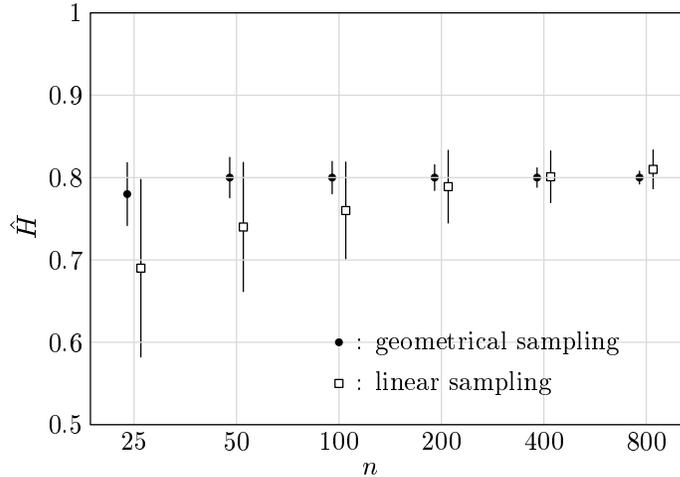


Figure 2: Estimates of H using geometrical and linear sampling.

estimates using geometrical sampling have much smaller variance and are unbiased for sample sizes larger than 25. As for the linear sampling, the bias is considerably larger although the estimates converge to the proper value of H as the sample size increases. However, the variance of the estimates is always higher than in the geometrical case. For example, the variance for 800 samples using linear sampling is nearly the same as for only 50 geometrically sampled points.

The next question was how the two different sampling methods affect the estimates for the variance parameter a . Figure 3 displays the results, assuming that H is known (or, equivalently, can be estimated exactly). These simulations were useful to test whether our approximations in calculating the inverse and determinant of the covariance matrices are adequate or not. Figure 2 presents two different approximations for the geometrical sampling. First, we used the simple approximate inverse covariance matrix of Eq.(23) in Eq.(10) using Eq.(20) (denoted by light gray dots and labeled ‘linear approximation’ in the figure). As can be seen, the estimates of a are strongly biased and the bias is getting larger as the number of samples increases. So this estimate is clearly inadequate, the approximation of Eq.(23) had to be refined. Next, we used the approximation of Eq.(28) for $\tilde{\Gamma}_H^{-1}$ with five parameters ($p = 5$). As we see from Figure 2, the strong bias from the \hat{a} estimates disappeared and the variance of the estimates is only slightly higher than the

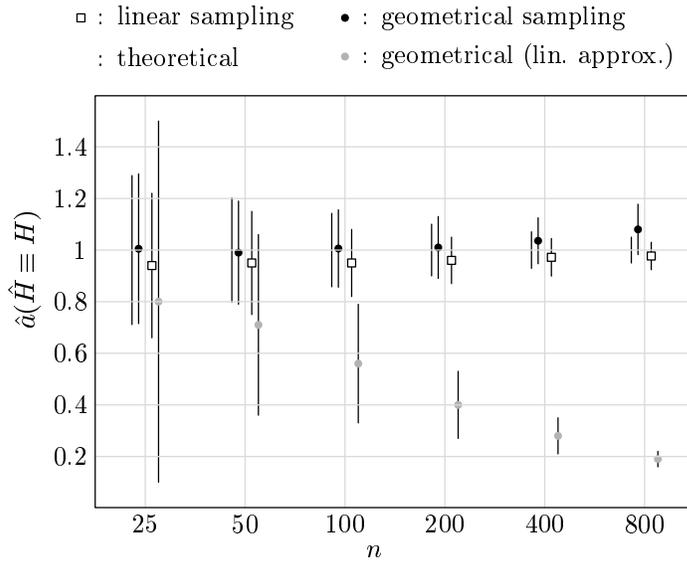


Figure 3: Estimates of a (when H is also estimated) using geometrical and linear sampling and different approximations, assuming H is known.

theoretical value that can be calculated using Eq.(11). (Note, however, that the bias for sample sizes of 400 and 800 seems to be slightly increased.) Finally, the linear sampling method was used. Its estimates are asymptotically unbiased and have approximately the same variance as expected. The approximate inverse matrix used was as in Eq.(28) with only two parameters ($p = 2$).

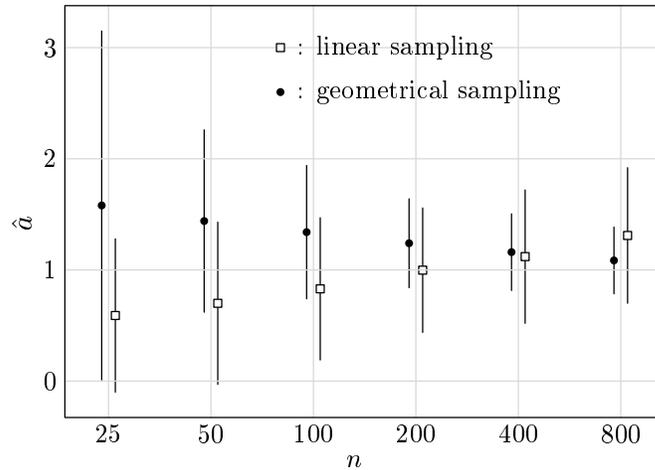


Figure 4: Estimates of a using geometrical and linear sampling and different approximations.

Figure 4 shows the MLE \hat{a} estimates without any *a priori* knowledge about the model parameters. All the approximations used here were the same as in the previous cases.

Since H is not known and can only be estimated with a given variance, the estimates of a have larger variances than in the previous simulations. The question is how robust those estimates are when \hat{H} can have a slight bias (see Figure 2). As for the geometrical sampling, the bias of \hat{a} gets smaller and its variance is also decreasing rapidly as the sample size increases. On the other hand, for the linear sampling case the estimates seem to be biased for larger sample sizes and their variance does not seem to decrease. The reason for this behaviour lies in the fact that the linear sampling for estimating H is less accurate than the geometrical sampling. The bias in \hat{H} together with its higher variance is responsible for the bias and variance of \hat{a} , even if the linear sampling seems to be a better choice to estimate a than the geometrical one for known H (see Figure 3).

As for the MLE estimates for m the geometrical sampling does not give any extra advantage or disadvantage compared to the linear sampling. In fact, the MLE estimate gives almost negligible reduction in the variance of \hat{m} when compared to the sample mean as an estimate for m , i.e., $X(1)$ in our case that corresponds to the last sample in both geometrical and linear sampling [1].

7 Conclusion

In this paper we have introduced the idea of using geometrical sampling for the ML estimation of the parameters of fractional Brownian traffic. The intention with this sampling is to reduce the number of sampling points required for a given predefined confidence level. Intuitively, the geometrical sampling distributes the sampling points advantageously at different time scales, whereas linear sampling stresses the finest time scale and contains redundant information.

We have derived expressions for the estimators of m and a and the log-likelihood function from which the estimator of H can be derived, both for the linear sampling and the geometrical sampling. Approximations were developed for the inverse and the determinant of the covariance matrix, needed for the calculation of the estimates. With these approximations the evaluation of the log-likelihood function is fast and the maximization with respect to H can easily be made. In this paper, however, we did not specifically address the question of what is the best numerical way of doing the maximization.

The experiments with simulated traffic showed that the geometrical sampling does indeed give a better estimate for H leading to a reduction of sample points. In one example the number of required points was reduced from 800 to 50. For the estimation of a the geometrical sampling does not give any direct advantage, with a known H , but as the estimator \hat{a} actually depends on the estimator \hat{H} , the overall accuracy obtained with geometrical sampling is better. For the estimation of m , different sampling schemes give essentially the same result, i.e., the estimate is basically the observed average rate. It should, however, be noted that the experiments were made only with simulated traffic with exact ‘measurements’. If the measured values are noisy then the descaling factor may amplify the noise of the points near the origin.

Though the geometrical sampling has been shown to give better results than the linear sampling, it is not claimed that it constitutes the optimal sampling scheme. So

there remains the theoretical question what is the best way of locating a given number of sampling points in the interval $[0, 1]$ with the constraint that the smallest distance between any pair of points is greater than or equal to a given minimum resolution.

A Appendix

Consider a symmetrically partitioned hypermatrix of the second order (see Eq.(38)) where the diagonal submatrices $\mathbf{C}_{q \times q}$ and $\mathbf{B}_{r \times r}$ are square. Assume further, that we know its inverse which can be similarly partitioned, i.e., $\mathbf{S}_{r \times r}$ and $\mathbf{L}_{q \times q}$ are square.

$$\begin{array}{c} r) \\ q) \end{array} \begin{array}{|c|c|} \hline \mathbf{A} & \mathbf{B} \\ \hline \mathbf{C} & \mathbf{D} \\ \hline \end{array}^{-1} = \begin{array}{|c|c|} \hline \mathbf{U} & \mathbf{L} \\ \hline \mathbf{S} & \mathbf{V} \\ \hline \end{array} \begin{array}{l} (q) \\ (r) \end{array} \quad (38)$$

The determinant of the \mathbf{C} submatrix can be calculated if we know the determinant of the hypermatrix as well as the determinant of the lower-left submatrix of the inverse:

$$|\mathbf{C}| = (-1)^{rq} \begin{vmatrix} \mathbf{A} & \mathbf{B} \\ \mathbf{C} & \mathbf{D} \end{vmatrix} \cdot |\mathbf{S}| \quad (39)$$

Our aim is to calculate the determinant of the correlation matrix. An approximate of this determinant can be given as the reciprocal of the determinant of the approximate inverse \mathbf{C} of the form Eq.(28). Since \mathbf{C} is a band matrix, the idea is to construct the hypermatrix in Eq.(38) to be a lower-triangular matrix, and as a result, its inverse will be also lower-triangular and its determinant can be easily calculated.

In the following we consider the case $p = 2$ when the matrix elements only in the diagonal and co-diagonals are nonzero. We do it just for simplicity, but the similar method can be used for different values of p . For $p = 2$, \mathbf{C} is symmetric and co-diagonal:

$$\mathbf{C} = \begin{pmatrix} a_1 & b_1 & 0 & \cdots & 0 \\ b_1 & a_2 & b_2 & \ddots & \vdots \\ 0 & b_2 & a_3 & \ddots & 0 \\ \vdots & \ddots & \ddots & \ddots & b_{n-1} \\ 0 & \cdots & 0 & b_{n-1} & a_n \end{pmatrix} \quad (40)$$

with $b_i \neq 0$. Note, that here we let the elements vary in the diagonal lines so this assumption makes it possible to incorporate any refinements such as the corner correction used in section 4.4. Adding one extra row to the top and one extra column to the right,

we get:

$$\left(\begin{array}{ccccc|c} 1 & 0 & 0 & \cdots & 0 & 0 \\ a_1 & b_1 & 0 & \cdots & 0 & 0 \\ b_1 & a_2 & b_2 & \ddots & \vdots & \vdots \\ 0 & b_2 & a_3 & \ddots & 0 & 0 \\ \vdots & \ddots & \ddots & \ddots & b_{n-1} & 0 \\ 0 & \cdots & 0 & b_{n-1} & a_n & 1 \end{array} \right)^{-1} = \left(\begin{array}{c|cccc} u_1 & 0 & \cdots & 0 & 0 \\ u_2 & l_{21} & \ddots & \vdots & \vdots \\ \vdots & \vdots & \ddots & 0 & 0 \\ u_n & l_{n1} & \cdots & l_{n(n-1)} & 0 \\ \hline s & v_1 & \cdots & v_{n-1} & v_n \end{array} \right) \quad (41)$$

(For the general case p extra rows and columns should be added.) To determine s (or $\mathbf{S}_{(p-1) \times (p-1)}$ in the general case) we need to solve

$$\begin{pmatrix} \mathbf{e}_1^t & 0 \\ \mathbf{C} & \mathbf{e}_n \end{pmatrix} \cdot \begin{pmatrix} \mathbf{u} \\ s \end{pmatrix} = \begin{pmatrix} \mathbf{e}_1 \\ 0 \end{pmatrix}, \quad (42)$$

where \mathbf{e}_i denotes a vector of length n with all its elements being zero except the i th which is 1. To do this, s can be calculated using the following recursive equations:

$$\begin{aligned} u_1 &= 1, \\ u_2 &= -\frac{1}{b_1} a_1, \\ u_{i+1} &= -\frac{1}{b_i} (a_i u_i + b_{i-1} u_{i-1}), \quad i = 2, 3, \dots, n-1, \\ s &= a_n u_n + b_{n-1} u_{n-1}. \end{aligned} \quad (43)$$

Since the determinant of a triangular matrix is equal to the products of its diagonal elements, the determinant of \mathbf{C} can be calculated as

$$|\mathbf{C}| = (-1)^n \cdot s \cdot \prod_i^{n-1} b_i. \quad (44)$$

(Again, in the general case the determinant $|\mathbf{S}|$ has to be calculated and used in the formula instead of s .)

Finally, we should note that only in the case $p = 2$ and for the case when \mathbf{C} approximates the inverse of $\tilde{\mathbf{\Gamma}}_H$ the determinant can be accurately approximated by the rather simple formula

$$|\mathbf{C}| \approx \prod_i^{n-1} |b_i|. \quad (45)$$

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