A practical method for calculating largest Lyapunov exponents from small data sets

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Abstract

Detecting the presence of chaos in a dynamical system is an important problem that is solved by measuring the largest Lyapunov exponent. Lyapunov exponents quantify the exponential divergence of initially close state-space trajectories and estimate the amount of chaos in a system. We present a new method for calculating the largest Lyapunov exponent from an experimental time series. The method follows directly from the definition of the largest Lyapunov exponent and is accurate because it takes advantage of all the available data. We show that the algorithm is fast, easy to implement, and robust to changes in the following quantities: embedding dimension, size of data set, reconstruction delay, and noise level. Furthermore, one may use the algorithm to calculate simultaneously the correlation dimension. Thus, one sequence of computations will yield an estimate of both the level of chaos and the system complexity.

1. Introduction

Over the past decade, distinguishing deterministic chaos from noise has become an important problem in many diverse fields, e.g., physiology [18], economics [11]. This is due, in part, to the availability of numerical algorithms for quantifying chaos using experimental time series. In particular, methods exist for calculating correlation dimension (D_2) [20], Kolmogorov entropy [21], and Lyapunov characteristic exponents [15, 17, 32, 39]. Dimension gives an estimate of the system complexity; entropy and characteristic exponents give an estimate of the level of chaos in the dynamical system.

The Grassberger-Procaccia algorithm (GPA) [20] appears to be the most popular method used to quantify chaos. This is probably due to the simplicity of the algorithm [16] and the fact that the same intermediate calculations are used to estimate both dimension and entropy. However, the GPA is sensitive to variations in its parameters, e.g., number of data points [28], embedding dimension [28], reconstruction delay [3], and it is usually unreliable except for long, noise-free time series. Hence, the practical significance of the GPA is questionable, and the Lyapunov exponents may provide a more useful characterization of chaotic systems.

For time series produced by dynamical systems, the presence of a positive characteristic exponent indicates chaos. Furthermore, in many applications it is sufficient to calculate only the largest Lyapunov exponent (λ_1). However, the existing methods for estimating λ_1 suffer from at least one of the following drawbacks: (1) unreliable for small data sets, (2) computationally intensive, (3) relatively difficult to implement. For this reason, we have developed a new method for calculating the largest Lyapunov exponent. The method is reliable for small data sets, fast, and easy to implement. "Easy to implement" is largely a subjective quality, although we believe it has had a notable positive effect on the popularity of dimension estimates.

The remainder of this paper is organized as follows. Section 2 describes the Lyapunov spectrum and its relation to Kolmogorov entropy. A synopsis of previous methods for calculating Lyapunov exponents from both system equations and experimental time series is also

given. In Section 3 we describe the new approach for calculating λ_1 and show how it differs from previous methods. Section 4 presents the results of our algorithm for several chaotic dynamical systems as well as several non-chaotic systems. We show that the method is robust to variations in embedding dimension, number of data points, reconstruction delay, and noise level. Section 5 is a discussion that includes a description of the procedure for calculating λ_1 and D_2 simultaneously. Finally, Section 6 contains a summary of our conclusions.

2. Background

For a dynamical system, sensitivity to initial conditions is quantified by the Lyapunov exponents. For example, consider two trajectories with nearby initial conditions on an attracting manifold. When the attractor is chaotic, the trajectories diverge, on average, at an exponential rate characterized by the largest Lyapunov exponent [15]. This concept is also generalized for the *spectrum* of Lyapunov exponents, λ_i (i=1, 2, ..., n), by considering a small n-dimensional sphere of initial conditions, where n is the number of equations (or, equivalently, the number of state variables) used to describe the system. As time (t) progresses, the sphere evolves into an ellipsoid whose principal axes expand (or contract) at rates given by the Lyapunov exponents. The presence of a positive exponent is sufficient for diagnosing chaos and represents local instability in a particular direction. Note that for the existence of an attractor, the overall dynamics must be dissipative, i.e., globally stable, and the total rate of contraction must outweigh the total rate of expansion. Thus, even when there are several positive Lyapunov exponents, the sum across the entire spectrum is negative.

Wolf *et al.* [39] explain the Lyapunov spectrum by providing the following geometrical interpretation. First, arrange the n principal axes of the ellipsoid in the order of most rapidly expanding to most rapidly contracting. It follows that the associated Lyapunov exponents will be arranged such that

$$\lambda_1 \geq \lambda_2 \geq \ \ldots \ \geq \lambda_n, \tag{1}$$

where λ_1 and λ_n correspond to the most rapidly expanding and contracting principal axes, respectively. Next, recognize that the length of the first principal axis is proportional to $e^{\lambda_1 t}$; the area determined by the first two principal axes is proportional to $e^{(\lambda_1+\lambda_2)t}$; and the volume determined by the first *k* principal axes is proportional to $e^{(\lambda_1+\lambda_2+\ldots+\lambda_k)t}$. Thus, the Lyapunov spectrum can be defined such that the exponential growth of a *k*-volume element is given by the sum of the *k* largest Lyapunov exponents. Note that information created by the system is represented as a change in the volume defined by the expanding principal axes. The sum of the corresponding exponents, i.e., the positive exponents, equals the Kolmogorov entropy (*K*) or mean rate of information gain [15]:

$$K = \sum_{\lambda_i > 0} \lambda_i \ . \tag{2}$$

When the equations describing the dynamical system are available, one can calculate the entire Lyapunov spectrum [5, 34]. (See [39] for example computer code.) The approach involves numerically solving the system's *n* equations for *n*+1 nearby initial conditions. The growth of a corresponding set of vectors is measured, and as the system evolves, the vectors are repeatedly reorthonormalized using the Gram-Schmidt procedure. This guarantees that only one vector has a component in the direction of most rapid expansion, i.e., the vectors maintain a proper phase space orientation. In experimental settings, however, the equations of motion are usually unknown and this approach is not applicable. Furthermore, experimental data often consist of time series from a single observable, and one must employ a technique for attractor reconstruction, e.g., method of delays [27, 37], singular value decomposition [8].

As suggested above, one cannot calculate the entire Lyapunov spectrum by choosing arbitrary directions for measuring the separation of nearby initial conditions. One must measure the separation along the *Lyapunov directions* which correspond to the principal axes of the ellipsoid previously considered. These Lyapunov directions are dependent upon the system flow and are defined using the Jacobian matrix, i.e., the tangent map, at each point of interest along the flow [15]. Hence, one must preserve the proper phase space orientation by using a suitable

approximation of the tangent map. This requirement, however, becomes unnecessary when calculating only the largest Lyapunov exponent.

If we assume that there exists an ergodic measure of the system, then the multiplicative ergodic theorem of Oseledec [26] justifies the use of arbitrary phase space directions when calculating the largest Lyapunov exponent with smooth dynamical systems. We can expect (with probability 1) that two randomly chosen initial conditions will diverge exponentially at a rate given by the largest Lyapunov exponent [6, 15]. In other words, we can expect that a random vector of initial conditions will converge to the most unstable manifold, since exponential growth in this direction quickly dominates growth (or contraction) along the other Lyapunov directions. Thus, the largest Lyapunov exponent can be defined using the following equation where $d(t)$ is the average divergence at time t and C is a constant that normalizes the initial separation:

$$d(t) = Ce^{\lambda_1 t}. \tag{3}$$

For experimental applications, a number of researchers have proposed algorithms that estimate the largest Lyapunov exponent [1, 10, 12, 16, 17, 29, 33, 38-40], the positive Lyapunov spectrum, i.e., only positive exponents [39], or the complete Lyapunov spectrum [7, 9, 13, 15, 32, 35, 41]. Each method can be considered as a variation of one of several earlier approaches [15, 17, 32, 39] and as suffering from at least one of the following drawbacks: (1) unreliable for small data sets, (2) computationally intensive, (3) relatively difficult to implement. These drawbacks motivated our search for an improved method of estimating the largest Lyapunov exponent.

3. Current Approach

The first step of our approach involves reconstructing the attractor dynamics from a single time series. We use the method of delays [27, 37] since one goal of our work is to develop a fast and easily implemented algorithm. The reconstructed trajectory, $\mathbf{X}$, can be expressed as a matrix where each row is a phase-space vector. That is,

$$\mathbf{X} = \begin{bmatrix} \mathbf{X}_1 & \mathbf{X}_2 & \ldots & \mathbf{X}_M \end{bmatrix}^{\mathrm{T}}, \tag{4}$$

where $\mathbf{X}_i$ is the state of the system at discrete time *i*. For an *N*-point time series, $\{x_1, x_2, \ldots, x_N\}$, each $\mathbf{X}_i$ is given by

$$\mathbf{X}_i = \begin{bmatrix} x_i & x_{i+J} & \ldots & x_{i+(m-1)J} \end{bmatrix}, \tag{5}$$

where *J* is the *lag* or *reconstruction delay*, and *m* is the *embedding dimension*. Thus, $\mathbf{X}$ is an $M \times m$ matrix, and the constants *m*, *M*, *J*, and *N* are related as

$$M = N - (m-1)J. \tag{6}$$

The embedding dimension is usually estimated in accordance with Takens' theorem, i.e., $m > 2n$, although our algorithm often works well when *m* is below the Takens criterion. A method used to choose the lag via the correlation sum was addressed by Liebert and Schuster [23] (based on [19]). Nevertheless, determining the proper lag is still an open problem [4]. We have found a good approximation of *J* to equal the lag where the autocorrelation function drops to $1 - \frac{1}{e}$ of its initial value. Calculating this *J* can be accomplished using the fast Fourier transform (FFT), which requires far less computation than the approach of Liebert and Schuster. Note that our algorithm also works well for a wide range of lags, as shown in Section 4.3.

After reconstructing the dynamics, the algorithm locates the *nearest neighbor* of each point on the trajectory. The nearest neighbor, $\mathbf{X}_{\hat{j}}$, is found by searching for the point that minimizes the distance to the particular *reference point*, $\mathbf{X}_j$. This is expressed as

$$d_j(0) = \min_{\mathbf{X}_{\hat{j}}} \left\| \mathbf{X}_j - \mathbf{X}_{\hat{j}} \right\|, \tag{7}$$

where $d_j(0)$ is the initial distance from the j^{th} point to its nearest neighbor, and $\|..\|$ denotes the Euclidean norm. We impose the additional constraint that nearest neighbors have a temporal separation greater than the mean period of the time series:[1]

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[1] We estimated the mean period as the reciprocal of the mean frequency of the power spectrum, although we expect any comparable estimate, e.g., using the median frequency of the magnitude spectrum, to yield equivalent results.

$$\left| j - \hat{j} \right| > \text{mean period}. \tag{8}$$

This allows us to consider each pair of neighbors as nearby initial conditions for different trajectories. The largest Lyapunov exponent is then estimated as the mean rate of separation of the nearest neighbors.

To this point, our approach for calculating λ_1 is similar to previous methods that track the exponential divergence of nearest neighbors. However, it is important to note some differences:

1) The algorithm by Wolf *et al.* [39] fails to take advantage of all the available data because it focuses on one "fiducial" trajectory. A single nearest neighbor is followed and repeatedly replaced when its separation from the reference trajectory grows beyond a certain limit. Additional computation is also required because the method approximates the Gram-Schmidt procedure by replacing a neighbor with one that preserves its phase space orientation. However, as shown in Section 2, this preservation of phase space orientation is unnecessary when calculating only the largest Lyapunov exponent.
2) If a nearest neighbor precedes (temporally) its reference point, then our algorithm can be viewed as a "prediction" approach. (In such instances, the predictive model is a simple delay line, the prediction is the location of the nearest neighbor, and the prediction error equals the separation between the nearest neighbor and its reference point.) However, other prediction methods use more elaborate schemes, e.g., polynomial mappings, adaptive filters, neural networks, that require much more computation. The amount of computation for the Wales method [38] (based on [36]) is also greater, although it is comparable to the present approach. We have found the Wales algorithm to give excellent results for discrete systems derived from difference equations, e.g., logistic, Hénon, but poor results for continuous systems derived from differential equations, e.g., Lorenz, Rössler.
3) The current approach is principally based on the work of Sato *et al.* [33] which estimates λ_1 as

$$\lambda_1(i) = \frac{1}{i \cdot \Delta t} \cdot \frac{1}{(M-i)} \sum_{j=1}^{M-i} \ln \frac{d_j(i)}{d_j(0)} \ , \tag{9}$$

where Δt is the sampling period of the time series, and $d_j(i)$ is the distance between the j^{th} pair of nearest neighbors after *i* discrete-time steps, i.e., $i \cdot \Delta t$ seconds. (Recall that *M* is the number of reconstructed points as given in Eq. (6).) In order to improve convergence (with respect to *i*), Sato *et al.* [33] give an alternate form of Eq. (9):

$$\lambda_1(i,k) = \frac{1}{k \cdot \Delta t} \cdot \frac{1}{(M-k)} \sum_{j=1}^{M-k} \ln \frac{d_j(i+k)}{d_j(i)} \ . \tag{10}$$

In Eq. (10), k is held constant, and λ_1 is extracted by locating the plateau of $\lambda_1(i,k)$ with respect to *i*. We have found that locating this plateau is sometimes problematic, and the resulting estimates of λ_1 are unreliable. As discussed in Section 5.3, this difficulty is due to the normalization by $d_j(i)$.

The remainder of our method proceeds as follows. From the definition of λ_1 given in Eq. (3), we assume the j^{th} pair of nearest neighbors diverge approximately at a rate given by the largest Lyapunov exponent:

$$d_j(i) \approx C_j e^{\lambda_1 (i \cdot \Delta t)} \ , \tag{11}$$

where C_j is the initial separation. By taking the logarithm of both sides of Eq. (11), we obtain

$$\ln d_j(i) \approx \ln C_j + \lambda_1 (i \cdot \Delta t) \ . \tag{12}$$

Eq. (12) represents a set of approximately parallel lines (for $j = 1, 2, \ldots, M$), each with a slope roughly proportional to λ_1. The largest Lyapunov exponent is easily and accurately calculated using a least-squares fit to the "average" line defined by

$$y(i) = \tfrac{1}{\Delta t} \left\langle \ln d_j(i) \right\rangle \ , \tag{13}$$

where $\langle .. \rangle$ denotes the average over all values of *j*. This process of averaging is the key to calculating accurate values of λ_1 using small, noisy data sets. Note that in Eq. (11), C_j performs the function of normalizing the separation of the neighbors, but as shown in Eq. (12), this

normalization is unnecessary for estimating λ_1. By avoiding the normalization, the current approach gains a slight computational advantage over the method by Sato *et al*.

**** FIGURE 1 NEAR HERE ****

The new algorithm for calculating largest Lyapunov exponents is outlined in Figure 1. This method is easy to implement and fast because it uses a simple measure of exponential divergence that circumvents the need to approximate the tangent map. The algorithm is also attractive from a practical standpoint because it does not require large data sets and it simultaneously yields the correlation dimension (discussed in Section 5.5). Furthermore, the method is accurate for small data sets because it takes advantage of all the available data. In the next section, we present the results for several dynamical systems.

4. Experimental Results

Table I summarizes the chaotic systems primarily examined in this paper. The differential equations were solved numerically using a fourth-order Runge-Kutta integration with a step size equal to Δt as given in Table I. For each system, the initial point was chosen near the attractor and the transient points were discarded. In all cases, the *x*-coordinate time series was used to reconstruct the dynamics. Figure 2 shows a typical plot (solid curve) of $\left\langle \ln d_j(i) \right\rangle$ versus $i \cdot \Delta t$;[2] the dashed line has a slope equal to the theoretical value of λ_1. After a short transition, there is a long linear region that is used to extract the largest Lyapunov exponent. The curve saturates at longer times since the system is bounded in phase space and the average divergence cannot exceed the "length" of the attractor.

**** TABLE I & FIGURE 2 NEAR HERE ****

The remainder of this section contains tabulated results from our algorithm under different conditions. The corresponding plots are meant to give the reader qualitative information about the facility of extracting λ_1 from the data. That is, the more prominent the

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[2] In each figure, "< ln(divergence) >" and "Time (s)" are used to denote $\left\langle \ln d_j(i) \right\rangle$ and $i \cdot \Delta t$, respectively.

linear region, the easier one can extract the correct slope. (Repeatability is discussed in Section 5.2.)

4.1. Embedding dimension

Since we normally have no *a priori* knowledge concerning the dimension of a system, it is imperative that we evaluate our method for different embedding dimensions. Table II and Figure 3 show our findings for several values of m. In all but three cases (m=1 for the Hénon, Lorenz, and Rössler systems), the error was less than ±10%, and most errors were less than ±5%. It is apparent that satisfactory results are obtained only when m is at least equal to the topological dimension of the system, i.e., $m \geq n$. This is due to the fact that chaotic systems are effectively stochastic when embedded in a phase space that is too small to accommodate the true dynamics. Notice that the algorithm performs quite well when m is below the Takens criterion. Therefore, it seems one may choose the smallest embedding dimension that yields a convergence of the results.

**** TABLE II & FIGURE 3 NEAR HERE ****

4.2. Length of time series

Next we consider the performance of our algorithm for time series of various lengths. As shown in Table III and Figure 4, the present method also works well when N is small (N=100 to 1000 for the examined systems). Again, the error was less than ±10% in almost all cases. (The greatest difficulty occurs with the Rössler attractor. For this system, we also found a 20-25% negative bias in the results for N=3000 to 5000.) To our knowledge, the lower limit of N used in each case is less than the smallest value reported in the literature. (The only exception is due to Briggs [7], who examined the Lorenz system with N=600. However, Briggs reported errors for λ_1 that ranged from 54% to 132% for this particular time series length.) We also point out that the literature [1, 9, 13, 15, 35] contains results for values of N that are an order of magnitude greater than the largest values used here.

**** TABLE III & FIGURE 4 NEAR HERE ****

It is important to mention that quantitative analyses of chaotic systems are usually sensitive to not only the data size (in samples), but also the observation time (in seconds). Hence, we examined the interdependence of N and $N \cdot \Delta t$ for the Lorenz system. Figure 5 shows the output of our algorithm for three different sampling conditions: (1) $N = 5000$, $\Delta t = 0.01$ s ($N \cdot \Delta t = 50$ s); (2) $N = 1000$, $\Delta t = 0.01$ s ($N \cdot \Delta t = 10$ s); and (3) $N = 1000$, $\Delta t = 0.05$ s ($N \cdot \Delta t = 50$ s). The latter two time series were derived from the former by using the first 1000 points and every fifth point, respectively. As expected, the best results were obtained with a relatively long observation time and closely-spaced samples (Case 1). However, we saw comparable results with the long observation time and widely-spaced samples (Case 3). As long as Δt is small enough to ensure a minimum number of points per orbit of the attractor (approximately n to $10n$ points [39]), it is better to decrease N by reducing the sampling rate and not the observation time.

**** FIGURE 5 NEAR HERE ****

4.3. Reconstruction delay

As commented in Section 3, determining the proper reconstruction delay is still an open problem. For this reason, it is necessary to test our algorithm with different values of J. (See Table IV and Figure 6.) Since discrete maps are most faithfully reconstructed with a delay equal to one, it is not surprising that the best results were seen with the lag equal to one for the logistic and Hénon systems (errors of -1.7% and -2.2%, respectively). For the Lorenz and Rössler systems, the algorithm performed well (error ≤ 7%) with all lags except the extreme ones (J=1, 41 for Lorenz; J=2, 26 for Rössler). Thus, we expect satisfactory results whenever the lag is determined using any common method such as those based on the autocorrelation function or the correlation sum. Notice that the smallest errors were obtained for the lag where the autocorrelation function drops to $1 - \frac{1}{e}$ of its initial value.

**** TABLE IV & FIGURE 6 NEAR HERE ****

4.4. Additive noise

Next, we consider the effects of additive noise, i.e., measurement or instrumentation noise. This was accomplished by examining several time series produced by a superposition of white noise and noise-free data (noise-free up to the computer precision). Before superposition, the white noise was scaled by an appropriate factor in order to achieve a desired signal-to-noise ratio (SNR). The SNR is the ratio of the power (or, equivalently, the variance) in the noise-free signal and that of the pure-noise signal. A signal-to-noise ratio greater than about 1000 can be regarded as low noise and a SNR less than about 10 as high noise.

The results are shown in Table V and Figure 7. We expect satisfactory estimates of λ_1 except in extremely noisy situations. With low noise, the error was smaller than ±10% in each case. At moderate noise levels (SNR ranging from about 100 to 1000), the algorithm performed reasonably well with an error that was generally near ±25%. As expected, the poorest results were seen with the highest noise levels (SNR less than or equal to 10). (We believe that the improved performance with the logistic map and low signal-to-noise ratios is merely coincidental. The reader should equate the shortest linear regions in Figure 7 with the highest noise and greatest uncertainty in estimating λ_1.) It seems one cannot expect to estimate the largest Lyapunov exponent in high-noise environments; however, the clear presence of a positive slope still affords one the qualitative confirmation of a positive exponent (and chaos).

**** TABLE V & FIGURE 7 NEAR HERE ****

It is important to mention that the adopted noise model represents a "worst-case" scenario because white noise contaminates a signal across an infinite bandwidth. (Furthermore, we consider signal-to-noise ratios that are substantially lower than most values previously reported in the literature.) Fortunately, some of the difficulties are remedied by filtering, which is expected to preserve the exponential divergence of nearest neighbors [39]. Whenever we remove noise while leaving the signal intact, we can expect an improvement in system predictability and, hence, in our ability to detect chaos. In practice, however, caution is

warranted because the underlying signal may have some frequency content in the stopband or the filter may substantially alter the phase in the passband.

4.5. Two positive Lyapunov exponents

As described in Section 2, it is unnecessary to preserve phase space orientation when calculating the largest Lyapunov exponent. In order to provide experimental verification of this theory, we consider the performance of our algorithm with two systems that possess more than one positive exponent: Rössler-hyperchaos [30] and Mackey-Glass [25]. (See Table VI for details.) The results are shown in Table VII and Figure 8. For both systems, the errors were typically less than ±10%. From these results, we conclude that the algorithm measures exponential divergence along the most unstable manifold and not along some other Lyapunov direction. However, notice the predominance of a negative bias in the errors presented in Sections 4.1-4.4. We believe that over short time scales, some nearest neighbors explore Lyapunov directions other than that of the largest Lyapunov exponent. Thus, a small underestimation (less than 5%) of λ_1 is expected.

**** TABLE VI NEAR HERE ****

**** TABLE VII & FIGURE 8 NEAR HERE ****

4.6. Non-chaotic systems

As stated earlier, distinguishing deterministic chaos from noise has become an important problem. It follows that effective algorithms for detecting chaos must accurately characterize both chaotic and non-chaotic systems; a reliable algorithm is not "fooled" by difficult systems such as correlated noise. Hence, we further establish the utility of our method by examining its performance with the following non-chaotic systems: two-torus, white noise, bandlimited noise, and "scrambled" Lorenz.

For each system, a 2000-point time series was generated. The two-torus is an example of a quasiperiodic, deterministic system. The corresponding time series, $x(i)$, was created by a superposition of two sinusoids with incommensurate frequencies:

$$x(i) = \sin(2\pi f_1 \cdot i \cdot \Delta t) + \sin(2\pi f_2 \cdot i \cdot \Delta t), \tag{14}$$

where $f_1 = 1.732051 \approx \sqrt{3}$ Hz, $f_2 = 2.236068 \approx \sqrt{5}$ Hz, and the sampling period was $\Delta t = 0.01$ s. White noise and bandlimited noise are stochastic systems that are analogous to discrete and continuous chaotic systems, respectively. The "scrambled" Lorenz also represents a continuous stochastic system, and the data set was generated by randomizing the phase information from the Lorenz attractor. This procedure yields a time series of correlated noise with spectral characteristics identical to that of the Lorenz attractor.

For quasiperiodic and stochastic systems we expect flat plots of $\left\langle \ln d_j(i) \right\rangle$ versus $i \cdot \Delta t$. That is, on average the nearest neighbors should neither diverge nor converge. Additionally, with the stochastic systems we expect an initial "jump" from a small separation at *t*=0. The results are shown in Figure 9, and as expected, the curves are mostly flat. However, notice the regions that could be mistaken as appropriate for extracting a positive Lyapunov exponent. Fortunately, our empirical results suggest that one may still detect non-chaotic systems for the following reasons:

1) The anomalous scaling region is not linear since the divergence of nearest neighbors is not exponential.
2) For stochastic systems, the anomalous scaling region flattens with increasing embedding dimension. Finite dimensional systems exhibit a convergence once the embedding dimension is large enough to accomodate the dynamics, whereas stochastic systems fail to show a convergence because they appear more ordered in higher embedding spaces. With the two-torus, we attribute the lack of convergence to the finite precision "noise" in the data set. (Notice the small average divergence even at $i \cdot \Delta t$=1.) Strictly speaking, we can only distinguish high-dimensional systems from low-dimensional ones, although in most applications a high-dimensional system may be considered random, i.e., infinite-dimensional.

**** FIGURE 9 NEAR HERE ****

5. Discussion

5.1. Eckmann-Ruelle requirement

In a recent paper, Eckmann and Ruelle [14] discuss the data-set size requirement for estimating dimensions and Lyapunov exponents. Their analysis for Lyapunov exponents proceeds as follows. When measuring the rate of divergence of trajectories with nearby initial conditions, one requires a number of neighbors for a given reference point. These neighbors should lie in a ball of radius r, where r is small with respect to the diameter (d) of the reconstructed attractor. Thus,

$$\frac{r}{d} = \rho << 1. \tag{15}$$

(Eckmann and Ruelle suggest ρ to be a maximum of about 0.1.) Furthermore, the number of candidates for neighbors, $\Gamma(r)$, should be much greater than one:

$$\Gamma(r) >> 1. \tag{16}$$

Next, recognize that

$$\Gamma(r) \approx \text{constant} \times r^D, \tag{17}$$

and

$$\Gamma(d) \approx N, \tag{18}$$

where D is the dimension of the attractor, and N is the number of data points. Using Eqs. (16)-(18), we obtain the following relation:

$$\Gamma(r) \approx N\left(\frac{r}{d}\right)^D >> 1. \tag{19}$$

Finally, Eqs. (15) and (19) are combined to give the Eckmann-Ruelle requirement for Lyapunov exponents:

$$\log N > D \log(1/\rho). \tag{20}$$

For $\rho = 0.1$, Eq. (20) directs us to choose N such that

$$N > 10^D. \tag{21}$$

This requirement was met with all time series considered in this paper. Notice that any rigorous definition of "small data set" should be a function of dimension. However, for comparative purposes we regard a small data set as one that is small with respect to those previously considered in the literature.

5.2. Repeatability

When using the current approach for estimating largest Lyapunov exponents, one is faced with the following issue of repeatability: Can one consistently locate the region for extracting λ_1 without a guide, i.e., without *a priori* knowledge of the correct slope in the linear region?[3] To address this issue, we consider the performance of our algorithm with multiple realizations of the Lorenz attractor.

Three 5000-point time series from the Lorenz attractor were generated by partitioning one 15000-point data set into disjoint time series. Figure 10 shows the results using a visual format similar to that first used by Abraham *et al.* [2] for estimating dimensions. Each curve is a plot of slope versus time, where the slope is calculated from a least-squares fit to 51-point segments of the $\left\langle \ln d_j(i) \right\rangle$ versus $i \cdot \Delta t$ curve. We observe a clear and repeatable plateau from about $i \cdot \Delta t$=0.6 to about $i \cdot \Delta t$=1.6. By using this range to define the region for extracting λ_1, we obtain a reliable estimate of the largest Lyapunov exponent: λ_1=1.57 ± 0.03. (Recall that the theoretical value is 1.50.)

**** FIGURE 10 NEAR HERE ****

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[3] In Tables II-IV, there appear to be inconsistent results when using identical values of *N*, *J*, and *m* for a particular system. These small discrepencies are due to the subjective nature in choosing the linear region and not the algorithm itself. In fact, the same output file was used to compute λ_1 in each case.

5.3. Relation to the Sato algorithm

As stated in Section 3, the current algorithm is principally based on the work of Sato *et al.* [33]. More specifically, our approach can be considered as a generalization of the Sato algorithm. To show this, we first rewrite Eq. (10) using $\langle..\rangle$ to denote the average over all values of *j*:

$$\lambda_1(i,k) = \frac{1}{k \cdot \Delta t}\left\langle \ln \frac{d_j(i+k)}{d_j(i)} \right\rangle . \tag{22}$$

This equation is then rearranged and expressed in terms of the output from the current algorithm, $y(i)$ (from Eq. (13)):

$$\begin{aligned} \lambda_1(i,k) &= \frac{1}{k \cdot \Delta t}\left[\left\langle \ln d_j(i+k)\right\rangle - \left\langle \ln d_j(i)\right\rangle\right] \\ &\approx \frac{1}{k}\left[y(i+k) - y(i)\right] \end{aligned} . \tag{23}$$

Eq. (23) is interpreted as a finite-differences numerical differentiation of $y(i)$, where *k* specifies the size of the differentiation interval.

Next, we attempt to derive $y(i)$ from the output of the Sato algorithm by summing $\lambda_1(i,k)$. That is, we define $y'(i')$ as

$$y'(i') = \sum_{i=0}^{i'} \lambda_1(i,k) = \frac{1}{k}\left[\sum_{i=0}^{i'} y(i+k) - \sum_{i=0}^{i'} y(i)\right]. \tag{24}$$

By manipulating this equation, we can show that Eq. (23) is not invertible:

$$\begin{aligned} y'(i') &= \frac{1}{k}\left[\sum_{i=0}^{i'+k} y(i) - \sum_{i=0}^{k-1} y(i) - \sum_{i=0}^{i'} y(i)\right] \\ &= \frac{1}{k}\left[\sum_{i=i'+1}^{i'+k} y(i) - \sum_{i=0}^{k-1} y(i)\right] \\ &= \frac{1}{k}\sum_{i=i'+1}^{i'+k} y(i) + \text{constant} \end{aligned} . \tag{25}$$

If we disregard the constant in Eq. (25), $y'(i')$ is equivalent to $y(i)$ smoothed by a *k*-point moving-average filter.

The difficulty with the Sato algorithm is that the proper value of k is not usually apparent *a priori*. When choosing k, one must consider the tradeoff between long, noisy plateaus of $\lambda_1(i,k)$ (for small k) and short, smooth plateaus (for large k). In addition, since the transformation from $y(i)$ to $\lambda_1(i,k)$ is not invertible, choosing k by trial-and-error requires the repeated evaluation of Eq. (22). With our algorithm, however, smoothing is usually unnecessary, and λ_1 is extracted from a least-squares fit to the longest possible linear region. For those cases where smoothing is needed, a long filter length may be chosen since one knows the approximate location of the plateau after examining a plot of $\left\langle \ln d_j(i) \right\rangle$ versus $i \cdot \Delta t$. (For example, one may choose a filter length equal to about one-half the length of the noisy linear region.)

5.4. Computational improvements

In some instances, the speed of the method may be increased by measuring the separation of nearest neighbors using a smaller embedding dimension. For example, we reconstructed the Lorenz attractor in a three-dimensional phase space and located the nearest neighbors. The separations of those neighbors were then measured in a one-dimensional space by comparing only the first coordinates of each point. There was nearly a threefold savings in time for this portion of the algorithm. However, additional fluctuations were seen in the plots of $\left\langle \ln d_j(i) \right\rangle$ versus $i \cdot \Delta t$, making it more difficult to locate the region for extracting the slope.

Similarly, the computational efficiency of the algorithm may be improved by disregarding every other reference point. We observed that many temporally adjacent reference points also have temporally adjacent nearest neighbors. Thus, two pairs of trajectories may exhibit identical divergence patterns (excluding a time shift of one sampling period), and it may be unnecessary to incorporate the effects of both pairs. Note that this procedure still satisfies the Eckmann-Ruelle requirement by maintaining the pool of nearest neighbors.

5.5. Simultaneous calculation of correlation dimension

In addition to calculating the largest Lyapunov exponent, the present algorithm allows one to calculate the correlation dimension, D_2. Thus, one sequence of computations will yield an estimate of both the level of chaos and the system complexity. This is accomplished by

taking advantage of the numerous distance calculations performed during the nearest-neighbors search.

The Grassberger-Procaccia algorithm [20] estimates dimension by examining the scaling properties of the correlation sum, $C_m(r)$. For a given embedding dimension, m, $C_m(r)$ is defined as

$$C_m(r) = \frac{2}{M(M-1)} \sum_{i \neq k} \theta\left[r - \left\|\mathbf{X}_i - \mathbf{X}_k\right\|\right], \tag{26}$$

where $\theta[..]$ is the Heavyside function. Therefore, $C_m(r)$ is interpreted as the fraction of pairs of points that are separated by a distance less than or equal to r. Notice that the previous equation and Eq. (7) of our algorithm require the same distance computations (disregarding the constraint in Eq. (8)). By exploiting this redundancy, we obtain a more complete characterization of the system using a negligible amount of additional computation.

6. Summary

We have presented a new method for calculating the largest Lyapunov exponent from experimental time series. The method follows directly from the definition of the largest Lyapunov exponent and is accurate because it takes advantage of all the available data. The algorithm is fast because it uses a simple measure of exponential divergence and works well with small data sets. In addition, the current approach is easy to implement and robust to changes in the following quantities: embedding dimension, size of data set, reconstruction delay, and noise level. Furthermore, one may use the algorithm to calculate simultaneously the correlation dimension.

Acknowledgments

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Captions

Table I. Chaotic dynamical systems with theoretical values for the largest Lyapunov exponent, λ_1. The sampling period is denoted by Δt.

Table II. Experimental results for several embedding dimensions. The number of data points, reconstruction delay, and embedding dimension are denoted by *N*, *J*, and *m*, respectively. We were unable to extract λ_1 with *m* equal to one for the Lorenz and Rössler systems because the reconstructed attractors are extremely noisy in a one-dimensional embedding space.

Table III. Experimental results for several time series lengths. The number of data points, reconstruction delay, and embedding dimension are denoted by *N*, *J*, and *m*, respectively.

Table IV. Experimental results for several reconstruction delays. The number of data points, reconstruction delay, and embedding dimension are denoted by *N*, *J*, and *m*, respectively. The asterisks denote the values of *J* that were obtained by locating the lag where the autocorrelation function drops to $1-\frac{1}{e}$ of its initial value.

Table V. Experimental results for several noise levels. The number of data points, reconstruction delay, and embedding dimension are denoted by *N*, *J*, and *m*, respectively. The signal-to-noise ratio (SNR) is the ratio of the power in the noise-free signal to that of the pure-noise signal.

Table VI. Chaotic systems with two positive Lyapunov exponents (λ_1, λ_2). To obtain a better representation of the dynamics, the numerical integrations were performed using a

step size 100 times smaller than the sampling period, Δt. The resulting time series were then downsampled by a factor of 100 to achieve the desired Δt.

Table VII. Experimental results for systems with two positive Lyapunov exponents. The number of data points, reconstruction delay, and embedding dimension are denoted by *N*, *J*, and *m*, respectively.

Figure 1. Flowchart of the practical algorithm for calculating largest Lyapunov exponents.

Figure 2. Typical plot of < ln(divergence) > versus time for the Lorenz attractor. The solid curve is the calculated result; the slope of the dashed curve is the expected result.

Figure 3. Effects of embedding dimension. For each plot, the solid curves are the calculated results, and the slope of the dashed curve is the expected result. See Table II for details. (a) Logistic map. (b) Hénon attractor. (c) Lorenz attractor. (d) Rössler attractor.

Figure 4. Effects of time series length. For each plot, the solid curves are the calculated results, and the slope of the dashed curve is the expected result. See Table III for details. (a) Logistic map. (b) Hénon attractor. (c) Lorenz attractor. (d) Rössler attractor.

Figure 5. Results for the Lorenz system using three different sampling conditions. Case 1: $N = 5000$, $\Delta t = 0.01$ s ($N \cdot \Delta t = 50$ s); Case 2: $N = 1000$, $\Delta t = 0.01$ s ($N \cdot \Delta t = 10$ s); and Case 3: $N = 1000$, $\Delta t = 0.05$ s ($N \cdot \Delta t = 50$ s). The slope of the dashed curve is the expected result.

Figure 6. Effects of reconstruction delay. For each plot, the solid curves are the calculated results, and the slope of the dashed curve is the expected result. See Table IV for details. (a) Logistic map. (b) Hénon attractor. (c) Lorenz attractor. (d) Rössler attractor.

Figure 7. Effects of noise level. For each plot, the solid curves are the calculated results, and the slope of the dashed curve is the expected result. See Table V for details. (a) Logistic map. (b) Hénon attractor. (c) Lorenz attractor. (d) Rössler attractor.

Figure 8. Results for systems with two positive Lyapunov exponents. For each plot, the solid curves are the calculated results, and the slope of the dashed curve is the expected result. See Table VII for details. (a) Rössler-hyperchaos. (b) Mackey-Glass.

Figure 9. Effects of embedding dimension for non-chaotic systems. (a) Two-torus. (b) White noise. (c) Bandlimited noise. (d) "Scrambled" Lorenz.

Figure 10. Plot of $\frac{d}{dt}\left\langle \ln d_j(i) \right\rangle$ versus $i \cdot \Delta t$ using our algorithm with three 5000-point realizations of the Lorenz attractor.

System [ref.]	Equations	Parameters	Δt (s)	Expected λ_1 [ref.]
logistic [15]	$x_{i+1} = \mu x_i (1 - x_i)$	μ = 4.0	1	0.693 [15]
Hénon [22]	$x_{i+1} = 1 - a x_i^2 + y_i$ $y_{i+1} = b x_i$	a = 1.4 b = 0.3	1	0.418 [39]
Lorenz [24]	$\dot{x} = \sigma(y - x)$ $\dot{y} = x(R - z) - y$ $\dot{z} = xy - bz$	σ = 16.0 R = 45.92 b = 4.0	0.01	1.50 [39]
Rössler [31]	$\dot{x} = -y - z$ $\dot{y} = x + ay$ $\dot{z} = b + z(x - c)$	a = 0.15 b = 0.20 c = 10.0	0.10	0.090 [39]

(TABLE I)

System	N	J	m	Calculated λ_1	% error
logistic	500	1	1	0.675	-2.6
			2	0.681	-1.7
			3	0.680	-1.9
			4	0.680	-1.9
			5	0.651	-6.1
Hénon	500	1	1	0.195	-53.3
			2	0.409	-2.2
			3	0.406	-2.9
			4	0.399	-4.5
			5	0.392	-6.2
Lorenz	5000	11	1	-	-
			3	1.531	2.1
			5	1.498	-0.1
			7	1.562	4.1
			9	1.560	4.0
Rössler	2000	8	1	-	-
			3	0.0879	-2.3
			5	0.0864	-4.0
			7	0.0853	-5.2
			9	0.0835	-7.2

(TABLE II)

System	N	J	m	Calculated λ_1	% error
logistic	100	1	2	0.659	-4.9
	200			0.705	1.7
	300			0.695	0.3
	400			0.692	-0.1
	500			0.686	-1.0
Hénon	100	1	2	0.426	1.9
	200			0.416	-0.5
	300			0.421	0.7
	400			0.409	-2.2
	500			0.412	-1.4
Lorenz	1000	11	3	1.751	16.7
	2000			1.345	-10.3
	3000			1.372	-8.5
	4000			1.392	-7.2
	5000			1.523	1.5
Rössler	400	8	3	0.0351	-61.0
	800			0.0655	-27.2
	1200			0.0918	2.0
	1600			0.0984	9.3
	2000			0.0879	-2.3

(TABLE III)

System	N	J	m	Calculated λ_1	% error
logistic	500	1 *	2	0.681	-1.7
		2		0.678	-2.2
		3		0.672	-3.0
		4		0.563	-18.8
		5		0.622	-10.2
Hénon	500	1 *	2	0.409	-2.2
		2		0.406	-2.9
		3		0.391	-6.5
		4		0.338	-19.1
		5		0.330	-21.1
Lorenz	5000	1	3	1.640	9.3
		11 *		1.561	4.1
		21		1.436	-4.3
		31		1.423	-5.1
		41		1.321	-11.9
Rössler	2000	2	3	0.0699	-22.3
		8 *		0.0873	-3.0
		14		0.0864	-4.0
		20		0.0837	-7.0
		26		0.0812	-9.8

(TABLE IV)

System	N	J	m	SNR	Calculated λ_1	% error
logistic	500	1	2	1	0.704	1.6
				10	0.779	12.4
				100	0.856	23.5
				1000	0.621	-10.4
				10000	0.628	-9.4
Hénon	500	1	2	1	0.643	53.8
				10	0.631	51.0
				100	0.522	24.9
				1000	0.334	-20.1
				10000	0.385	-7.9
Lorenz	5000	11	3	1	0.645	-57.0
				10	1.184	-21.1
				100	1.110	-26.0
				1000	1.273	-15.1
				10000	-1.470	-2.0
Rössler	2000	8	3	1	0.0106	-88.2
				10	0.0394	-56.2
				100	0.0401	-55.4
				1000	0.0659	-26.8
				10000	0.0836	-7.1

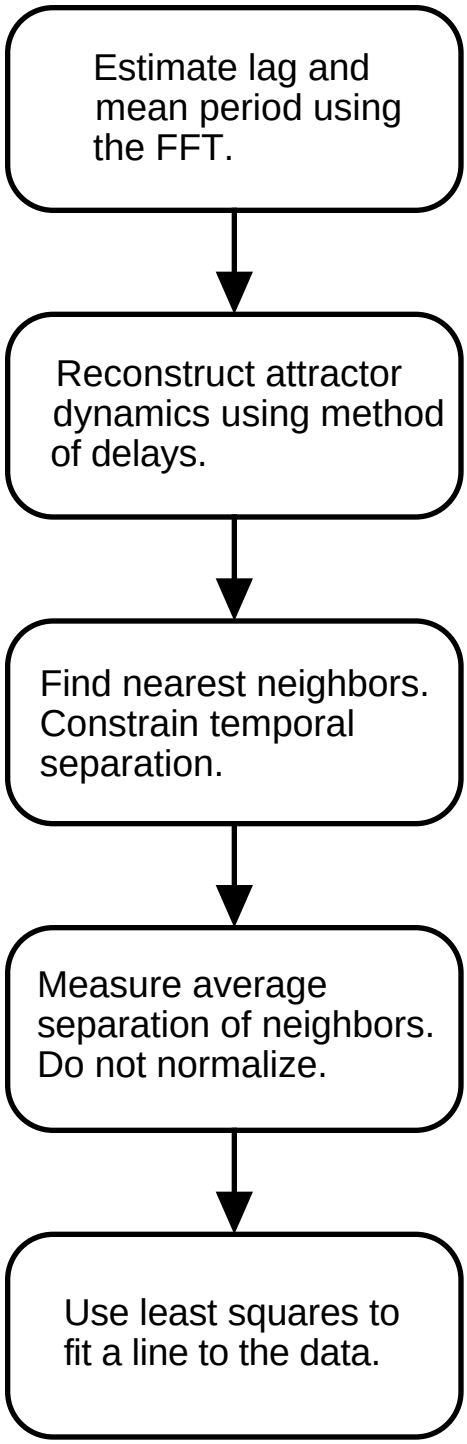
(TABLE V)

System [ref.]	Equations	Parameters	Δt (s)	Expected λ_1, λ_2 [ref.]
Rössler-hyperchaos [30]	$\dot{x} = -y - z$	a=0.25	0.1	λ_1=0.111 [39]
	$\dot{y} = x + ay + w$	b=3.0		λ_2=0.021 [39]
	$\dot{z} = b + xz$	c=0.05		
	$\dot{w} = cw - dz$	d=0.5		
Mackey-Glass [25]	$\dot{x} = \frac{ax(t+s)}{1+[x(t+s)]^c} - bx(t)$	a=0.2	0.75	λ_1=4.37E-3 [39]
		b=0.1		λ_2=1.82E-3 [39]
		c=10.0		
		s=31.8		

(TABLE VI)

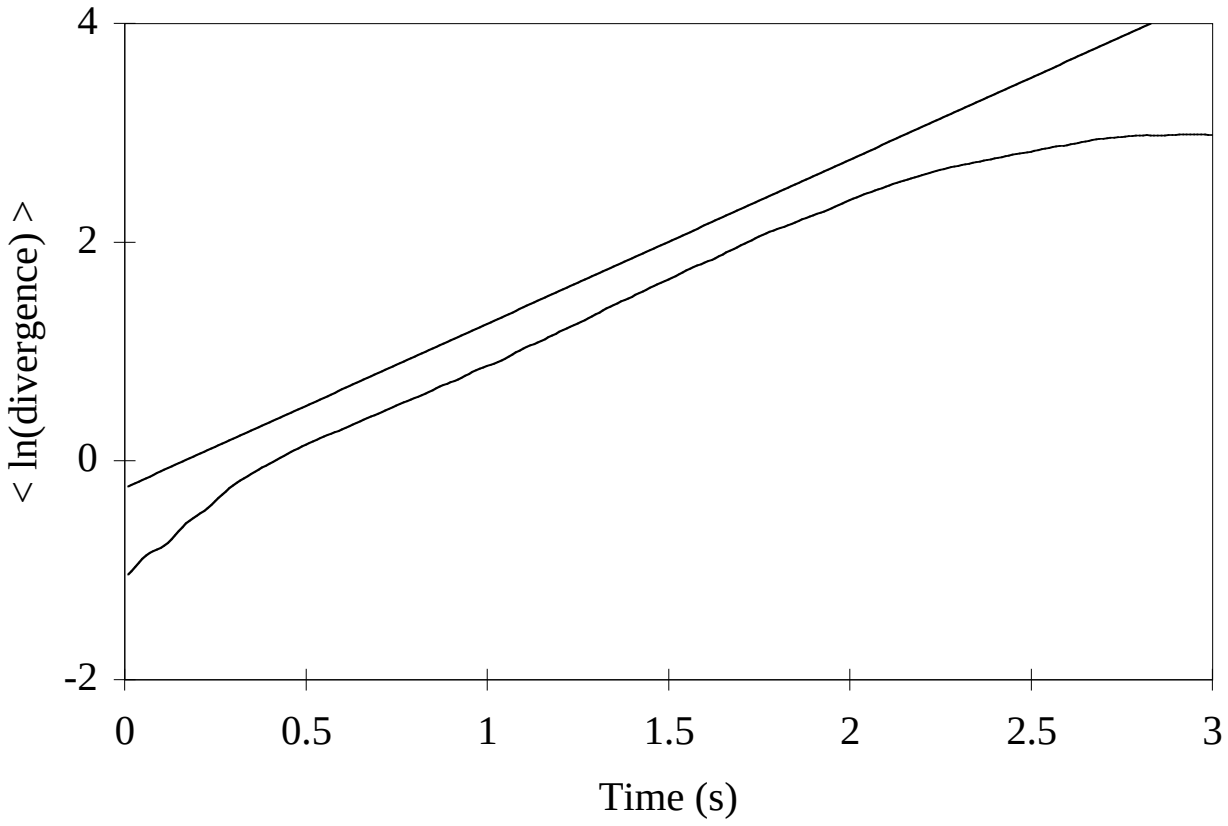
System	N	J	m	Calculated λ_1	% error
Rössler-hyperchaos	8000	9	3	0.048	-56.8
			6	0.112	0.9
			9	0.112	0.9
			12	0.107	-3.6
			15	0.102	-8.1
Mackey-Glass	8000	12	3	4.15E-3	-5.0
			6	4.87E-3	11.4
			9	4.74E-3	8.5
			12	4.80E-3	9.8
			15	4.85E-3	11.0

(TABLE VII)

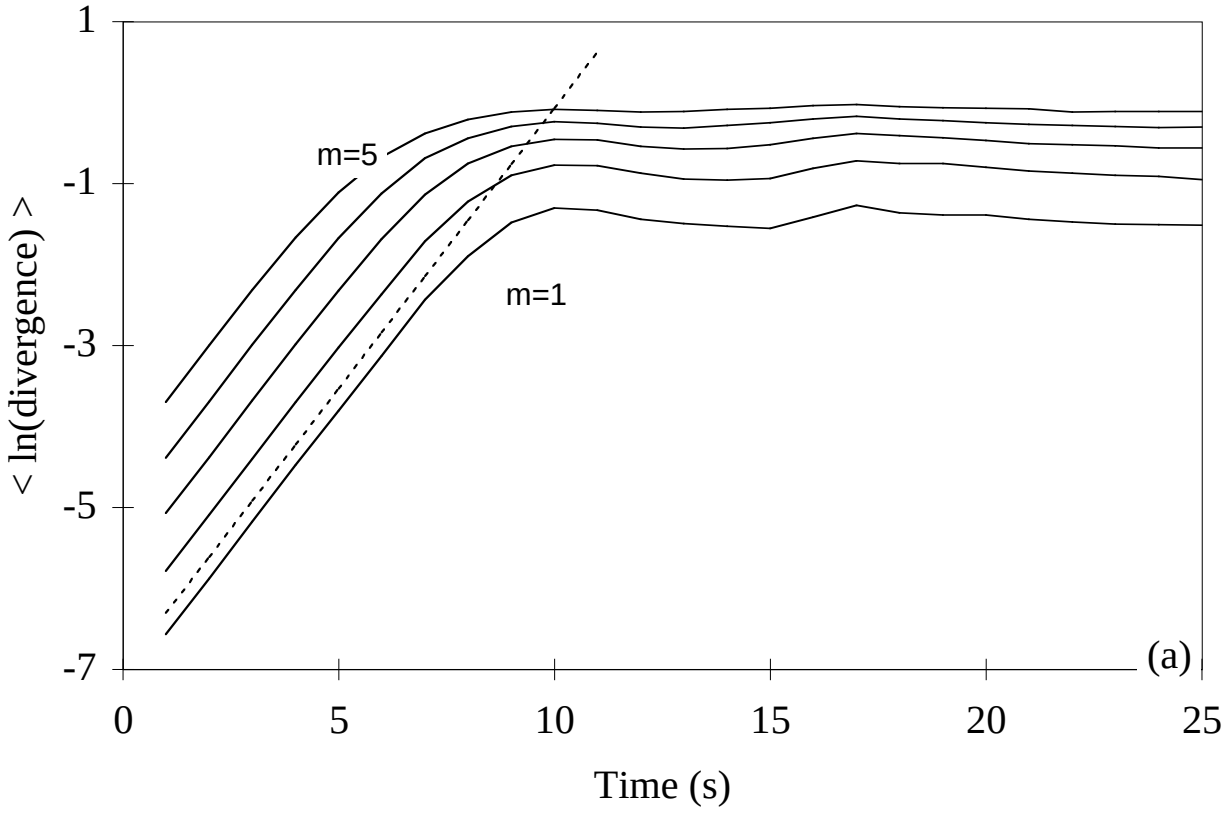

Estimate lag and mean period using the FFT.
Reconstruct attractor dynamics using method of delays.
Find nearest neighbors. Constrain temporal separation.
Measure average separation of neighbors. Do not normalize.
Use least squares to fit a line to the data.

(FIGURE 1)

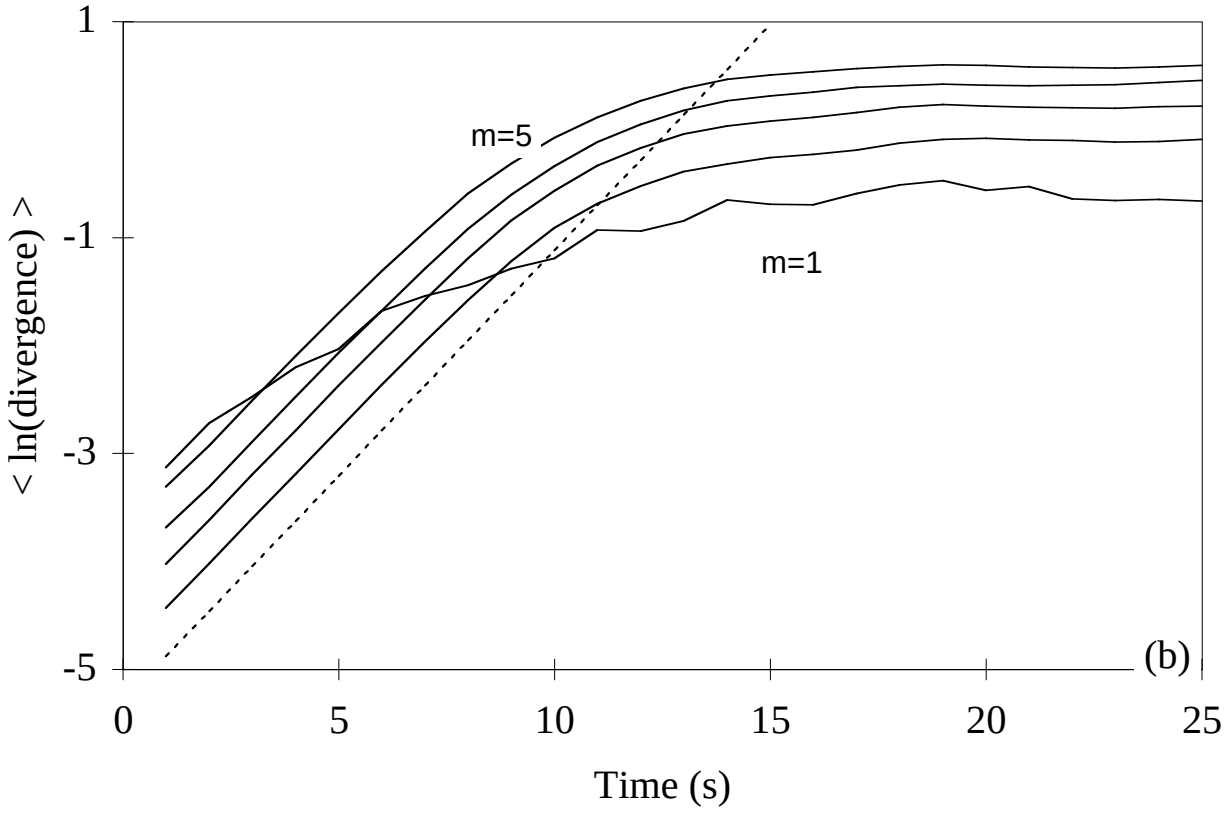
(FIGURE 2)



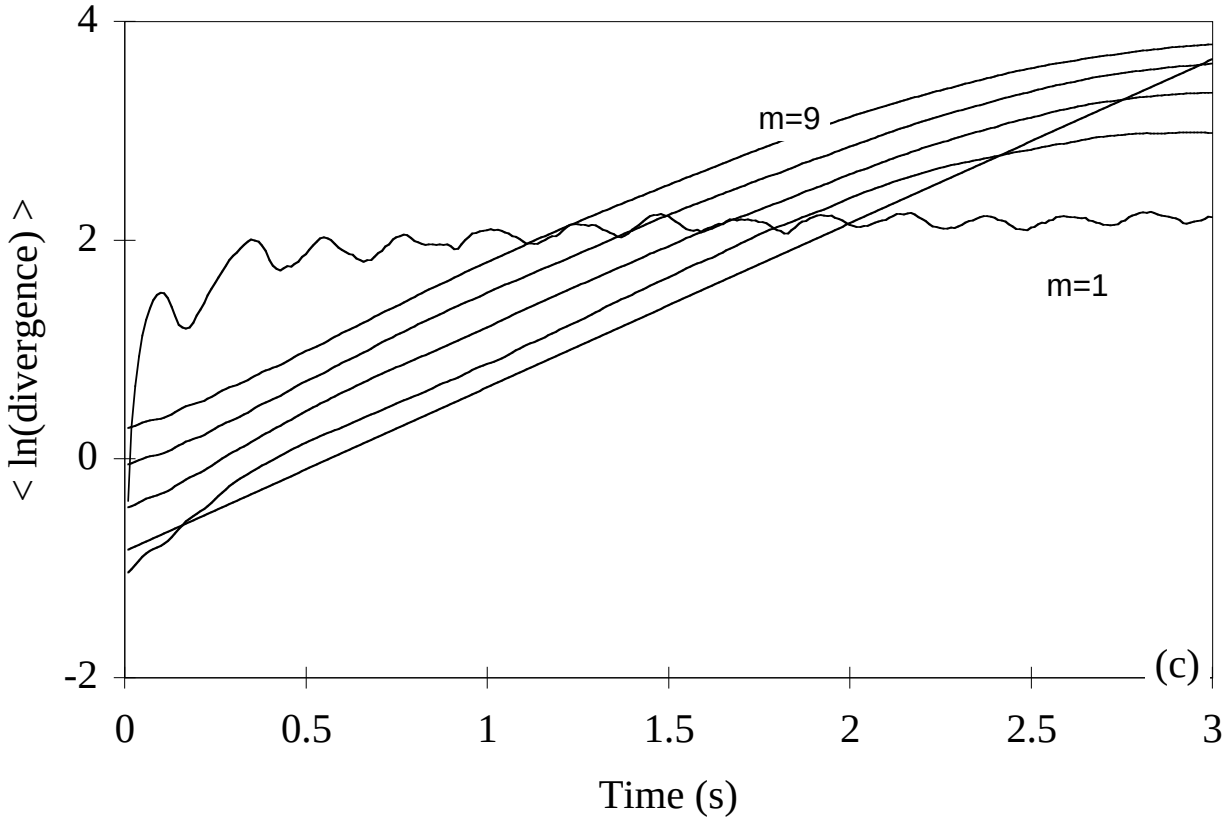
(FIGURE 3A)



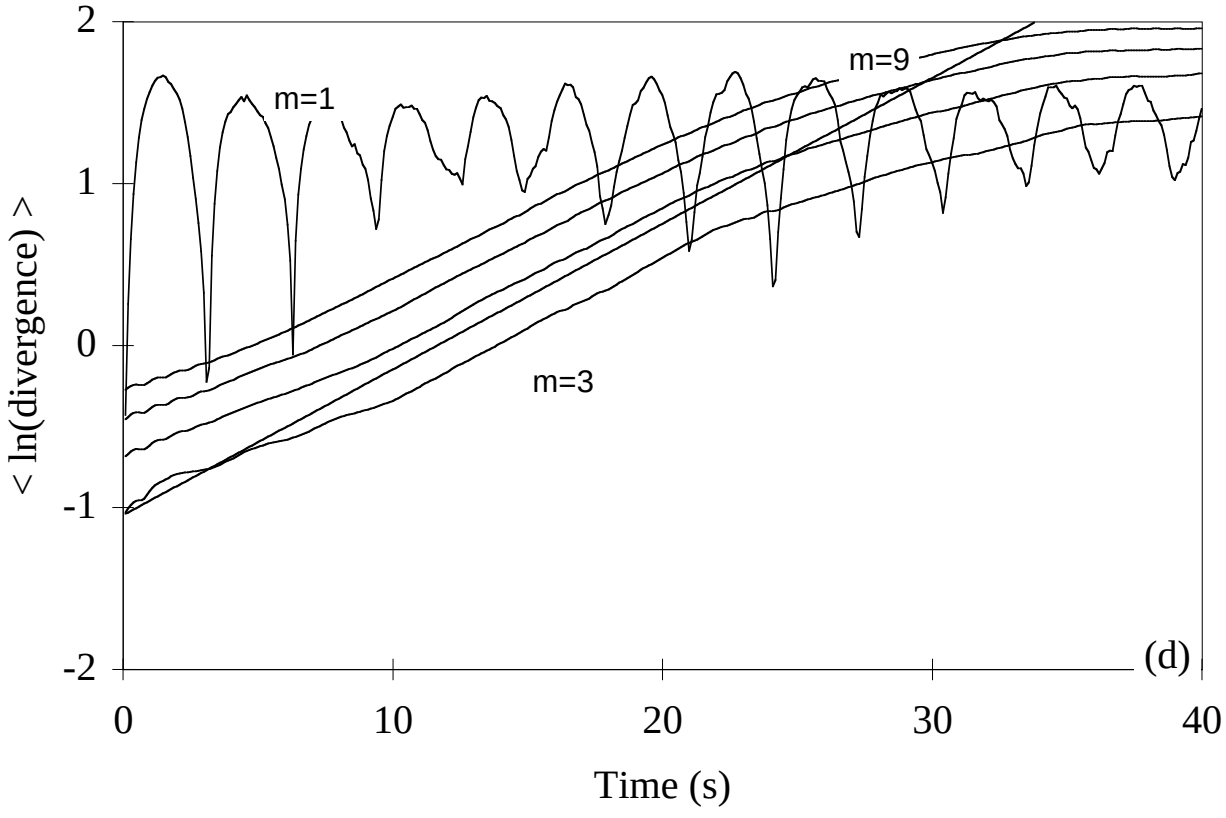
(FIGURE 3B)



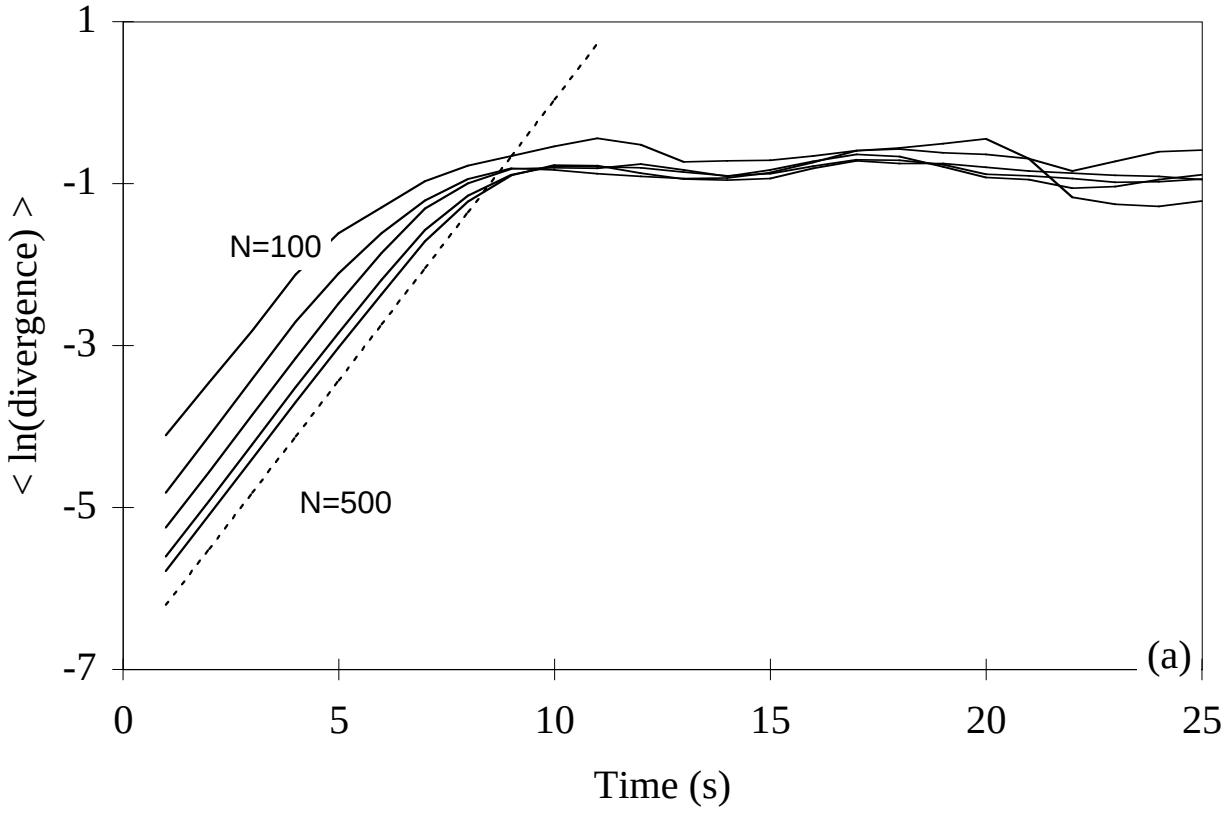
(FIGURE 3C)



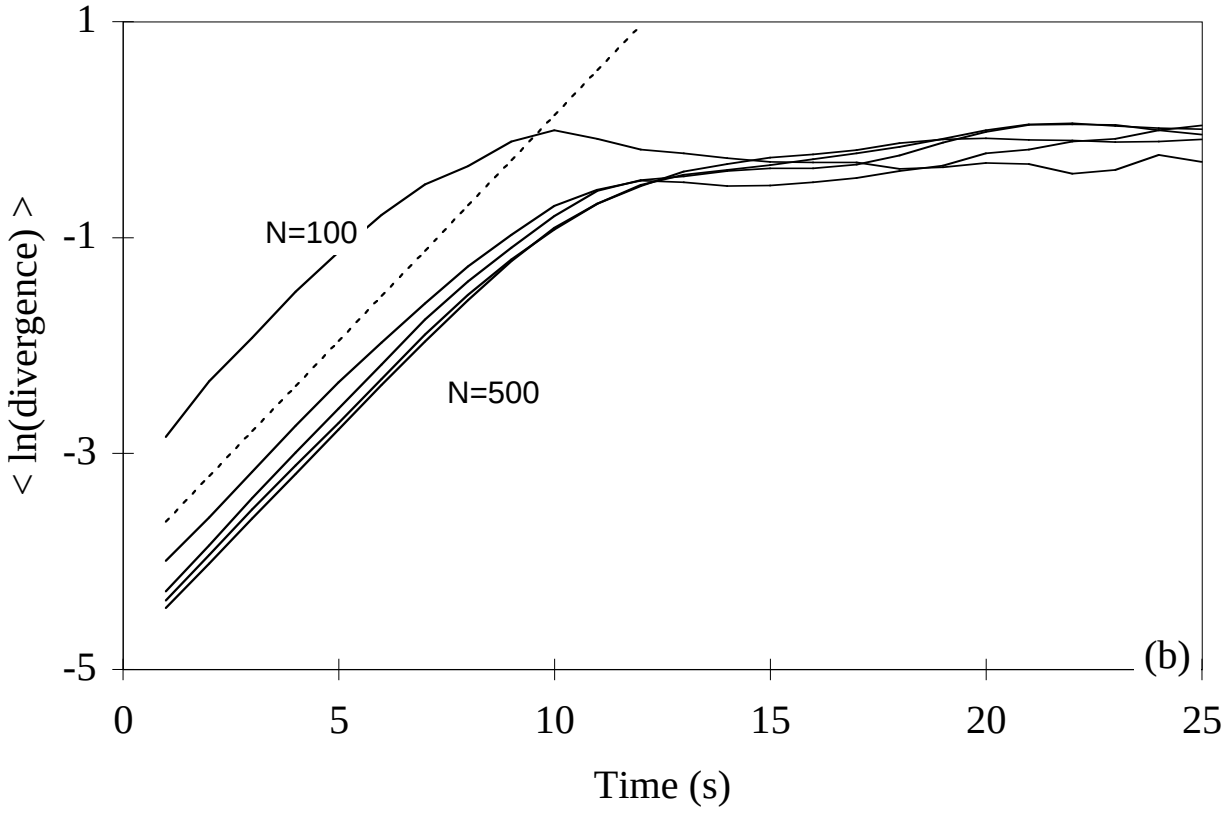
(FIGURE 3D)



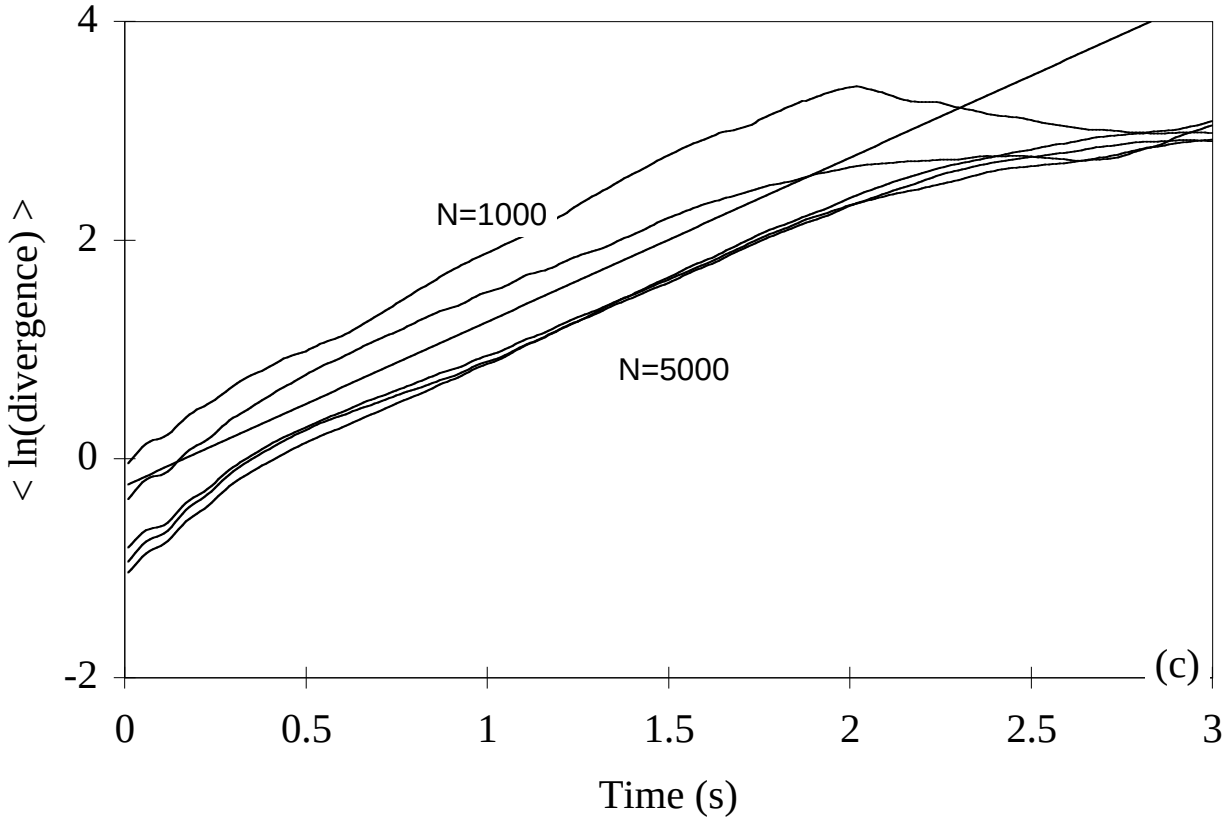
(FIGURE 4A)



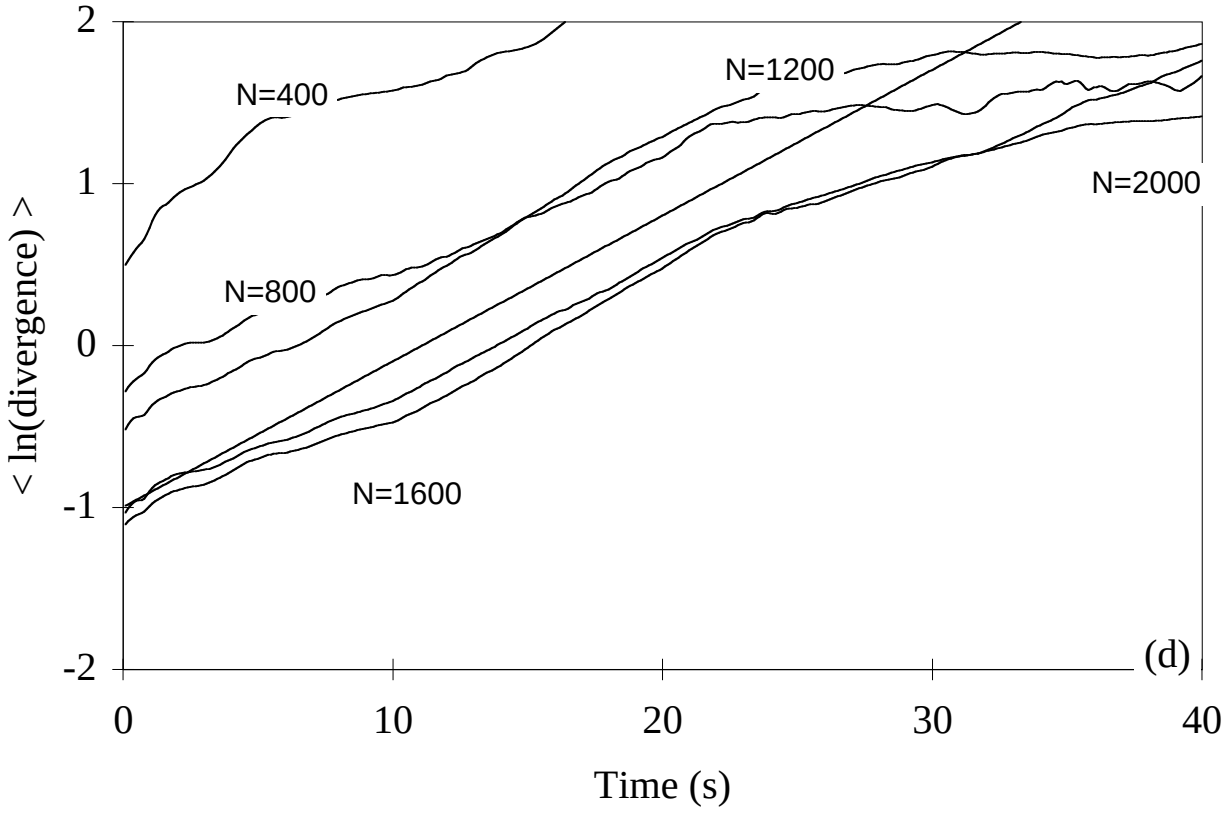
(FIGURE 4B)



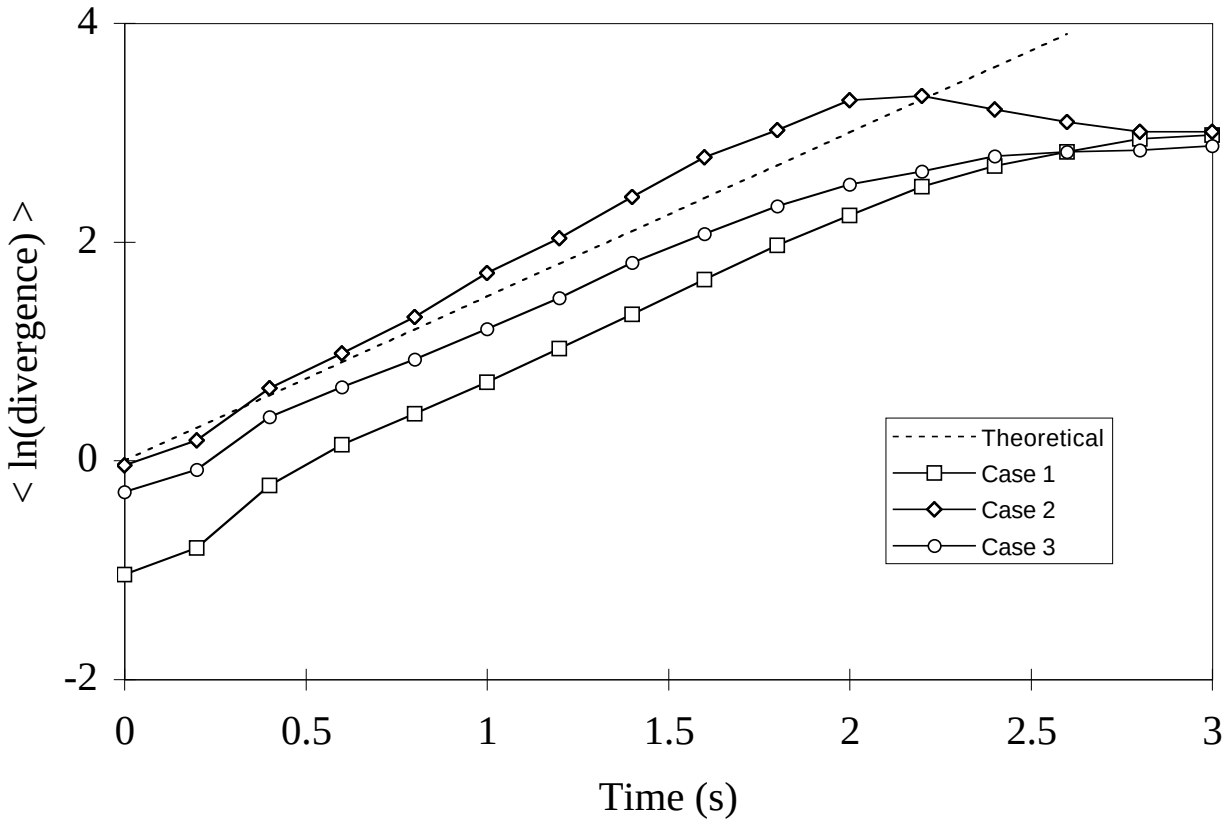
(FIGURE 4C)



(FIGURE 4D)


2
1
0
-1
-2
< ln(divergence) >
N=400
N=1200
N=2000
N=800
N=1600
(d)
0
10
20
30
40
Time (s)

(FIGURE 5)


4
2
0
-2
< ln(divergence) >
0
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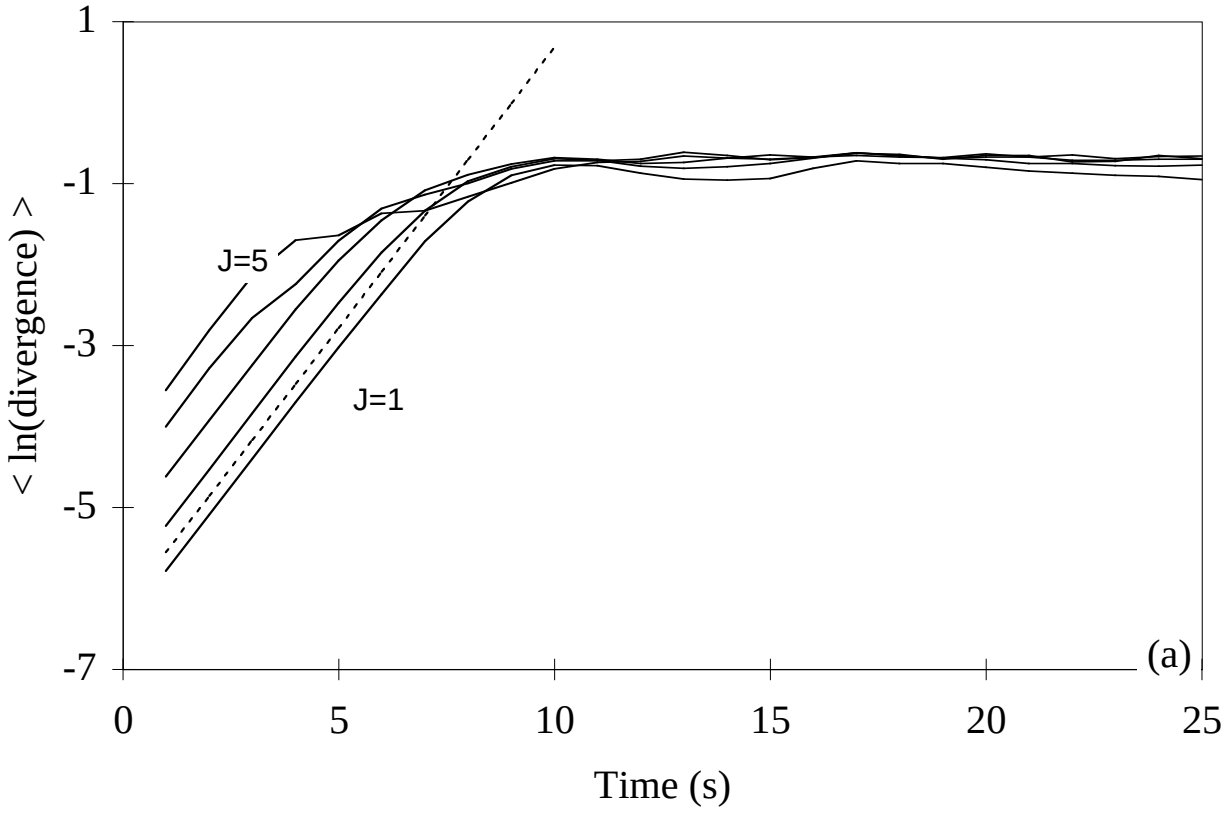
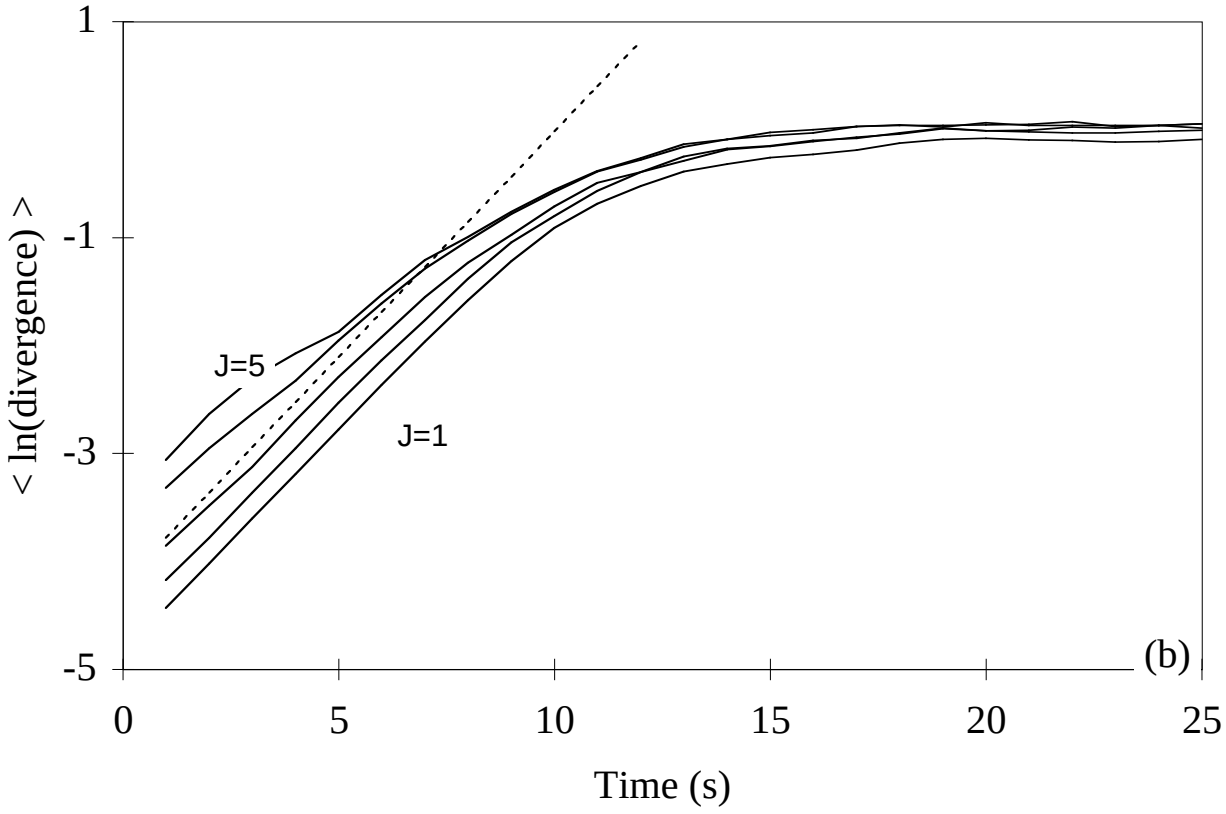
1
1.5
2
2.5
3
Time (s)
Theoretical
Case 1
Case 2
Case 3

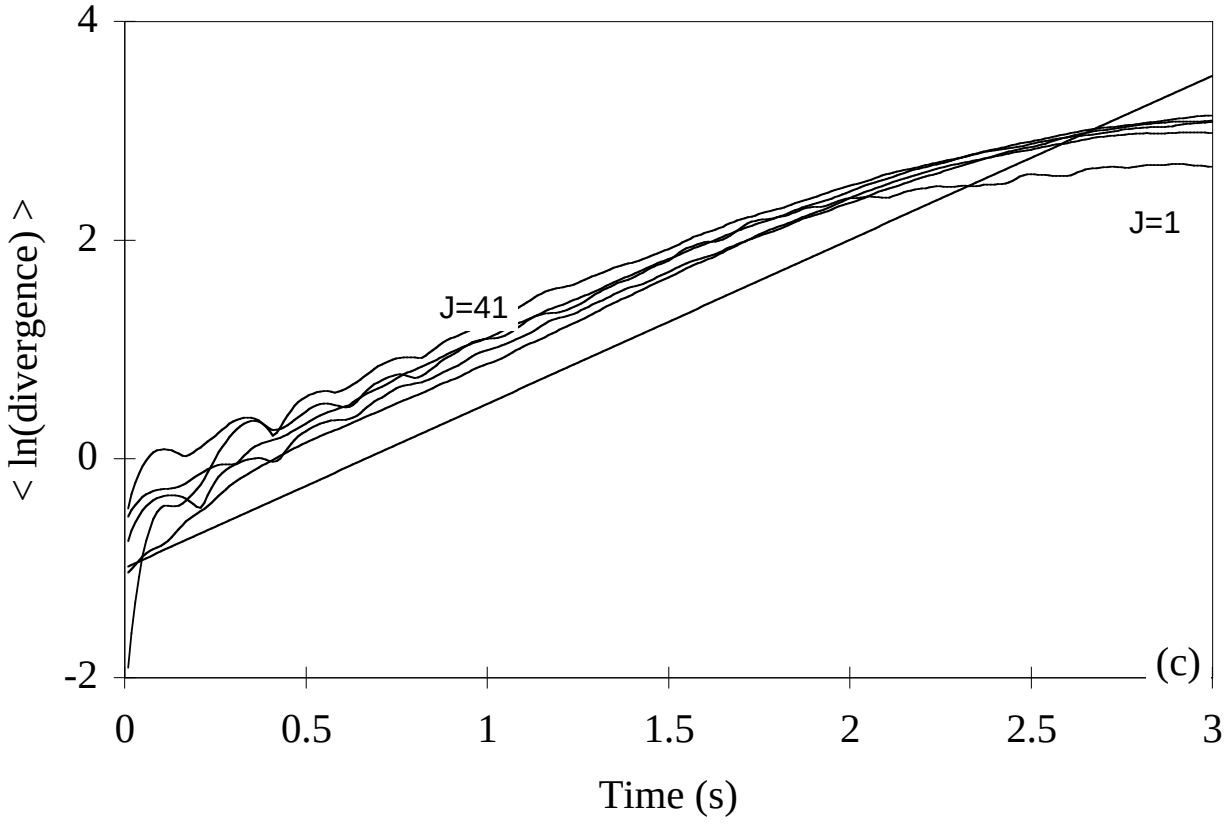
(FIGURE 6A)



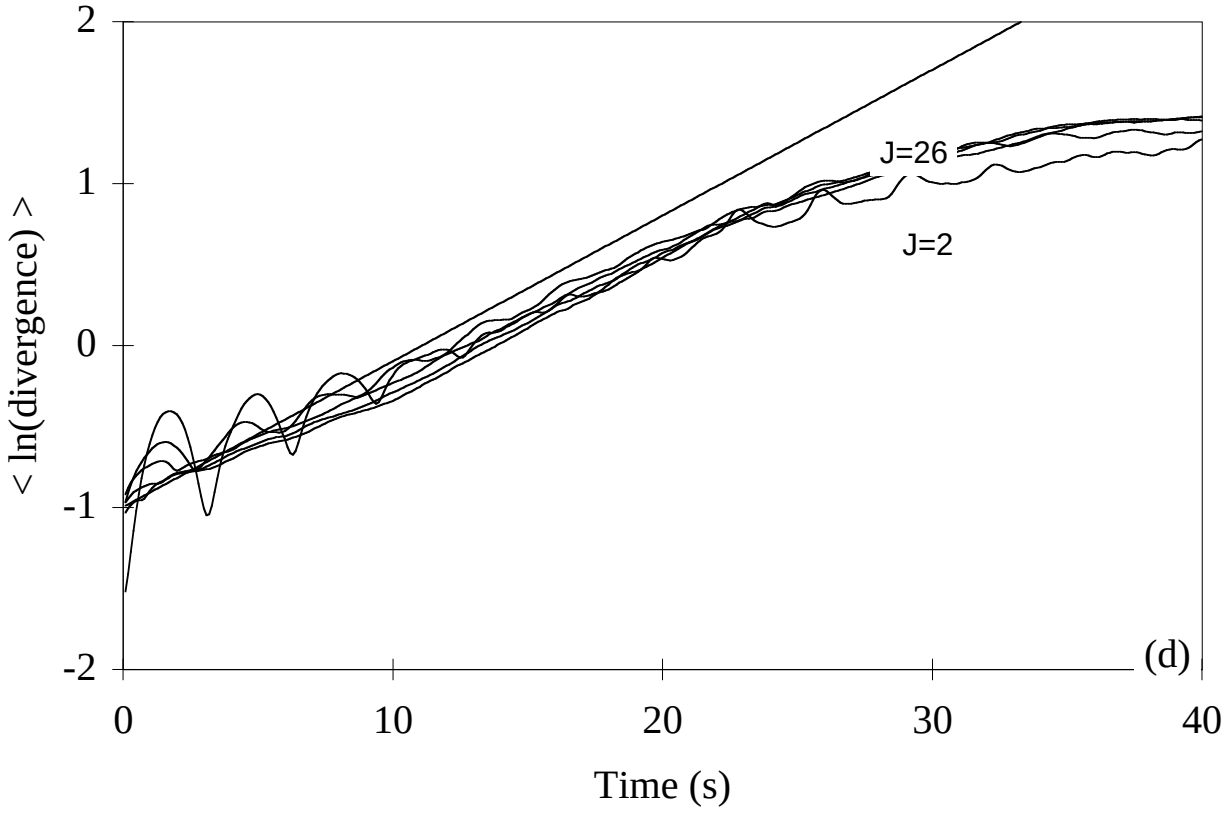
(FIGURE 6B)



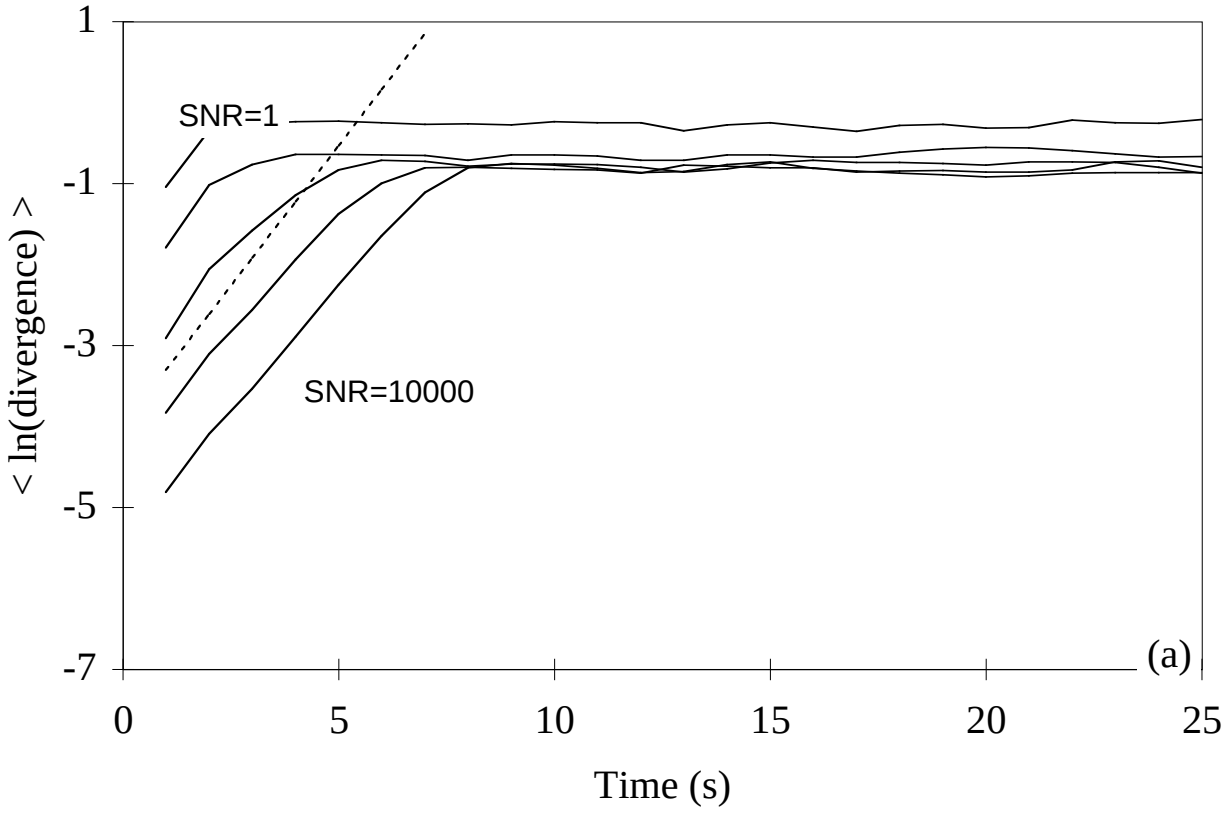
(FIGURE 6C)



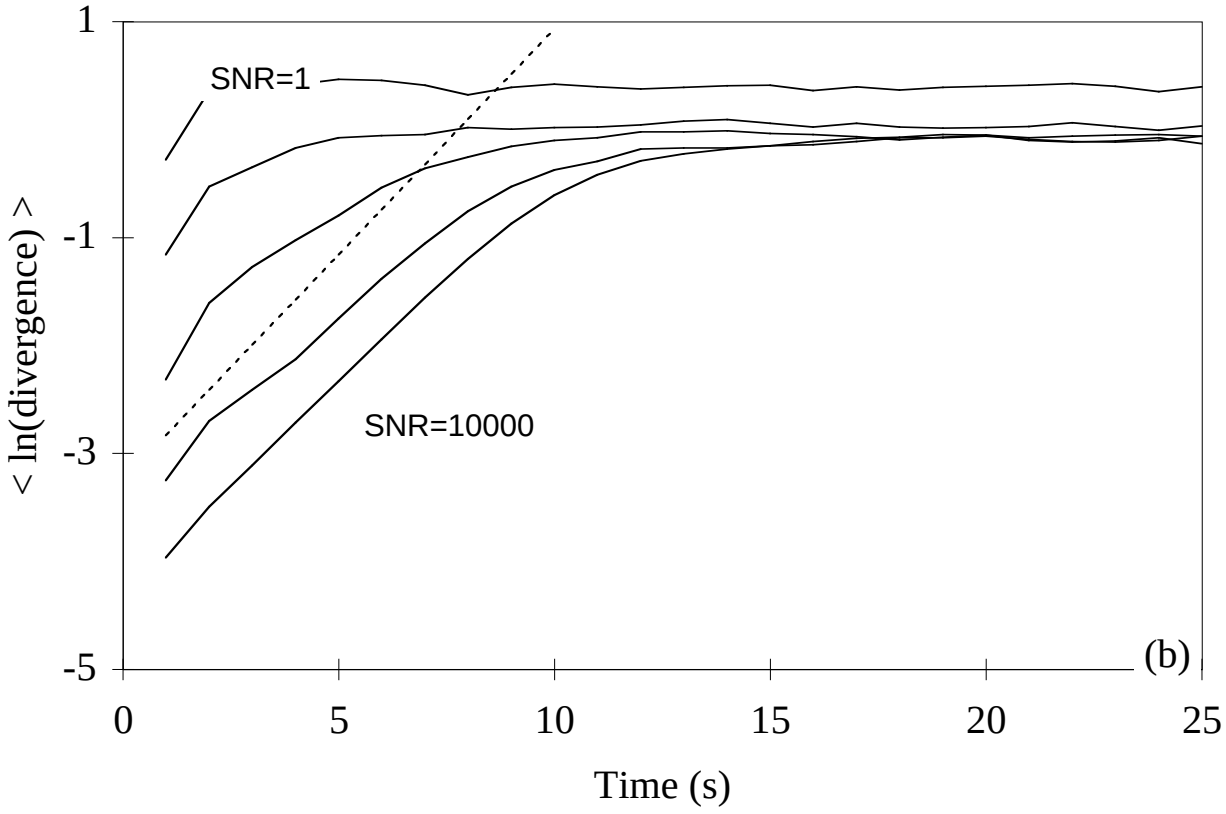
(FIGURE 6D)



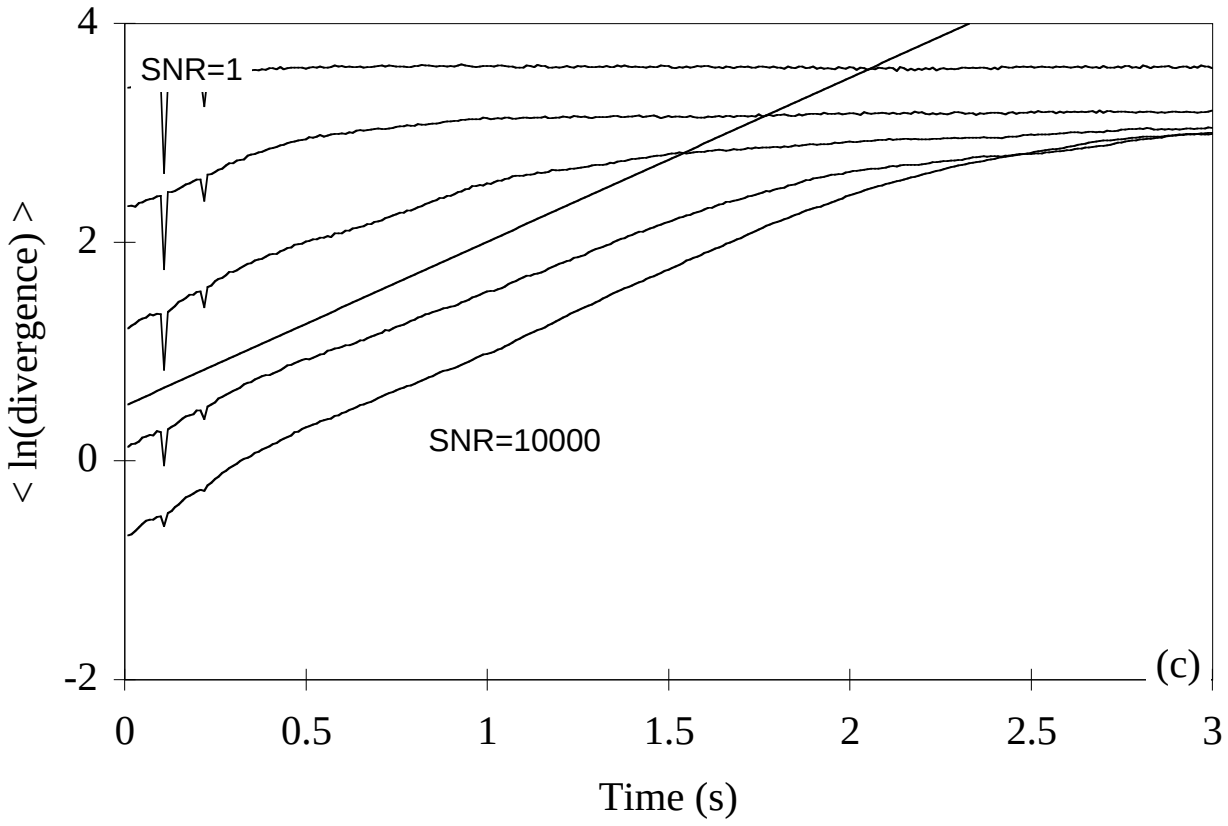
(FIGURE 7A)



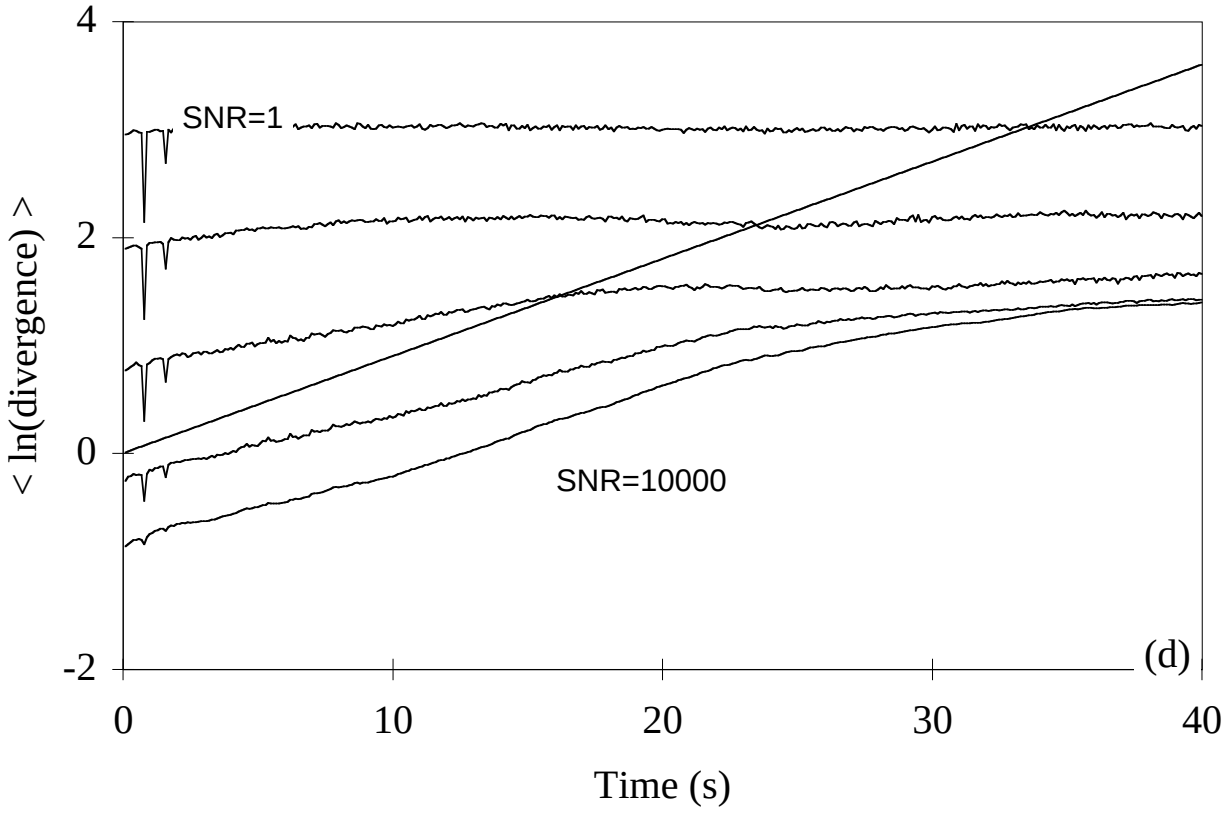
(FIGURE 7B)



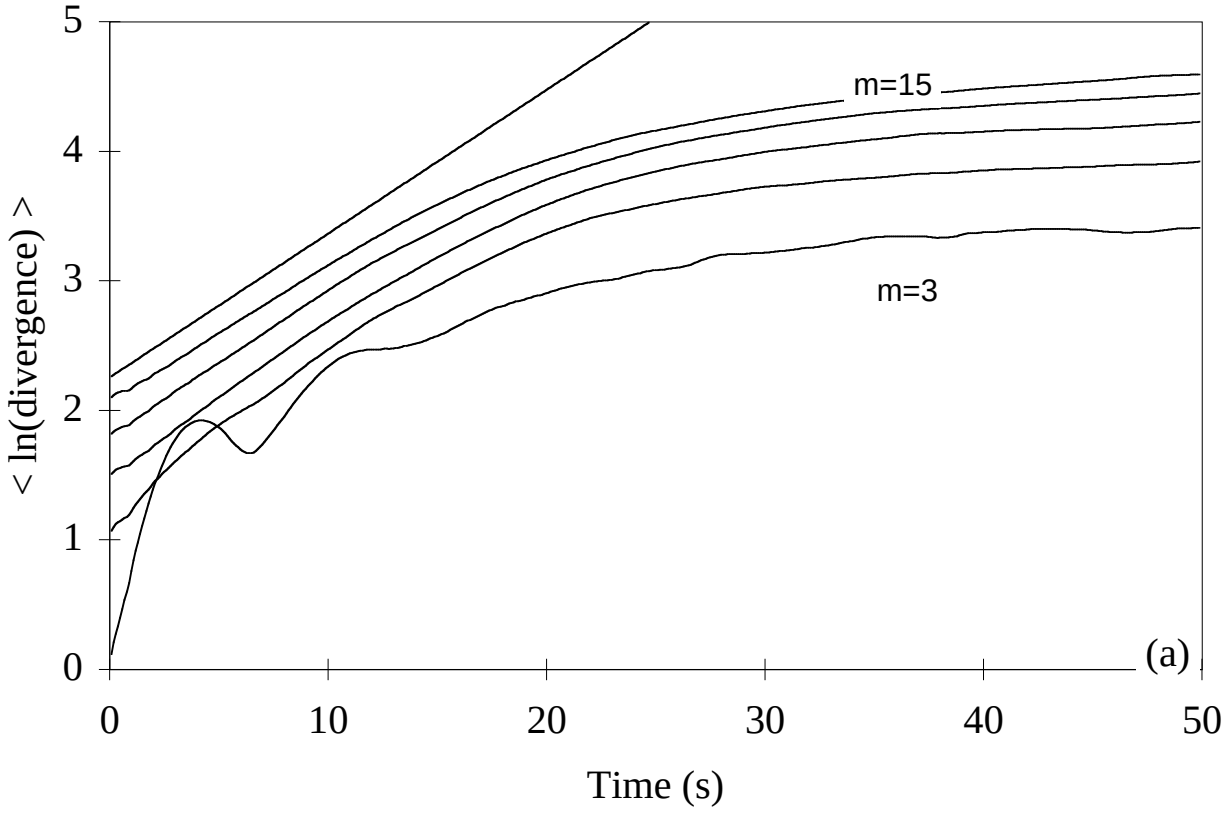
(FIGURE 7C)



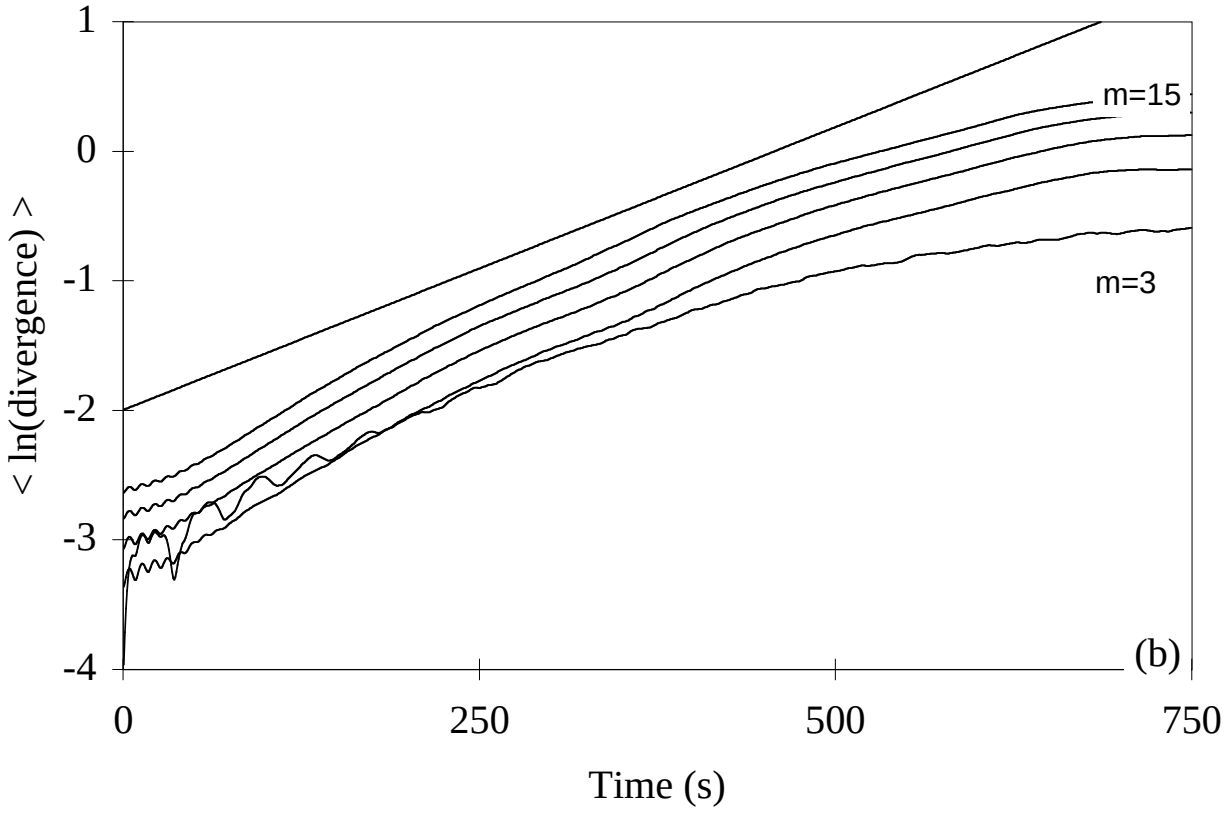
(FIGURE 7D)



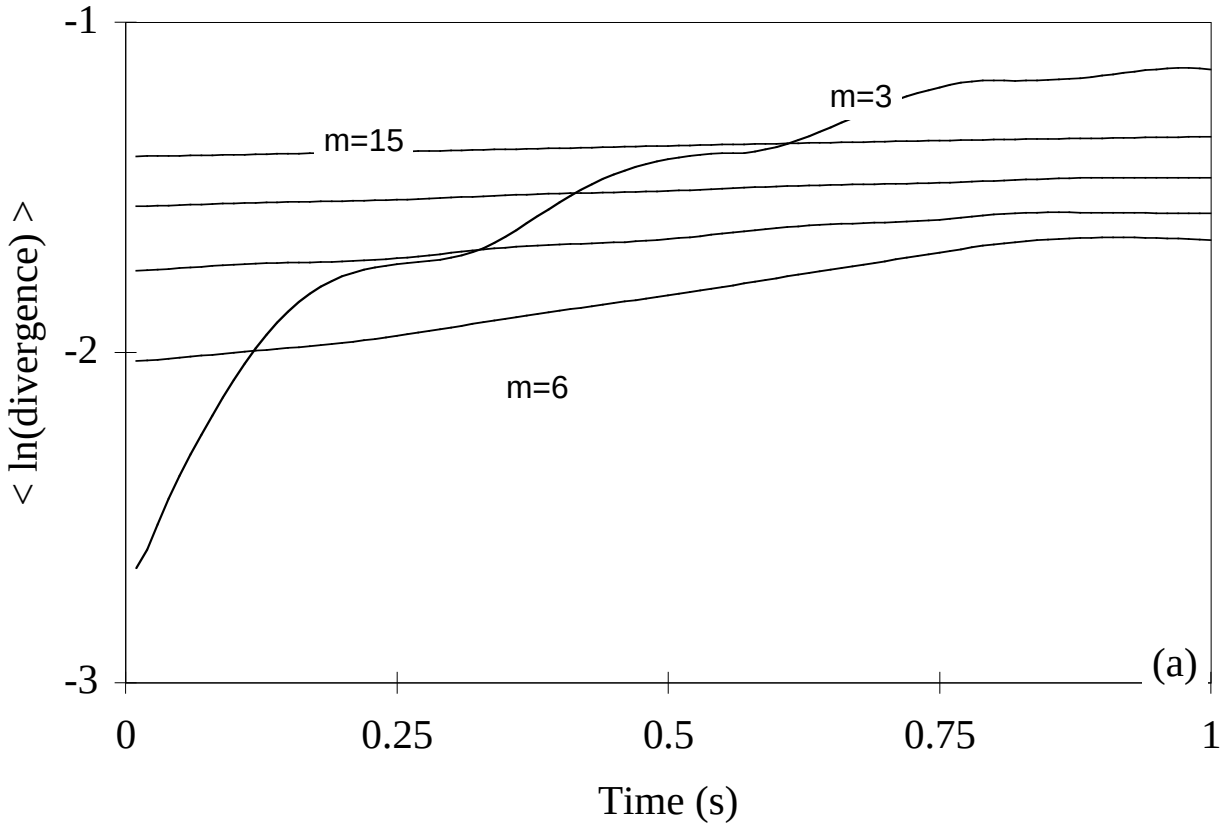
(FIGURE 8A)



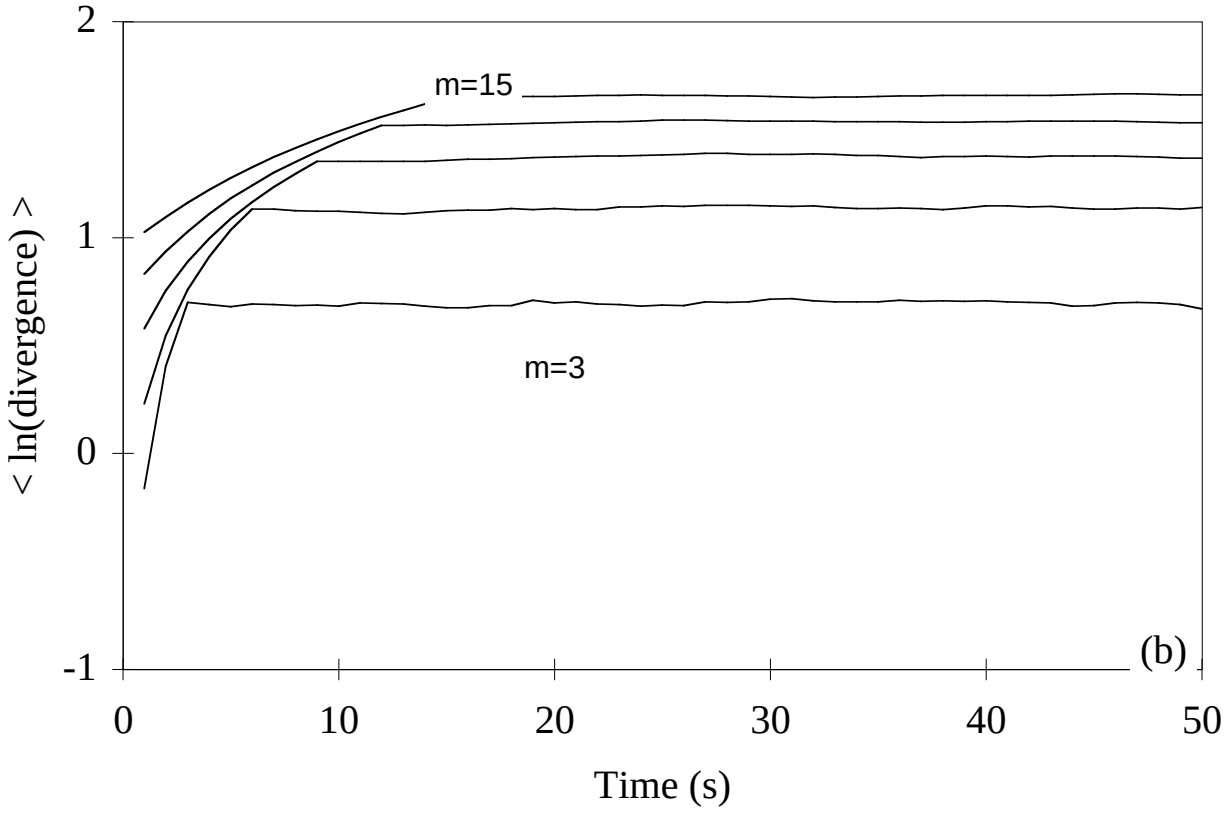
(FIGURE 8B)



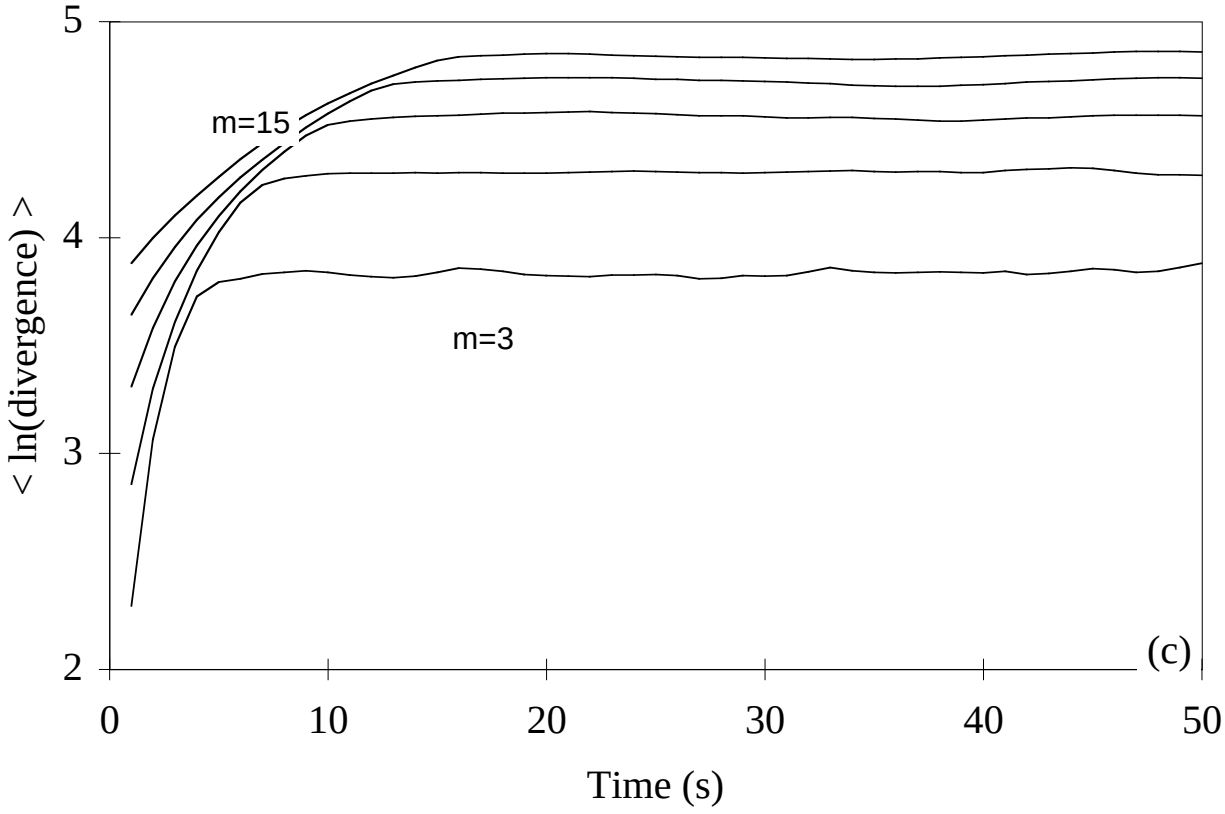
(FIGURE 9A)



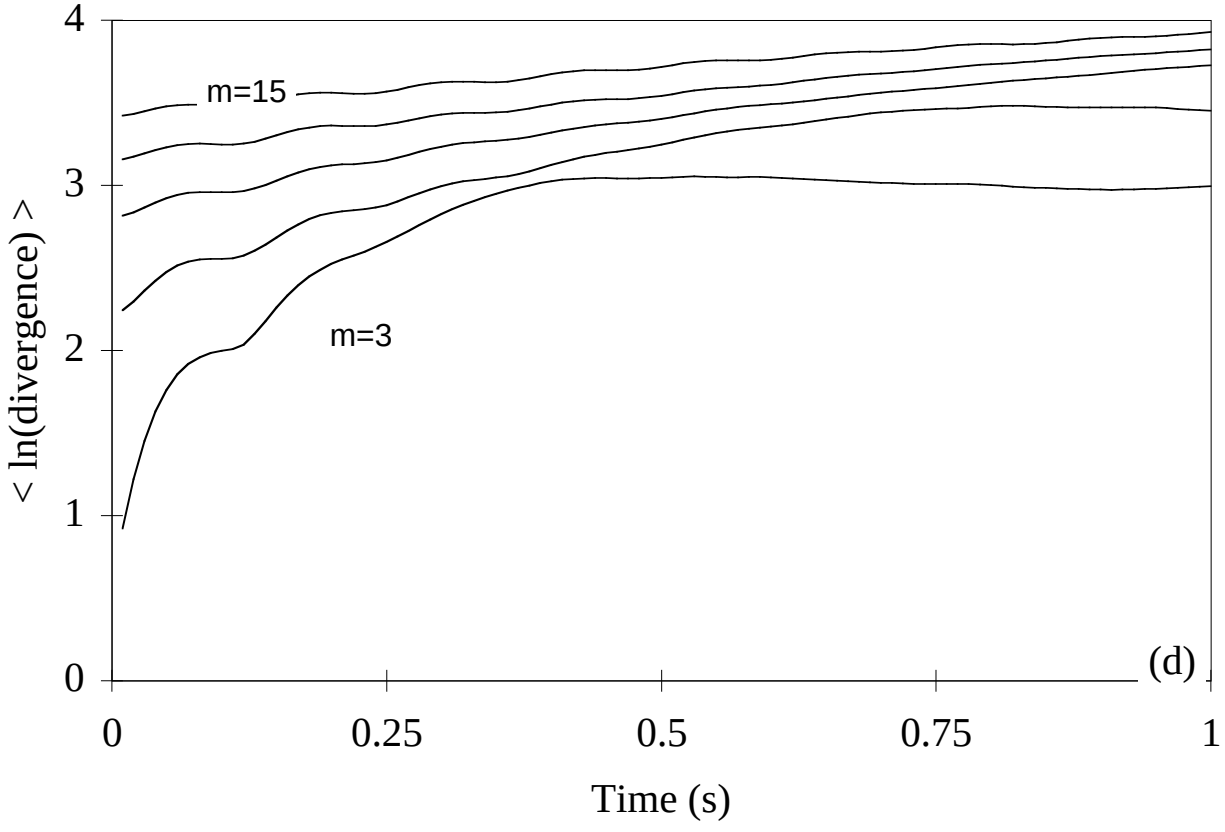
(FIGURE 9B)



(FIGURE 9C)



(FIGURE 9D)



(FIGURE 10)