

Minimum Required Data Length to Reconstruct GC Network

November 10, 2013

1 Task

Compute the minimum required data length when calculating GC. Use IF neural model as an example.

2 Analysis

Suppose there are two random variables x and y .

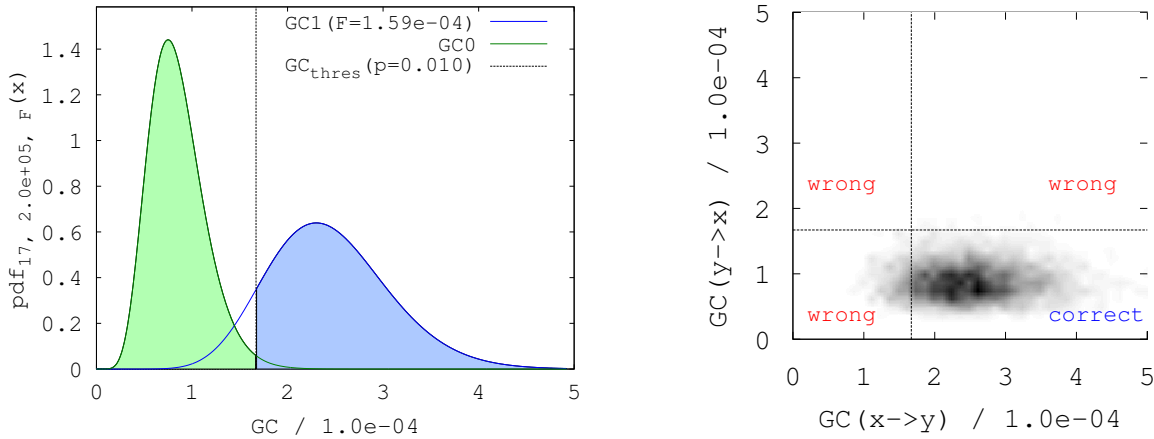
Recall: Distribution of GC obey

$$L \cdot \hat{F}_{x \rightarrow y} \stackrel{a}{\sim} \chi'^2(m, L \cdot F_{x \rightarrow y})$$

in the large L limit ($\stackrel{a}{\sim}$). Where L is number of data samples, $\hat{F}_{x \rightarrow y}$ is calculation value of true GC $F_{x \rightarrow y}$, m is fitting order. χ'^2 is Noncentral chi-squared distribution (see ref. [1] for definition and properties).

For convenience, we denote the probability density function (pdf) of \hat{F} as $\rho_{m,L,F}(x)$.

Now we want to know: for a given L , m and true $F_{x \rightarrow y}$, $F_{y \rightarrow x}$, what's the expected correct rate? Obviously, 25% will be the lowest bound (random guess, without any other knowledge about the two neuron network).



(a) theoretical asymptotic pdf of $\hat{F}_{x \rightarrow y}$ and $\hat{F}_{y \rightarrow x}$ (separately).

(b) density of simultaneous distribution of $\hat{F}_{x \rightarrow y}$ and $\hat{F}_{y \rightarrow x}$. Obtained from 4000 experiment and counting by divide each axis into 40 uniform bins.

Figure 1: GC pdf under one set of typical parameter: $\mu = 1.0\text{kHz}$, $f = 0.012$, $S = 0.01$, using $L = 2.0 \times 10^5$ ($T = 1 \times 10^5\text{ms}$), $m = 17$, the true GC is $F_{x \rightarrow y} \approx 1.592 \times 10^{-4}$ and $F_{y \rightarrow x} \approx 0.006 \times 10^{-4}$ (obtained by $L = 1 \times 10^8$). The black line represent the GC thresholding value (GC_{thres}) that we used to judge whether there is connection or not. Here GC_{thres} satisfies $P(\hat{F}_{y \rightarrow x} < \text{GC}_{\text{thres}}) = 0.01$ and $F_{y \rightarrow x} = 0$ (our null hypothesis), i.e. false positive error rate is 1%.

In order to measure the correctness of GC analysis, here define the correct rate p_{correct} as following: in a two-variable GC test, the true GC $F_{x \rightarrow y} > 0$, $F_{y \rightarrow x} = 0$, the GC test use a fixed p-value (false positive error rate, say 0.01) and fixed fitting order m , sample number L to determine a GC threshold GC_{thres} . Then compare calculated GC \hat{F} to GC_{thres} to guess whether there is connection or not. We denote the ratio between number of correct guesses (network is $x \rightarrow y$) and number of total guess as p_{correct} .

If $\hat{F}_{x \rightarrow y}$ and $\hat{F}_{y \rightarrow x}$ are independent, then the expression for p_{correct} will be very simple:

$$p_{\text{correct}} = \int_0^{F_{\text{thres}}} \rho_{m,L,F0}(F) dF \left(1 - \int_0^{F_{\text{thres}}} \rho_{m,L,F1}(F) dF \right), \quad (1)$$

that is the product of areas of green and blue region in Fig.(1a). Otherwise, we have to count the volumn of lower right part of Fig.(1b).

2.1 Is $\hat{F}_{x \rightarrow y}$ and $\hat{F}_{y \rightarrow x}$ are independent?

Geweke said (Ref.[2]) $\hat{F}_{x \rightarrow y}$ and $\hat{F}_{y \rightarrow x}$ are asymptotically independent. But how ‘‘asymptotically’’.

First, is the asymptotic pdf of $\rho_{m,L,F}(x)$ accurate?

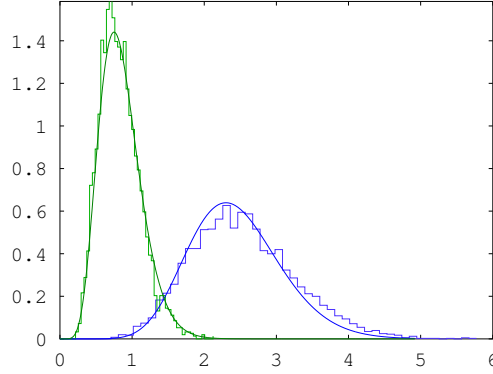
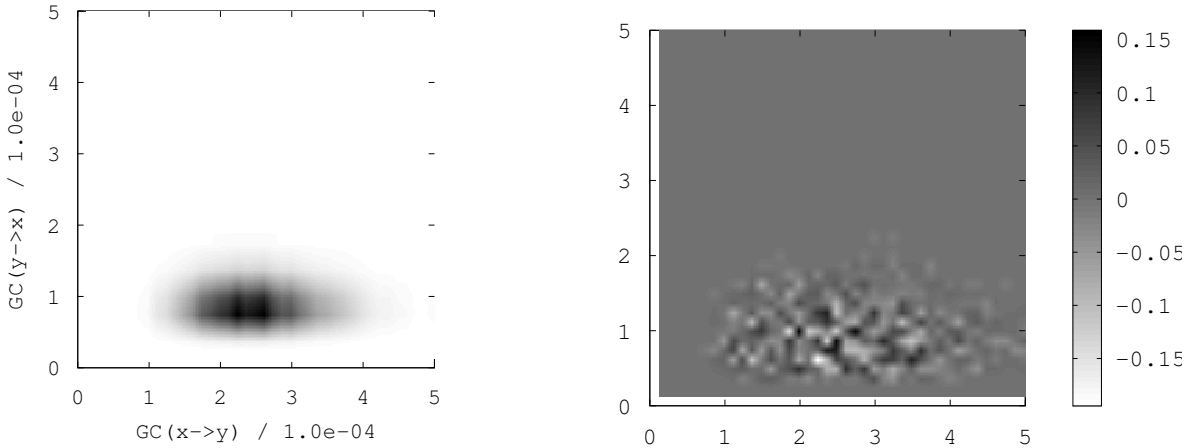


Figure 2: Comparison of statistic data and asymptotic pdf of $\rho_{m,L,F}(x)$. Same parameter as Fig.(1).

Second, by the 4000 GC data point mensioned in Fig.(1b), we can calculate the correlation of $\hat{F}_{x \rightarrow y}$ and $\hat{F}_{y \rightarrow x}$. The result is -0.012 , which can be explained by statistic error ($1/\sqrt{4000} \approx 0.016$).

Further, we compare the joint distribution to the product of marginal distribution.



(a) product of marginal distribution, looks similar to Fig.(1b).

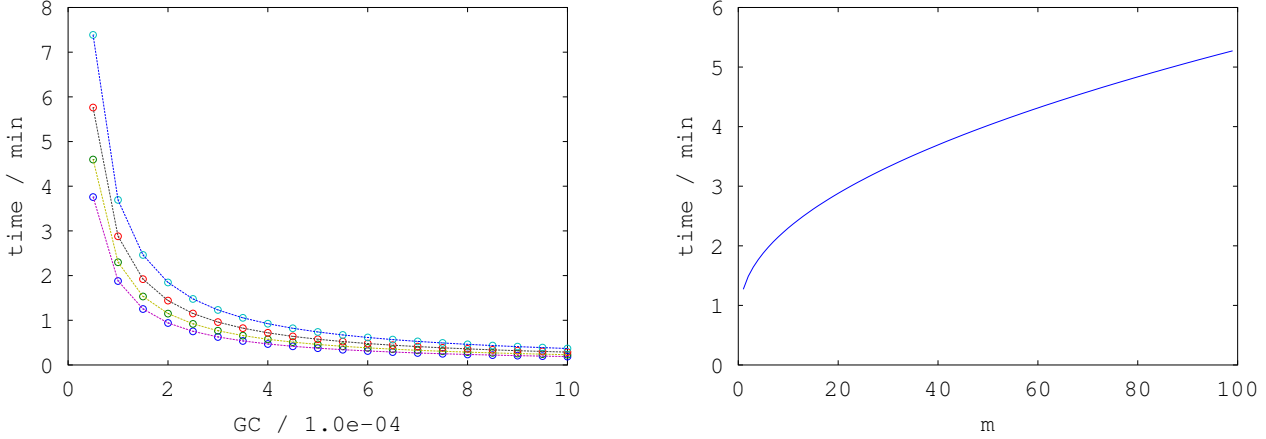
(b) pdf of $(\hat{F}_{x \rightarrow y}, \hat{F}_{y \rightarrow x})$ subtract product of marginal distribution, normalized to distribution peak equals one.

Figure 3

In this parameter (See description of Fig.(1)), the 4000 experiments tell us the correct rate is about 87.1%, while using asymptotic pdf and assume $\hat{F}_{x \rightarrow y}$ and $\hat{F}_{y \rightarrow x}$ are independent, we get 88.9% from Eq.(1) through $m, L, F_{x \rightarrow y}, F_{y \rightarrow x} = 0$ in Fig.(1). Several more experiments are done, they're all matched good enough ($< 5\%$).

2.2 Minimum required data length

From Eq.(1) it's now possible to solve the minimum required data length L_{\min} or data time length T_{\min} (instead of doing a lot of numerical experiments), if m and $F_{x \rightarrow y}$ are known.



(a) Required time length v.s. GC value. Four curves from up to down corresponding to $m = \{5, 10, 20, 40\}$, circle dot is obtained by solving Eq.(1), dashed line is obtained from Eq.(2)(see below).

(b) Required time length v.s. fitting order. Fix GC value to $F_{\text{true}} = 1.0 \times 10^{-4}$

Figure 4: False positive error rate set to 0.01, required correct ratio set to 90%.

In the case of false positive error rate set to 0.01, required correct ratio set to 90%, there is a good approximation of minimum length (relative error of T_{\min} is about 0.1%):

$$T_{\min} \approx \frac{10.00}{F_{\text{true}}} \left(1.153 + \frac{\sqrt{m - 0.513}}{1.917} \right) \Delta t, \quad (m \in \{1, 2, \dots, 100\}) \quad (2)$$

where $1/\Delta t$ is sample rate, $\Delta t = 0.5$ ms in all above cases.

Recall that $F_{\text{true}}/\Delta t \rightarrow \text{const.}$ in the limit $\Delta t \rightarrow 0$, so Eq.(2) tell us that there is no benefit to use small Δt , because in that case $m \propto 1/\Delta t$ which make T_{\min} larger.

Bigger positive error rate will lead to smaller T_{\min} , but effect is limited. e.g. set positive error rate to 0.05 will decrease T_{\min} about 12%, while set positive error rate to 0.005 will increase T_{\min} about 7%.

Now remaining problem is: what is the true GC and corresponding m .
We once have done that:

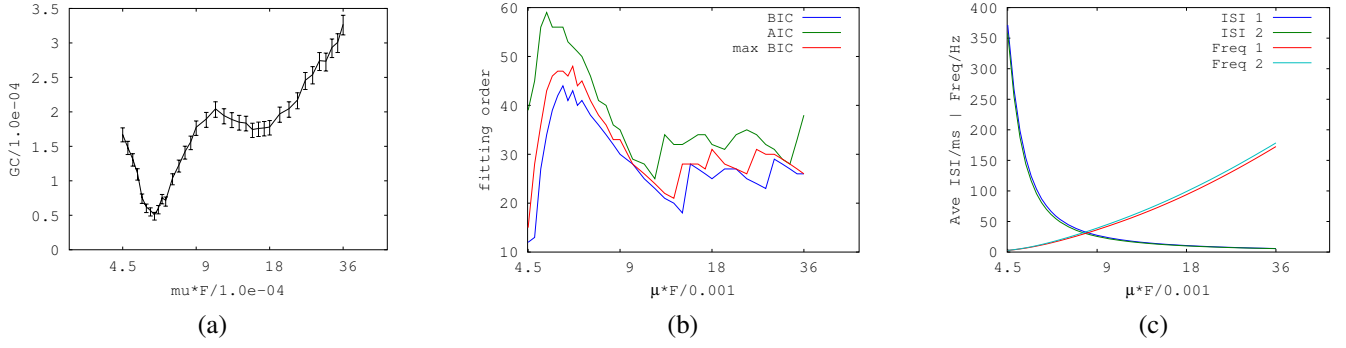


Figure 5: Scan F . $\mu = 0.01$, $S = 0.01$, $T = 1.0 \times 10^4$ sec, $\Delta t = 0.5$ ms

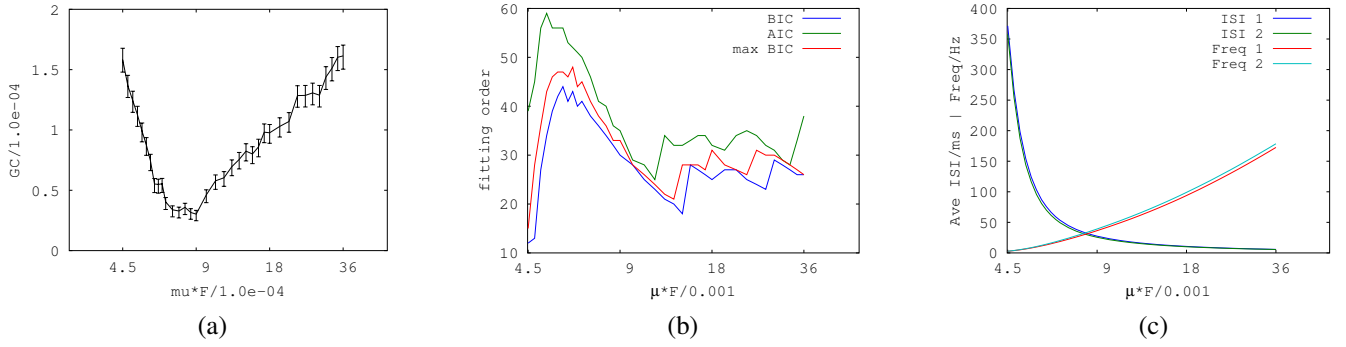


Figure 6: Scan F . $\mu = 0.02$, $S = 0.01$, $T = 1.0 \times 10^4$ sec, $\Delta t = 0.5$ ms

Importing these data to Eq.(2) we get:

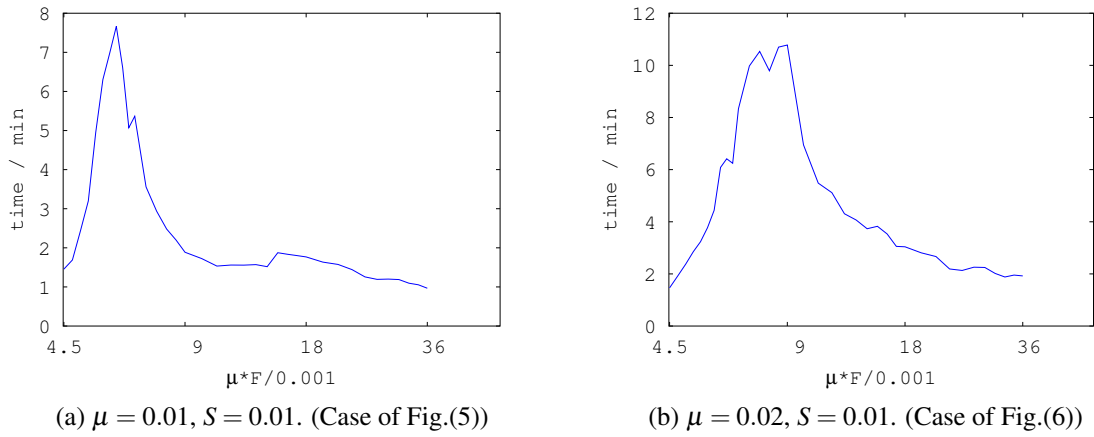


Figure 7: Required time length

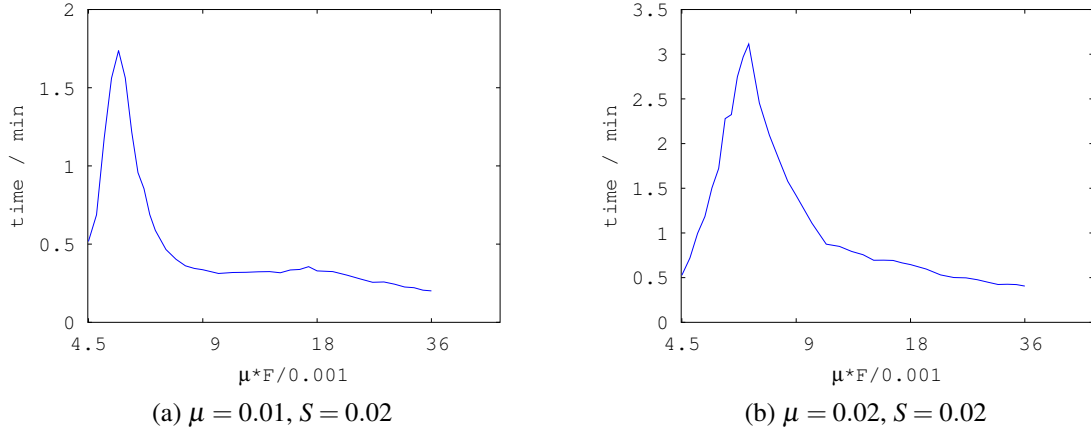


Figure 8: Required time length

As what we expected, twice the S , roughly quarter the required time.

Case of using spike train:

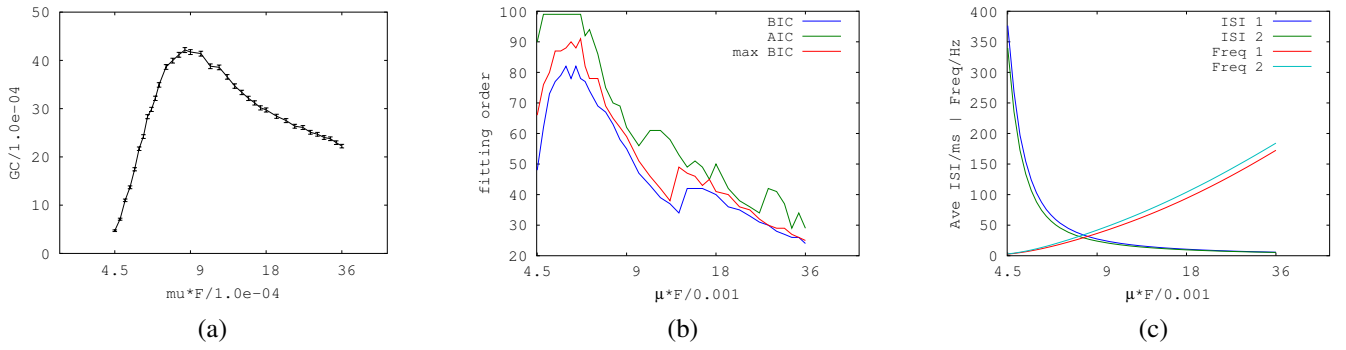


Figure 9: Using spike train. Scan F . $\mu = 0.01, S = 0.02, T = 1.0 \times 10^4 \text{ sec}, \Delta t = 0.5 \text{ ms}$

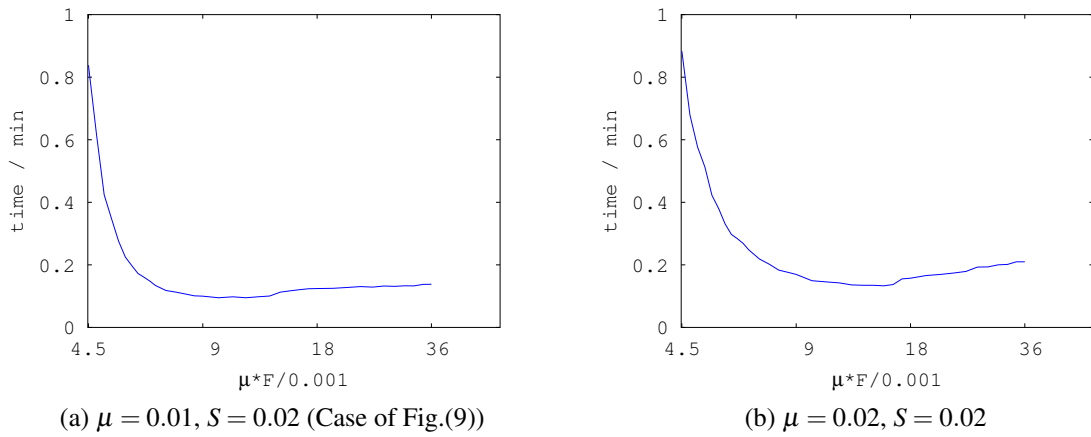


Figure 10: Required time length. (Use spike train)

Note: $EPSP \approx 107 S \text{ mV}$ or $EPSP \approx 7 S$ in the normalized unit.

3 Big ISI Case

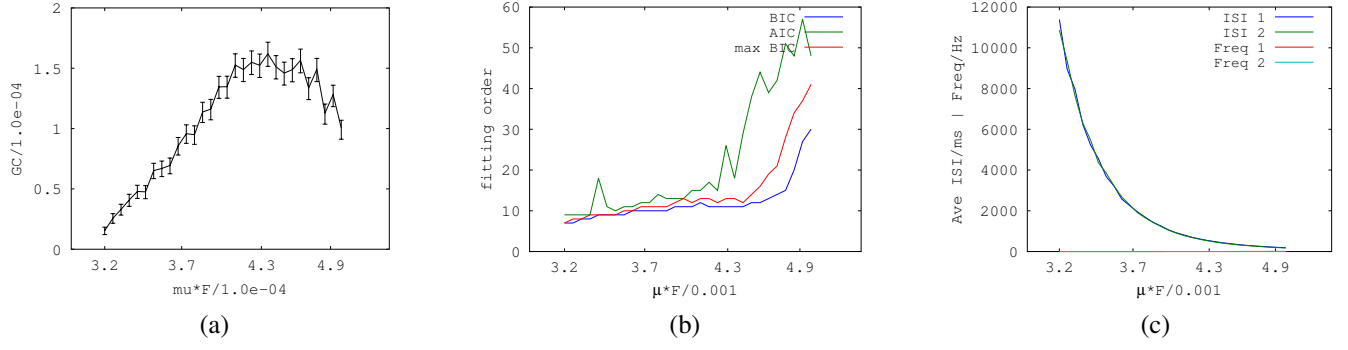


Figure 11: Scan F . $\mu = 0.01, S = 0.01, T = 1.0 \times 10^4$ sec, continue Fig.(5)

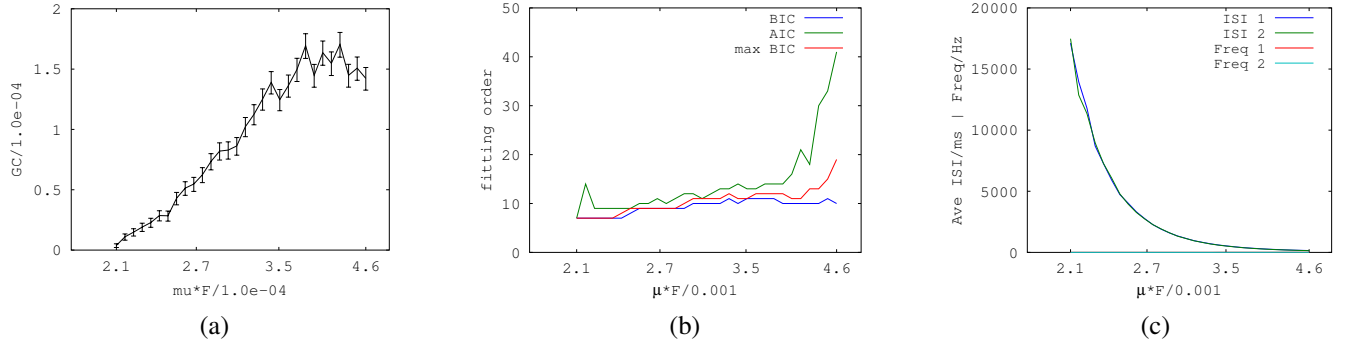


Figure 12: Scan F . $\mu = 0.02, S = 0.01, T = 1.0 \times 10^4$ sec, continue Fig.(6)

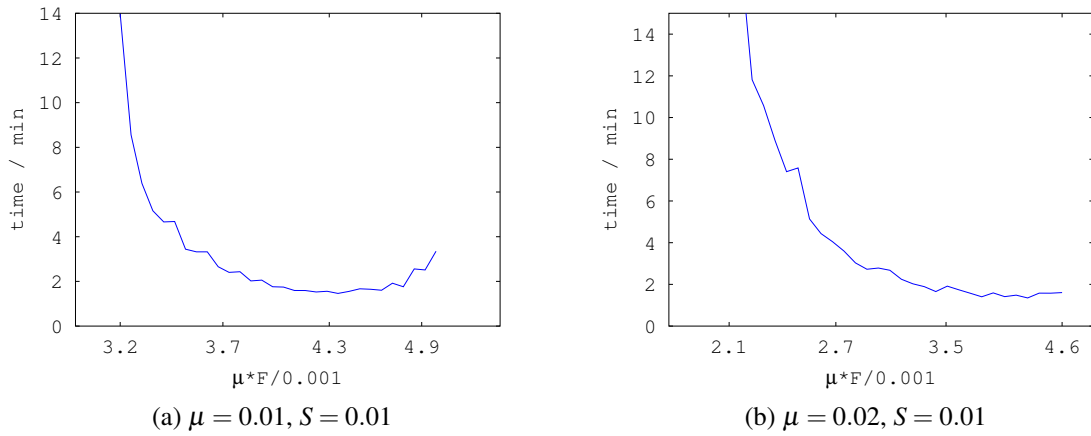


Figure 13: Required time length

References

- [1] http://en.wikipedia.org/wiki/Non-central_chi-square_distribution

- [2] Measurement of Linear Dependence and Feedback Between Multiple Time Series, John Geweke, Journal of the American Statistical Association, Vol.77, No.378 (1982)