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Note on Steady-state Approximation of Enzyme Kinetics

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Recently, it has been gradually believed that enzymatic reactions generally involve several enzyme-substrate complexes in their pathways. For such the reactions, an approximation technique, for instance, steady-state approximation or Michaelis-Menten approximation, may not be simply applied to analyze the experimental data. In the present study, a three-step scheme of enzymatic reaction was subjected as a typical example of enzymatic reaction to the digital computer simulation to establish the practical experimental conditions which allow the proper application of approximation technique in handling the experimental data.

INTRODUCTION

It has been well recognized that the deterministic rate equation of enzymatic reaction has the form of simultaneous nonlinear differential equation, and some mathematical skillful tasks are required for obtaining the analytical solution of the rate equation. On the practical kinetics for enzymatic reactions, therefore, a suitable approximation technique such as steady-state approximation has been generally adopted to avoid the mathematical difficulties. The application of approximation technique will depend upon the experimental conditions which had been used for measurement of the behavior of the reaction. The rate parameters such as K_m or V_{max} calculated by the approximation technique may provide meaningless values and conduct one's judgement to erroneous direction unless the experimental conditions were properly assembled.

The limitation on the use of such an approximation technique had been clearly revealed by several investigators for only the case of relatively simple enzymatic reaction as typically represented by Michaelis-Menten scheme:

$$E+S \rightleftharpoons ES \rightarrow E+P$$

In recent years, it came to know that general enzymatic reactions involve a sequence of the enzyme-substrate complexes as shown by following equation:

$$E+S \rightleftharpoons ES. \rightleftharpoons ES. \rightleftharpoons \cdots \rightleftharpoons ES. \rightarrow E+P$$

Thus, the progress of enzyme chemistry enabled the description of detailed parts of the enzymatic reaction scheme. Now, enzymatic reactions are discussed in the submolecular level. Contrary to the above situation, practical kinetic analyses seem to be performed frequently with ignoring the fine mechanisms of enzymatic reactions; in practical experiments, the data on complicated reaction are simply delt

with approximation technique by assuming rather simple reaction scheme without regard to the limitations of the approximation techniques.

Walter et *al.* (1964) have clarified the limitations on the application of the steady-state approximation for three-step enzymatic reaction by means of analog computer simulation. However, rate constants assumed in the simulation, that control the rates of formation and isomerization of enzyme-substrate complexes, were quite different from those observed on the real enzymatic reactions. Heineken et *al.* (1967 a, 1967 b) studied further the behaviors of three-step enzymatic reaction. However, the results did not seem to give clear information for the establishment of the practical experimental conditions.

The present study was designed to clarify the experimental conditions for enzymatic reaction that allow the application of the steady-state approximation on the analysis of enzymatic reaction involving the several enzyme-substrate complexes. The authors set up the three-step enzymatic reaction:

$$E+S \rightleftharpoons ES_1 \rightleftharpoons ES_2 \rightarrow E+P$$

and the behaviors of this reaction were simulated by digital computer in real time scale. The rate constants were suitably chosen with referring to those observed on the real enzymatic reactions.

COMPUTATION

The three-step reaction scheme may be represented by following equation (scheme):

$$E + S \underset{k_{-1}}{\overset{k_{+1}}{\rightleftharpoons}} ES_1 \underset{k_{-2}}{\overset{k_{+2}}{\rightleftharpoons}} ES_2 \xrightarrow{k_{+3}} E + P$$
 (1)

where E, S, ES and P represent the enzyme, substrate, enzyme-substrate complex and the product, k_{+1} and k_{-1} are the constants for the formation of the complex ES,, k_{+2} and k_{-2} are for the isomerization of ES, to ES,, and k_{-3} for the formation of the product. E(t), S(t), $X_1(t)$, $X_2(t)$ and P(t) represent the concentrations of enzyme, substrate, ES₁, ES, and product respectively, at the reaction time t. E(O), S(O), $X_2(O)$, and P(9) are the initial concentrations of the reaction species.

The rate equation of the three-step reaction scheme in form of simultaneous differential equation may be written as:

$$\begin{cases} \frac{dE(t)}{dt} = -k_{+1} \cdot E(t) \cdot S(t) + k_{-1} \cdot X_1(t) + k_{+3} \cdot X_2(t) \\ \frac{dS(t)}{dt} = -k_{+1} \cdot E(t) \cdot S(t) + k_{-1} \cdot X_1(t) \\ \frac{dX_1(t)}{dt} = k_{+1} \cdot E(t) \cdot S(t) - (k_{-1} + k_{+2}) \cdot X_1(t) + k_{-2} \cdot X_2(t) \\ \frac{dX_2(t)}{dt} = k_{+2} \cdot X_1(t) - (k_{-2} + k_{+3}) \cdot X_2(t) \end{cases}$$
(2)

$$\frac{dP(t)}{dt} = k_{+3} \cdot X_2(t)$$

Since $E(t) \leq E(0)$, $S(t) \leq S(0)$, $X_1(t) \leq E(0)$, $X_2(t) \leq E(0)$, and $P(t) \leq S(0)$, the concentrations in the rate equation were converted into the relative value. Then the equation (2) may be replaced by equation (3):

$$\begin{cases}
\frac{dE'(t)}{dt} = -k_{+1}S(0) \cdot E'(t) \cdot S'(t) + k_{-1} \cdot X_{1}'(t) + k_{+3}X_{2}'(t) \\
\frac{dS'(t)}{dt} = -k_{+1}E(0) \cdot E'(t) \cdot S'(t) + k_{-1}\frac{E(0)}{S(0)}X_{1}'(t) \\
\frac{dX_{1}'(t)}{dt} = k_{+1}S(0) \cdot E'(t) \cdot S'(t) - (k_{-1} + k_{+2})X_{1}'(t) + k_{-2}X_{2}'(t)
\end{cases}$$

$$\frac{dX_{2}'(t)}{dt} = k_{+2}X_{1}'(t) - (k_{-2} + k_{+3})X_{2}'(t) \\
\frac{dP'(t)}{dt} = k_{+3}\frac{E(0)}{S(0)}X_{2}'(t)$$
(3)

where

$$E'(t) = E(t)/E(0)$$
S' (t) = S(t) /S(O),
 $X_1'(t) = X_1(t) /E(O)$
 $X_2'(t) = X_2(t) /E(0)$
 $P'(t) = P(t) /S(0)$

On the simulation, the initial concentrations were assumed as ;

$$E(0) = 1.0 \times 10^{-6} \text{M}.$$

 $S(0) = \mathbf{a} \times E(0)$
 $X_1(0) = X_2(0) = \mathbf{P}(0) = 0$

where a is a fixed number $(a=10, 20, \ldots, 100)$.

The values of rate constants were chosen as,

$$10^4 < k_{+1} < 10^6 (\mathrm{M}^{-1} \, \mathrm{sec}^{-1}), \ 1 < k_{-1} < 10 (\mathrm{sec}^{-1})$$

 $10^3 < k_{+2} < 10^4 (\mathrm{sec}^{-1}), \ 10 < k_{-2} < 10^2 (\mathrm{sec}^{-1})$
 $0 < k_{+3} < 0.2 (\mathrm{sec}^{-1})$

These values were chosen with referring to the experimentally obtained values on enzymatic reaction of lysozyme (Sykes, 1969 a; Holler *et al.*, 1969; Chipman, 1971) and chymotrypsin (Sykes, 1969 b).

The numerical solution of the simultaneous differential equation was obtained by MRKGM subprogram. The maximum initial pitch in solving the differential equation was 0.06 sec. The MRKGM is a minor modification of SRKGD that is the FACOM SSL program for the numerical calculation of differential equation (Runge-Kutta-Gill method). When an overflow occurs at an operational register of computer during the calculation (a), and when the solutions (calculated values of concentration) are negative (b), the pitch is reduced to one-fifth of the original one and the calculation is repeated (recalculation) by this revised sub-

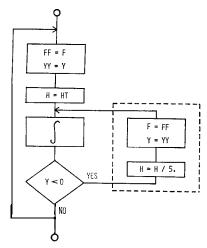


Fig. 1. Flow chart for avoiding obstruction in computation of differential equation (part of MRKGM).

program MRKGM. The flow chart of re-calculation part is shown in Fig. I. F and Y are differential value and integrated value (solution) respectively, FF and YY are memories to reserve the values of F and Y calculated before the occurrence of the obstruction ((a) or (b)), respectively. H is the corrected pitch and HT is the initial pitch. Symbol \int represents the operation by which the simultaneous differential equation is solved.

The program MRKGM has an advantage of that the obstruction caused by a wrong selection of the initial pitch can be avoided. Therefore, this program seems to be one of the simple and available programs of setting an unfeasibility caused by the exceedingly great difference between rate constants contained in the same simultaneous equation.

On the three-step enzymatic reaction, $dX_1(t)/dt=0$ and $dX_2(t)/dt=0$ are the essential conditions for the steady-state of the reaction. In practical reaction, the steady-state in the mathematical definition will not be attained, even if the reaction takes place under the most complete conditions. In the present study, however, it was assumed for convenience that the steady-state is attained when the trajectory of $X_2(t)$ with respect to the reaction time t is visibly judged to be parallel to the horizontal axis. Since the calculated values of $X_1(t)$ contain the computational errors to some extent that are due to its very small value, it was not used for the judgement of the steady-state. Furthermore, the time periods in which the steady-state approximation could be adopted is defined as that in which the values of $X_2(t)$ deviate only 5 % up- or downward from the linear trajectory. This time periods is called steady-state time and the time ratio is difined by the percent ratio of steady-state time to the entire reaction time.

All computations were performed by FACOM 230-75 Digital Computer and some FACOM SSL in the Computer Center, Kyushu University.

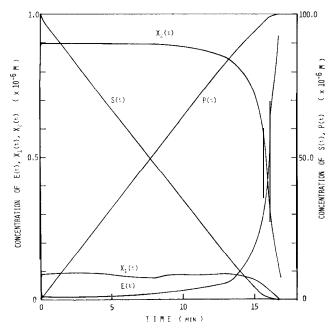


Fig. 2. Calculated time-courses of enzymatic reaction with scheme (1). Rate constants used for simulation are: k_{+1} =5×10⁵ M⁻¹ sec⁻¹, k_{-1} =5 sec⁻¹, k_{+2} =10³ sec⁻¹, k_{-2} =10² sec⁻¹, k_{+3} =0.12 sec⁻¹. Molar ratio of S(0)/E(0) is assumed to be 100.

RESULTS

Relation between k_{+3} and steady-state

A typical calculated time-course of three-step enzymatic reaction is shown in Fig. 2. The concentration of ES, was held in nearly constant value in the time periods from 0.1 sec to 8 min. A considerably irregular shape of $X_1(t)$ was due to the errors in the computation. The changes in S(t) and P(t) exhibit the linearity until 13 min of the reaction time where about 85% of the substrate had been converted to the product. The time-ratio on steady-state approximation defined before was roughly estimated to be 74%.

The value of k_{+3} controls the overall rate of the enzymatic reaction. The change in the k_{+3} value under the fixing of the other rate constants did not affect the time-ratio as shown in Table 1.

Table 1. Effect of k_{+3} value on time ratio.

k_{+3} (sec ⁻¹)	Time ratio (%)
0.12	74
0.16	73
0.20	75

A typical example in the case where the steady-state was not attained is shown in Fig. 3. All the rate constants except for k_{+3} were common for Figs. 2

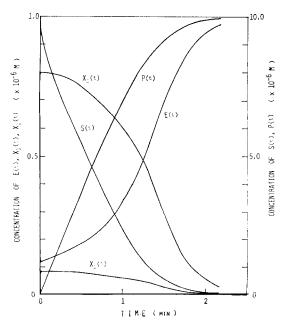


Fig. 3. Calculated time-courses of the reaction with scheme (1). Rate constants are: $k_{+1}=5\times10^5~\rm M^{-1}~sec^{-1}$, $k_{-1}=5~sec^{-1}$, $k_{+2}=10^3~sec^{-1}$, $k_{-2}=10^2~sec^{-1}$, $k_{+3}=0.16~sec^{-1}$. Molar ratio of S(0)/E(0) is assumed to be 10.

and 3. In the case of Fig. 3, S(0)/E(0) was assumed to be 10, differing from that for Fig. 2.

The decrease in the k_{+3} value in legend of Fig. 3 did not lead to the appearance of steady-state as shown in Fig. 4; the time-course on X_{+3} 0 with $k_{+3}=0.16\,\mathrm{sec^{-1}}$ 1 is simply extended along the time axis. Thus, the value of k_{+3} does not correlate to the appearance of steady-state or the time-ratio.

S(0)/E(0) Ratio and steady-state

It is obvious from the time-courses of $X_2(t)$ shown in Figs. 2 and 3 that the S(0)/E(0) ratio plays the vital role in the appearance of the steady-state in the enzymatic reaction. The time-courses of $X_2(t)$ with changing the S(0)/E(0) ratio are shown in Fig. 5. The values of rate constants used for the calculation were $k_{+1}=5\times10^5~\mathrm{M^{-1}\,sec^{-1}}, k_{-1}=5~\mathrm{sec^{-1}}, k_{+2}=10^3~\mathrm{sec^{-1}}, k_{-2}=10^2~\mathrm{sec^{-1}}, k_{+3}=0.16~\mathrm{sec^{-1}}$. The estimated time-ratios from the curves in Fig. 5 are listed in Table 2. It is necessary for the application of the steady-state approximation to choose the S(0)/E(0) ratio of more than 50. Fig. 6 shows the time-courses drawn by elongation of time axis of rapid enzymatic reaction with smaller S(0)/E(0) ratios up to the unity (relative time). For the reaction with S(0)/E(0) ratio of 30, early one-third periods may allow the application of the steady-state approximation, if one may expect a considerably large calculation error.

Rate constants at first step and steady-state

The effect of the k_{+1} value on the appearance of steady-state is shown in

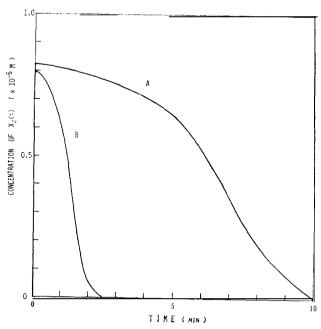


Fig. 4. Calculated time-courses of $X_2(t)$, Curve A; $k_{+3}=0.03\,\mathrm{sec^{-1}}$. Curve B; $k_{+3}=0.16\,\mathrm{sec^{-1}}$. Other parameters are the same as Fig. 3.

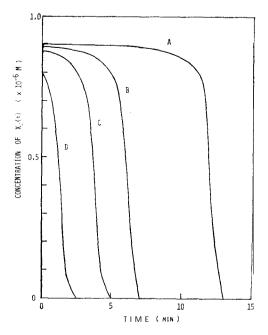


Fig. 5. Calculated time-courses of $X_2(t)$ with changing the S(0)/E(0) ratio. Curve A; S(0)/E(0)=100. Curve B; S(0)/E(0)=50. Curve C; S(0)/E(0)=30. Curve D; S(0)/E(0)=10.

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Table 2.	Relation	between	S(O)/E(O)	ratio	and	steady-state.

S(O)/E(O)	Time ratio (%)
100	73
50	57
30	0
10	0

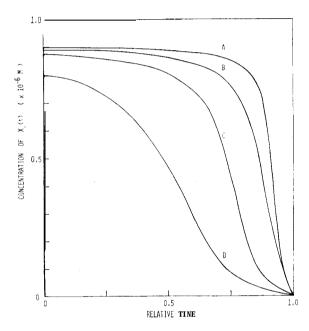


Fig. 6. Time-course of X,(t) drawn by elongation of time axis in Fig. 5. Time axis shows the relative scale (the end of reaction time is assumed to be 1.0). Curve A; S(0)/E(0)=100. Curve B; S(0)/E(0)=50. Curve C; S(0)/E(0)=30. Curve D; S(0)/E(0)=10.

Fig. 7. For the calculation of the curves in the figure, the other rate constants we re fixed at $k_1=5\,\mathrm{sec^{-1}},\,k_{+2}=10^3\,\mathrm{sec^{-1}},\,k_{-2}=10^2\,\mathrm{sec^{-1}},\,k_{+3}=0.16\,\mathrm{sec^{-1}}$ and the S(0)/E(0) ratio of 50 was used. The estimated time-ratios are listed in Table 3. Thus, it is clear that the k_{+1} value controls remarkably the periods of steady-state. However, the degree of change in the steady-state with changing the k_{+1} value depends upon the original value of k_{+1} ; as shown in Fig 7, the changes in the k_{+1} value from 5×10^6 to 5×10^5 and from 5×10^5 to 5×10^4 , cause a quite different changes in the time-course of $X_2(t)$. This will be due to the nonlinearity of the rate equation. In this connection, the contribution of the combination of k_{+1} and k_{-1} to the steady-state under fixing the values of the equilibrium constant K_1 to 10^5 was estimated. As shown in Fig. 8 and Table 4, the combination of forward and reverse rate constants in the equilibrium step seems to affect the steady-state behavior. It is clear from the comparison of the lowest

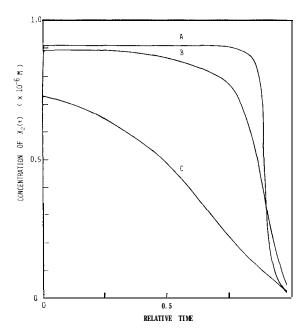


Fig. 7. Calculated time-courses of $X_2(t)$ with changing the k_{+1} value (time axis shows the relative scale). Curve A; $k_{+1} = 5 \times 10^6 \,\mathrm{M}^{-1}\,\mathrm{sec}^{-1}$. Curve B; $k_{+1} = 5 \times 10^5 \,\mathrm{M}^{-1}\,\mathrm{sec}^{-1}$. Curve C; $k_{+1} = 5 \times 10^4 \,\mathrm{M}^{-1}\,\mathrm{sec}^{-1}$.

 $k_{+1} \; (\mathrm{M^{-1}sec^{-1}})$ Time ratio (%) 5×10^6 85 5×10^5 57 5×10^4 0

Table 3. Effect of k_{+1} value on time ratio.

curve $(k_{+1}=5 \ x \ 10^4, k_{-1}=5, K_1=10^4)$ in Fig. 7 and lower curve $(k_{+1}=5 \ x \ 10^4, k_{-1}=0.5, K_1=10^5)$ in Fig. 8 that parallel changes (the latter case) in the both rate constants with constant value of K_1 cause the time-course of $X_2(t)$ much closed to the steady-state pattern. This infers that the K_1 value has also some influence to the time-course o-f the amount of the enzyme-substrate complex.

Rate constants at second step and steady-state

Since both forward and reverse rate constants at the second step are the first-order constant different from those at the first step, the effects of the rate constants k_{+2} and k_{-2} and the equilibrium constant K_2 on the steady-state behavior seem to have rather simple fashion. Consequently, only the effect of reverse rate constant was examined. As shown in Fig. 9 and Table 5, ten times increase in the k_{-2} value caused a small decrease in the steady-state periods. It should be noticed that a small value of k_{-2} caused the elongation of the steady-

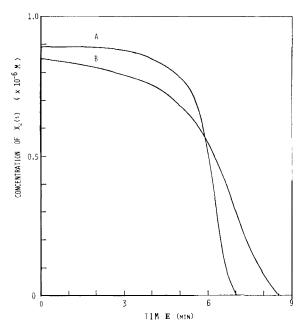


Fig. 8. Calculated time-course of $X_2(t)$ with changing the combination of k_{+1} and k_{-1} under fixing the value of the equilibrium constant. Curve A; $k_{+1} = 5 \times 10^5 \,\mathrm{M}^{-1}$ sec⁻¹, $k_{-1} = 5 \,\mathrm{sec}^{-1}$. Curve B; $k_{+1} = 5 \times 10^4 \,\mathrm{M}^{-1} \,\mathrm{sec}^{-1}$, $k_{-1} = 0.5 \,\mathrm{sec}^{-1}$. Other parameters are: S(0)/E(0) = 50, $k_2 = 10^3 \,\mathrm{sec}^{-1}$, $k_{-2} = 10^2 \,\mathrm{sec}^{-1}$, $k_{+3} = 0.16 \,\mathrm{sec}^{-1}$.

Table 4. Influence of combination of k_1 and k_{-1} on time ratio under fixing value of 4.

$k_{+1} (\mathrm{M^{-1}sec^{-1}})$	k_{-1} (sec ⁻¹)	Equilibrium const. (K_1)	Time ratio (%)
5 × 10 ⁵	5	105	57
5×10^4	0.5	105	0

state periods in contrast to the effects of other rate constants.

Factors affecting steady-state behavior

Summarized results of the simulation on the three-step enzymatic reaction are listed in Table 6. The most influential factor is the ratio of the initial concentration of substrate to that of enzyme. The ratio below 50 cannot establish the steady-state during enzymatic reactions, even if varied combination of the rate constants is adopted for simulation. The rate constants at the first step participating in the formation and the breakdown of ES, also play a role for the appearance of the steady-state. The rate constants at the second step influence only the extent of the time periods of the steady-state; if the S(0)/E(0) ratio and the rate constants at the first step have sufficient value for the appearance of the steady-state, k_{+2} and k_{-2} may show the elongation effect on the time of the steady-state. This consideration suggests that the behavior of the three-step enzymatic reaction with respect to the steady-state is essentially similar

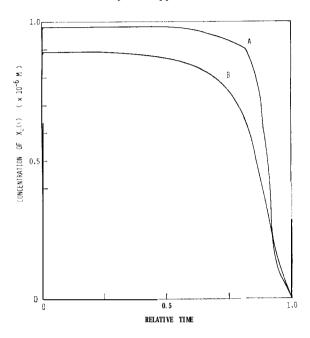


Fig. 9. Calculated time-course of $X_2(t)$ with changing the k_{-2} value (time axis shows the relative scale). Curve A ; $k_{-2}=10\,\mathrm{sec^{-1}}$. Curve B; $k_2=10^2\,\mathrm{sec^{-1}}$. Other parameters are: S(0)/E(0)=50, $k_1=5\times10^5\,\mathrm{M^{-1}\,sec^{-1}}$, $k_1=5\,\mathrm{sec^{\,1}}$, $k_{-3}=0.16\,\mathrm{sec^{-1}}$.

Table 5. Effect of k_{-2} value on time ratio.

k-, (sec-1)	Time ratio (%)
10	57
10	75

Table 6. Summarized results of simulation.

C(O) /E(O)	400	-1001	00—50			~ ~		~^	0.0	4.0	
S(O)/E(O)	100	5	5 50	50)	50	50	50	30	10	10
$k_{+1}(10^5 \mathrm{M}^{-1} \mathrm{sec}^{-1})$. =	<u>ر</u>	3 30	5		0.5		0.5	5	5	5
$k_{-1}(\sec^{-1})$	5	5	5	5	5	5	5	0.5	5	5	5
$k_{+2}(10^3 { m sec}^{-1})$	1	1	1	1	1	1	1	1	1	1	1
$k_{-2}(10^2 { m sec}^{-1})$	1	1	1	1 1	0	•	1 1	1	1	1	1
$k_{+3}(\sec^{-1})$	0.12	0.16	I1.20	0.16	0.16	0.16	0.16	0.10	0.16	0.16	0.03
Whole reaction (time)	17	13	10	6.5	7	6	12	8.5	5	2.5	10
Steady-state time (min)	12.5	9.5	7.	5 5	.5	4	4.5	() ()	0	0	()
Time ratio (%)	74	73	75	5 8	5	57	75	() 0	0	0	ii
X2 teady-state (10-6M)	0.90	0.90	0.90	0.91	0.89	0.98	0	0	0	0	0

to that of the two-step reaction (Michaelis-Menten scheme). The k_{+3} which controls the rate of overall reaction does not change the profile of the time-course of $X_2(t)$. This may infer that when the original enzymatic reaction does not exhibit the steady-state periods, the appearance of the steady-state may not be expected by the artificial slow-down of the reaction with changing the milieu conditions such as reaction temperature.

DISCUSSION

Analog computer has been widely used for simulation of chemical reaction including enzymatic reactions because computation of rate equation can easily be performed. In most cases, however, simulation results were not represented by the real time-scale; the time-scale in the plotting of time-course has usually second units with the whole time range of order of several ten seconds. In another words, the computation has been performed by choosing suitable values of rate constants so that substrate was completely consumed within several ten seconds. The reason why such the computation has widely been performed is the difficulty of the real time computation by analog comput-Since enzymatic reactions are represented in nature by nonlinear differential equation, the judgement on the steady-state from the simulated results by simply replacing the calculated arbitrary time-scale to the real time-scale should be avoided. For exact simulation and following proper judgement, therefore, it is preferable that the computation is performed with the real time-scale. rate equation of multi-step enzymatic reaction contains rate constants of various order, from 10⁻¹ to 10⁷. The differential equation containing such a great difference in the values of rate parameters is not solved by usual analog computer, when the real time-scale is required.

In digital simulation, the great difference in the values of rate constants is not critical factor for the numerical computation of the rate equation. However, in practical computation, the steep gradients of E(t), X,(t) and $X_2(t)$ at an early stage (see Fig. 2) caused an overflow at the operational register and the very small value of E(t) in following stages induced frequently the negative calculated value due to the computational error, suspending the flow of computation. To avoid such the obstructions, MRKGM program was adopted (see Fig. 1). weak point of MRKGM is that the computational error with this program could not be predicted by the theoretical background. The amplitude of computational error with MRKGM must be estimated by comparing the computational results with those obtained by other programs. In comparison with the computational result obtained by HAMPCD program (FACOM SSL program, Hamming method), it was evidenced that computed values with MRKGM contain within 1% error. Thus, the computations with digital computer enabled the real-time simulation of the three-step enzymatic reaction.

In present study, the rate constants were assumed with referring to the experimentally obtained values on hydrolytic enzyme, lysozyme and chymotrypsin. Lysozyme catalyzes the cleavage of β -1,4-glucosaminide linkages of substrate larger than (NAG),. The k_{+1} value is independent of the polymerization degree of substrate, being held at fixed value, whereas the k_{-1} value is reversely pro-

Enzyme	Substrate	$k_{+1} (\mathrm{M}^{-1} \mathrm{sec}^{-1})$	$k_{-1} (\sec^{-1})$	Reference
Malate dehydrogenase	NADH	5×1()8	50	Czeriinski and Schreck (1963)
Pyruvate carboxylase-Mn²+	Pyruvate	4.5×10 ⁶	2.1×10^{4}	Mildvan, Leigh and Cohn (1967)
Ribonuclease	Cytidine 2',3'- cyclic phosphate	2-4× 1 0 ⁷	$1-2\times10^{4}$	Erman and Hammes (1966)
Lysozyme	$(NAG)_2$ (NAG) ,	$^{4.6\times10^{6}}_{4.4\times10^{6}}$	$\left. egin{array}{l} 9.5 \hspace{0.1cm} imes 10^2 \ 2.8 \hspace{0.1cm} imes 10^2 \end{array} ight\}$	Chipman and Schimmel (1968)
	n-Methyl NAG β-Methyl NAG	$1.4\pm0.7\times10^{5}\ 1.6\pm0.8\times10^{5}$	$\left\{ \begin{array}{l} 5.5 \pm 2.1 \times 10^{3} \\ 4.5 \pm 1.7 \times 10^{3} \end{array} \right\}$	Sykes (1969a)
Chymotrypsin	3-Fluoro acetyl 2-phenyl alanine	1.0×10 ⁴	4.9×10^{2}	Sykes (1969b)

Table 7. Rate constants for enzyme-substrate complex (ES,) formation.

portional to the polymerization degree. Thus, in the present study, the k_{-1} value was assumed to be 5 with considering the experimental finding that the binding constant of substrate $(NAG)_5$ is of order of 10^5 . Several examples of rate constants participating in the formation of ES, are collected and listed in Table 7. The k_{+1} and k_{-1} values of hydrolytic enzymes are not always average. Both values are distributing to a wide range depending upon the combination of enzyme and substrate. The rate constants participating in the isomerization of the enzyme-substrate complex had been estimated experimentally for a few cases. Examples are shown in Tables 8 and 9. Since the isomerization may be accompanying the some conformational change, a large value as much as the value of k_{+1} will not be expected and the values adopted to the present simulation seem to be average of those of general enzymatic reactions.

Table 8. Rate constants for lysozyme-NAG complex isomerization.

Substrate	pН	$k_{+2} ({ m sec}^{-1})$	$k_{-2}(\sec^{-1})$	Reference
(NAG) ₂	6.0	2 ×104	1×10³	Holler,
$(NAG)_2$	7.0	1.5×10^{4}	2×10^{3}	Rupley and Hess
$(NAG)_3$	7.0	4×10^{2}	50	(1969)

Table 9. Rate constants for adenosine triphosphate-creatine phosphotransferase complex isomerization.

$\mathbf{\textit{k}}_{+2}(\sec^{-1})$	$k_{-2}(\mathrm{sec^{-1}})$	k-,/ k +,(M)	Reference
1.67×104	2.4X103	6.0×10 ⁻⁴	Hammes and Hurst (1969)

As described above, the rate constants of enzymatic reactions distribute to wide range. The application of the results of the present study to design the experimental conditions for enzymatic reactions should be bounded to those reactions which contain the rate constants close to the values assumed in the present study. For the reactions that have quite different rate constants, the time-range

where the steady-state approximation is permitted should be examined by another simulation.

REFERENCES

- Chipman, D. M. and P. R. Schimmel 1968 Dynamics of Lysozyme-Saccharide Interactions. *J. Biol. Chem.*, 243:3771-3774
- Chipman, D. M. 1971 A Kinetic Analysis of Lysozyme with Oligosaccharides from Bacterial Cell Walls. *Biochemistry*, **10**: 1714-1722
- Czerlinski, G. and G. Schreck 1964 Chemical Relaxation of the Reaction of Malate Dehydrogenase with Reduced Nicotinamide Adenine Dinucleotide Determined by Fluoroescence Detection. *Biochemistry*, 3:89-100
- Erman, J. E. and G. G. Hammes 1966 Relaxation Spectra of Ribonuclease. IV. The Interaction of Ribonuclease with Cytidine 2':3'-Cyclic Phosphate. *J. Amer. Chem. Soc.*, 88: 5607-5614
- Hammes, G. G. and J. K. Hurst 1969 Relaxation Spectra of Adenosine Triphosphate-Creatine Phosphotransferase. *Biochemistry*, 8:1083–1094
- Heineken, F. G., H. M. Tsuchiya and R. Aris 1967 a On the Mathematical Status of the Pseudo-steady State Hypothesis of Biochemical Kinetics. *Math. Biosci.*, 1: 95-113
- Heineken, F. G., H. M. Tsuchiya and R. Aris 1967b On the Accuracy of Determining Rate Constants in Enzymatic Reactions. *Math. Biosci.*, **1:** 115-141
- Holler, E., J. A. Rupley and G. P. Hess 1969 Kinetics of Lysozyme-Substrate Interactions. *Biochem. Biophys. Res. Commun.*, 37: 423-429
- Mildvan, A. S., J. S. Leigh and M. Cohn 1967 Kinetic and Magnetic Resonance Studies of Pyruvate Kinase III. The Enzyme-Metal-Phosphoryl Bridge Complex in the Fluorokinase Reaction. *Biochemistry*, 6:1805-1818
- Sykes, B. D. 1969 a A Trasient Nuclear Magnetic Resonance Study of Kinetics of Methyl N-Acetyl-D-glucosaminide Inhibition of Lysozyme. Biochemistry, 8:1110-1116
- Sykes, B. D. 1969 b An Application of Trasient Nuclear Magnetic Resonance Method to the Measurement of Biological Exchange Rates with the Chymotrypsins. *J. Amer. Chem. Soc.*, 91: 949-955
- Walter, C. F., M, F. Morales 1964 An Analogue Computer Investigation of Certain Issues in Enzyme Kinetics. *J. Biol. Chem.*, 239: 1277-1283