

Simplifying the Analysis of Enzyme Kinetics of Cytochrome c Oxidase by the Lambert-W Function

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ABSTRACT

Conventional analysis of enzyme-catalyzed reactions uses a set of initial rates of product formation or substrate decay at a variety of substrate concentrations. Alternatively to the conventional methods, attempts have been made to use an integrated Michaelis-Menten equation to assess the values of the Michaelis-Menten K_M and turnover $k_{\rm cat}$ constants directly from a single time course of an enzymatic reaction. However, because of weak convergence, previous fits of the integrated Michaelis-Menten equation to a single trace of the reaction have no proven records of success. Here we propose a reliable method with fast convergence based on an explicit solution of the Michaelis-Menten equation in terms of the Lambert-W function with transformed variables. Tests of the method with stopped-flow measurements of the catalytic reaction of cytochrome c oxidase, as well as with simulated data, demonstrate applicability of the approach to determine K_M and $k_{\rm cat}$ constants free of any systematic errors. This study indicates that the approach could be an alternative solution for the characterization of enzymatic reactions, saving time, sample and efforts. The single trace method can greatly assist the real time monitoring of enzymatic activity, in particular when a fast control is mandatory. It may be the only alternative when conventional analysis does not apply, e.g. because of limited amount of sample.

Keywords: Cytochrome c Oxidase; Enzyme Kinetics; Enzymes; Lambert-W Function; Michaelis-Menten Model

1. Introduction

The conventional Michaelis-Menten model is commonly used in biochemistry to assess the values of the catalytic constant k_{cat} and the Michaelis-Menten constant $K_{\text{M}} = (k_{\text{cat}} + k_{-1})/k_1$ of irreversible enzyme-substrate reactions [1-5]:

$$E + S \xrightarrow{k_1 \atop k_{-1}} ES \xrightarrow{k_{\text{cat}}} E + P \tag{1}$$

where k_1 and k_{-1} are the forwards and backwards rate coefficients and E is the enzyme, S the free substrate, ES the enzyme-substrate complex, and P the released product. In the case when k_1 , $k_{-1} >> k_{\text{cat}}$, which is known as the approximation of quasi-equilibrium between E and ES [1], and under the assumption of a quasi-steady state, dES/dt = 0, which holds for $S_0 >> E$, the Michaelis-Menten equation is given by [1,6]:

$$\frac{dP(t)}{dt} = k_{cat} E_0 \frac{S_0 - P(t)}{K_M + S_0 - P(t)}$$
 (2)

The initial rate $v_0 = dP(t)/dt$ at $t \rightarrow 0$ is a nonlinear function of starting substrate concentrations S_0 , where E_0 is the total concentration of enzyme, and P(t = 0) = 0 is the initial condition for the product concentration. If v_0 is plotted as a function of S_0 , the parameters k_{cat} and K_M

have to be assessed with Equation (2) at $t\rightarrow 0$ by nonlinear regression. Another way is to linearize Equation (2) using inverted variables like the $1/v_0$ and $1/S_0$ and then apply linear regression [7]. The disadvantage of the inverted plots is that they are sensitive to errors for small or large values of v_0 or S_0 [5]. Eadie [8] and Hofstee [5] have proposed a linearization of Equation (2) using non inverted variables v_0 and v_0/S_0 to overcome this limitation. Independent of the particular linearization procedure employed to analyze enzyme kinetics all the methods rely on the determination of the initial rates v_0 in a wide range of starting concentrations S_0 . We will refer to these methods as v_0 -plots. In cases, when measurements in a wide range of S_0 are not possible or not feasible because of restricted availability of the samples, it would be advantageous to establish a new method for assessment of the enzymatic kinetic parameters for a single value of S_0 .

Several attempts have been made to assess the values of K_M and k_{cat} directly from a time course S(t) of an enzymatic reaction at a single S_0 . Different forms of parametrically integrated Michaelis-Menten equation, e.g. $(S_0 - S) = E_0 k_{\text{cat}} t - K_M \ln(S_0/S)$, have been fitted to progress reaction curves for enzyme catalyzed reactions to deter-

mine k_{cat} and K_M [9-12]. However, the fit of the implicit parametric solution to the explicit S(t)-curves by linear or nonlinear regression was a demanding numerical problem and often resulted in uncertain values of k_{cat} and K_M . Only beginning with the work of Schnell and Medoza [13], the explicit solution of the Michaelis-Menten equation for P(t) in terms of Lambert-W function became available, and could provide more accurate results, since no more approximations, than already made in the Michaelis-Menten model, were required. For instance, Goudar and colleagues [14-16] have used the explicit solution of the Michaelis-Menten equation, Equation (3), to analyze single progress curves of enzymatic reactions. The new method promised to be fast and sample saving. Yet surprisingly, the explicit solution of the Michaelis-Menten equation has up to now only occasionally been adopted to estimate k_{cat} and K_M , even though its theory is well documented [13].

Walsh et al. [17] have pointed out that attempts to use the explicit solution of the Michaelis-Menten Equation (2) to describe reaction time courses have so far met with little success. It is well known, that nonlinear regression, in our case with the Lambert-W function, cannot be started without specifying the initial input parameters $K_{M \text{ input}}$ and $k_{\text{cat,input}}$, even if the whole progress curve is accurately measured. If $K_{M,input}$ and $k_{cat,input}$ are significantly different to the real K_M and k_{cat} , the nonlinear fit converges very slowly, if at all, and the estimated values of K_M and k_{cat} are imprecise. To our knowledge, the problem of the strong dependency of the fitted constants $K_{\rm M}$ and k_{cat} on the input parameters $K_{M,\text{input}}$ and $k_{\text{cat,input}}$ has not be properly addressed so far. In the present paper we examine the problem and demonstrate, that because of a weak convergence, previous fits of the explicit solution of the Michaelis-Menten equation to a single trace P(t)could not provide a reliable estimation of K_M and k_{cat} . We propose an alternative fitting method using an explicit solution of the Michaelis-Menten equation in terms of the transformed Lambert-W function, Equation (8), which is much less prone to uncertainties in input parameters than previous single-trace methods. We compare the method in simulated and experimental conditions with conventional v_0 -plots to demonstrate that the new approach could be developed to a standard method for a time and sample saving characterization of enzymatic reactions. For that we use simulated data following the ideal irreversible Michaelis-Menten kinetics as well as experimental data on the reaction of cytochrome c oxidase (CcO). At high salt conditions and in the presence of an excess of oxygen, the CcO-reaction meets the single substrate Michaelis-Menten model and no product inhibition takes place. Additionally, the CcO-reaction guarantees full irreversibility of the enzymatic reaction due to formation of water from oxygen during the catalysis [18].

2. Material and Methods

2.1. Preparation of Enzyme Cytochrome *c*Oxidase

Cytochrome c oxidase (CcO) is the terminal complex of the membrane-bound respiratory chain and spends electrons from ferrous cytochrome c for the reduction of oxygen to water. CcO from Rhodobacter sphaeroides was expressed and purified, using 21 cell culture flasks in a gyratory shaker followed by 12 h solubilisation in detergent [19]. Purified protein was stored in phosphate buffer (50 mM, pH 8 and 0.01% dodecylmaltoside) at -80°C until use. The enzyme concentration was determined from UV/VIS spectra of air-oxidised and sodium dithionite reduced samples, using the difference of the differential extinction coefficients $\Delta \varepsilon_{605} - \Delta \varepsilon_{630} = 24 \text{ mM}^{-1} \cdot \text{cm}^{-1}$ at the wavelengths $\lambda = 605$ nm and 630 nm [19]. Ferrous cytochrome c from horse heart (purity > 95% from Fluka) was used without further purification. A solution of cvtochrome c was reduced in a fivefold molar excess of sodium dithionite. Reductant was separated via a 5 ml HiTrap desalting column (GE Healthcare) on an ÄktaPurifier FPLC (GE Healthcare) and the protein was stored at -80° C. The concentration of cytochrome c was determined spectrophotometrically by recording the reducedminus-oxidized difference spectrum of the sodium dithionite reduced and ferricyanide oxidized sample and using the differential extinction coefficient $\Delta \varepsilon_{550} = 19$ $mM^{-1} \cdot cm^{-1}$ at $\lambda = 550$ nm [20].

2.2. Measurements of Enzyme Kinetics

Kinetic experiments of electron transfer from cytochrome c to CcO were performed by the stopped-flow technique [19]. The oxidation of ferrous cytochrome c was monitored at $\lambda = 550$ nm by a miniature fibre-optic spectrometer (USB 2000 from Ocean Optics). A 50 nM solution of oxidase was mixed with a 5 to 120 μ M solution of cytochrome c, both buffered in a 50 mM phosphate buffer, pH = 6.5, containing 0.05% dodecylmaltoside and 100 mM KCl.

2.3. Simulation of Enzyme Kinetics

The normalized time trace $p(t) = P(t)/S_0$ of an enzymatic reaction was simulated with the integrated Michaelis-Menten equation, Equation (3), for 425 equidistant time points in the range $0 \le t \le 65$ s for the parameters $K_M = 220 \, \mu\text{M}$, $k_{\text{cat}} = 650 \, \text{s}^{-1}$ and $E_0 = 50 \, \text{nM}$, *i.e.* for the parameters close to those of the CcO reaction. Pseudorandom noise was generated by the "rnd"-function of the software Mathcad 2001 Professional, MathSoft, Inc., and added to the analytical trace p(t). The amplitude of the noise was ranging from 0 up to 4% of the maximum value p = 1.

2.4. Description of the Linear and Nonlinear Regressions

For the linear regression we used the "neigung"-function of the Mathcad 2001 Professional, MathSoft, Inc. The initial rates $v_{0,\rm exp}$ were determined by linear regression using the first ten data points of the experimental and the computer simulated trace p(t). The same linear regression was applied to Lineweaver-Burk and EadieHofstee plots for estimation of the values of K_M and $k_{\rm cat}$. The conventional nonlinear regression with Equation (2), as well as the nonlinear regression with the integrated Michaelis-Menten equation was based on the iterative Levenberg-Marquardt method implemented in the "minfehl"-function of the Mathcad 2001 Professional.

3. Results and Discussion

3.1. Simulation of Enzymatic Reactions

Our simulations of enzymatic reactions employ an integration of the Michaelis-Menten Equation (2) yielding a closed analytical formula for P(t) in terms of the Lambert-W function W(x) [13], in contrast to frequently applied implicit parametric solutions:

$$P(t) = S_0 - K_M W \left(\frac{S_0}{K_M} \exp \left(\frac{S_0}{K_M} - \frac{k_{cat}}{K_M} E_0 t \right) \right)$$
(3)

Note that Equation (3) refers to the fully irreversible enzymatic reaction as constituted by Equation (1). The progress curve P(t) is simulated by Equation (3) at $S_0 = 30 \, \mu\text{M}$, $K_M = 220 \, \mu\text{M}$, $k_{\text{cat}} = 650 \, \text{s}^{-1}$ and $E_0 = 50 \, \text{nM}$. The result is an ideal product P(t)-curve following the one-substrate Michaelis-Menten model. Then Equation (3) is fitted with K_M and k_{cat} to P(t)-curve by nonlinear regression as described in Section 2.4. To study the robustness of the method against deviations of initial input parameters from the true values, we started the nonlinear regression with $K_{M,\text{input}} \neq K_M = 220 \, \mu\text{M}$ and $k_{\text{cat,input}} \neq k_{\text{cat}} = 650 \, \text{s}^{-1}$. The extents of the relative variations of the input parameters $\Delta K_{M,\text{input}}/K_M$ and $\Delta k_{\text{cat,input}}/k_{\text{cat}}$ are quantified by

$$\Delta K_{M,\text{input}} / K_M = \left(K_{M,\text{input}} - K_M \right) / K_M \tag{4}$$

$$\Delta k_{\text{cat input}} / k_{\text{cat}} = \left(k_{\text{cat input}} - k_{\text{cat}} \right) / k_{\text{cat}}$$
 (5)

The relative deviations of the estimated parameters $\delta K_{M,\text{est}}/K_M$ and $\delta k_{\text{cat,est}}/k_{\text{cat}}$ are characterized by

$$\delta K_{M,\text{est}} / K_M = \left(K_{M,\text{est}} - K_M \right) / K_M \tag{6}$$

$$\delta k_{\text{cat,est}}/k_M = (k_{\text{cat,est}} - k_{\text{cat}})/k_{\text{cat}}$$
 (7)

The calculation showed, that the relative deviations of the input parameters in the range of $-0.5 \le \Delta K_{M,\text{input}}/K_M \le 0.5$ and $-0.5 \le \Delta k_{\text{cat,input}}/k_{\text{cat}} \le 0.5$ cause practically equal relative deviations of both estimated parameters:

 $\delta K_{M,\rm est}/K_M \approx \delta k_{\rm cat,est}/k_{\rm cat}$. Therefore, we combined the results for relative deviations of the estimated parameters $K_{M,\rm est}$ and $k_{\rm cat,est}$ in a single plot, **Figure 1**.

The results of a direct fit of Equation (3) to the time course P(t) are summarized in **Figure 1(a)**. If the value of $k_{\text{cat,input}}$ is somewhat different to the exact value, both $K_{M,\text{est}}$ and $k_{\text{cat,est}}$ cannot be determined correctly. On the other hand, $K_{M,\text{est}}$ and $k_{\text{cat,est}}$ are very little sensitive to variations of $K_{M,\text{input}}$. It means that the fit of Equation (3) to P(t) cannot provide convergence to correct values of K_{M} and k_{cat} , if the initial input parameter $k_{\text{cat,input}}$ somenwhat deviates from the true value of 650 s⁻¹. As far as we know, all previous attempts to determine $K_{M,\text{est}}$ and $k_{\text{cat,est}}$ explicitly using Equation (3) were based on the similar approach as tested above [14-16].

We solve this problem by employing the fact that the apparent rate constant in Equation (3) is given by $m = k_{\text{cat}} \cdot E_0 / K_M$ [2]. In terms of m, Equation (3) takes the form:

$$p(t) = \frac{P(t)}{S_0} = 1 - \frac{1}{s}W(s\exp(s - mt))$$
 (8)

where $s = S_0 / K_M$ is the dimensionless parameter. Applying Equation (8) to fit kinetics of an enzymatic reaction by nonlinear regression, we render the procedure robust against errors in initial input values of K_M and $k_{\rm cat}$. Using the transformed Equation (8), we obtain very small relative errors for the estimated parameters $\delta K_{M,\rm est} / K_M \approx \delta k_{\rm cat,est} / k_{\rm cat}$ in the range between -7.5×10^{-3} and 1.5×10^{-2} , **Figure 1(b)**. For the most values of $K_{M,\rm input}$ and $k_{\rm cat,input}$ studied here we calculate even zero deviations. For characterization of the quality of the fit, we use the standard deviation:

$$SD = \sqrt{\frac{1}{N-1} \sum_{i=1}^{N} \left(p\left(t_{i}, K_{M,\text{est}}, k_{\text{cat,est}}\right) - p\left(t_{i}, K_{M}, k_{\text{cat}}\right) \right)^{2}}$$
(9)

where N = 425 is the number of the equidistant time points t_i at which the function $p(t_i)$ is sampled. Note that when the non-transformed Equation (3) is used, the location of local minima of SD is on a line diagonal to $\Delta K_{M,\text{input}}/K_M$ and $\Delta k_{\text{cat,input}}/k_{\text{cat}}$ axis, see **Figure 2(a)**. This is very unfavorable for nonlinear regression and leads to the weak convergence and mainly to a wrong local minimum. The large deviation from the global minimum causes large uncertainty in $K_{M,est}$ and $k_{cat,est}$. The transformed Equation (8) applied in the new (m, s)-coordinate system exhibits much faster convergence, as Equation (3) in the (K_M, k_{cat}) -coordinates, **Figure 2(b)**. Contrary to Equation (3), the fit of the apparent rate constant m at a constant parameter s yields the correct m-coordinate of the global minimum of SD. Fitting s at a correct m-coordinate of the minimum greatly facilitates convergence to the well-defined global minimum of SD shown by the arrow in **Figure 2(b)**. The estimated coordinates m_{est} , s_{est} of the global minimum can be easily recalculated into the parameters $K_{M,\text{est}}$ and $k_{\text{cat,est}}$.

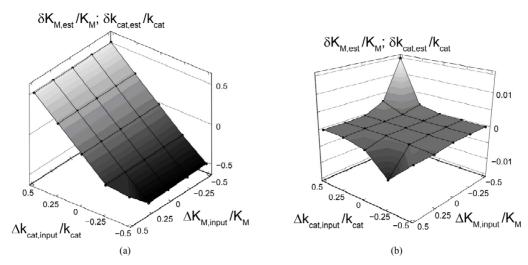


Figure 1. Hyperplane of the relative deviations of the estimated parameters $K_{M,\text{est}}$ and $k_{\text{cat,est}}$ from the actual values as a function of the relative deviations of the input parameters $\Delta K_{M,\text{input}}/K_M$ and $\Delta k_{\text{cat,input}}/k_{\text{cat}}$. Black spheres represent calculated points connected by eye-help lines and surfaces. (a) $K_{M,\text{est}}$ and $k_{\text{cat,est}}$ are estimated using Equation (3). (b) $K_{M,\text{est}}$ and $k_{\text{cat,est}}$ are estimated using the transformed Equation (8) in the form $P(t) = p(t) \cdot S_0$. Note the large difference in the scales of the vertical axes of (a) and (b). For both plots $K_{M,\text{est}}$ and $k_{\text{cat,est}}$ are determined by nonlinear regression applied to the error-free function P(t) computed by Equation (3).

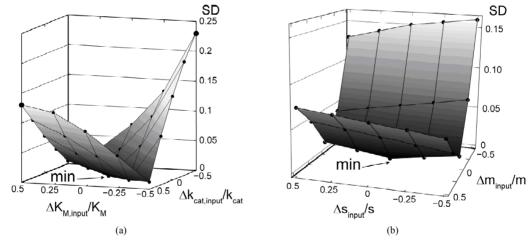


Figure 2. Standard deviation SD, computed with Equation (9), as a function of the relative deviation of input parameters: $\Delta K_{M,\text{input}}/K_M$ and $\Delta k_{\text{cat,input}}/k_{\text{cat}}$, (a), and $\Delta s_{\text{input}}/s = (s_{\text{input}} - s)/s$, $\Delta m_{\text{input}}/m = (m_{\text{input}} - m)/m$, (b). Black spheres represent calculated values of SD connected by eye-help lines and surfaces. The arrows show the position of the global minima of SD: (a) regression with Equation (3), (b) regression with Equation (8). The error-free functions P(t) and $p(t) = P(t)/S_0$ are computed as specified in Figure 1.

3.2. Advanced Algorithm for Single Trace Regression

The impact of the precision of the input parameters $K_{M,\text{input}}$ and $k_{\text{cat},\text{input}}$ on the errors on the estimated values of $K_{M,\text{est}}$ and $k_{\text{cat},\text{est}}$ becomes stronger with increasing noise level of data. Previously, $K_{M,\text{input}}$ and $k_{\text{cat},\text{input}}$ were determined through a linearization of Equation (3) resulting in uncertain estimates [15,16]. Here we avoid the linearization by using the fact that the simple exponential function

$$p_{\text{appr}}(t) = 1 - \exp(-m_{\text{input}}t)$$
 (10)

is an approximation to Equation (8), [2]. First, we estimate the input value of the apparent rate constant $m_{\rm input}$ via fitting Equation (10) by regression to experimental or simulated p(t) curves. Calculations with simulated p(t) have shown (data are not presented) that for a reliable estimation of $m_{\rm input}$ it is recommendable to sample p(t) up to the maximum time $t_{\rm max} \approx 7/m_{\rm input}$. The second input parameter for the dimensionless parameter $s_{\rm input}$ is obtained using the first derivative of Equation (8) for $t \to 0$:

$$\frac{\mathrm{d}p(t)}{\mathrm{d}t}\bigg|_{t\to 0} = \frac{m}{1+s_{\mathrm{input}}} \tag{11}$$

The value $(dp(t)/dt)|t\rightarrow 0 \approx v_{0,exp}$ is estimated by usual linear regression applied to ten or twenty initial points of the p(t) curve. Rearrangement of Equation (11) gives the second input parameter s_{input} in the form:

$$s_{\text{input}} \approx m_{\text{input}} / v_{0,\text{exp}} - 1$$
 (12)

It is recalled that linear regression for $v_{0,\rm exp}$ and nonlinear regression of the exponential Equation (10) for $m_{\rm input}$ yield intrinsically wrong estimates of $m_{\rm input}$ and $s_{\rm input}$. However, the estimates are good enough to serve as input parameters for the nonlinear regression with the transformed Equation (8); cf. next Section 3.3. On this way, the determination of $m_{\rm input}$ and $s_{\rm input}$ can be easily automated; see the reference in the Appendix 1.6. Finally, the searched kinetic constants are given by $K_{M,\rm est} = S_0/s_{\rm est}$ and $k_{\rm cat,est} = m_{\rm est} \cdot K_{M,\rm est}/E_0$.

3.3. Robustness of Single Trace Regression to Noise

For testing of robustness of single trace regression to error-prone data, noise was added to the error-free function p(t), Equation (8), in form of the pseudo random error:

$$p_{\rm err}(t) = p(t) + e \cdot x \tag{13}$$

where e is the noise amplitude ranging from zero to 0.04, and x is the pseudo-random variable varying in the range of -1 to 1. The function $p_{\rm err}(t)$ is calculated in N=425 equidistant points in the time range of 0 to 65 s. The estimates $m_{\rm est}$ and $s_{\rm est}$ are determined as averages of five samples of noise of $p_{\rm err}(t)$. As a measure for the quality of the estimations we use the standard deviation, Equation (9). The resulting SD values are small for all noise levels: SD=0 at e=0, $SD=5.8\times 10^{-3}$ at e=0.01 and SD=0.024 at e=0.04. Symmetric distribution of the relative deviations of the estimated parameters $\delta K_{M, \rm est}/K_M$ and $\delta k_{\rm cat, est}/k_{\rm cat}$ around zero suggests no systematic error in the resulting $K_{M, \rm est}$ and $k_{\rm cat, est}$ values; see **Figure 3**, right panels.

As mentioned above, the values of $\delta K_{M,\rm est}/K_M$ and $\delta k_{\rm cat,est}/k_{\rm cat}$ are practically equal to each other in the whole range of $\Delta K_{M,\rm input}/K_M$ and $\Delta k_{\rm cat,input}/k_{\rm cat}$ studied here; therefore, they are presented together in the same graph, **Figure 3** (right panels). The estimates of $K_{M,\rm est}$ and $k_{\rm cat,est}$, averaged over the results for 25 input pairs of $K_{M,\rm input}$ and $k_{\rm cat,input}$, are summarized in **Table 1**. Note that the estimates are very close to the actual values $K_M = 220~\mu{\rm M}$ and $k_{\rm cat} = 650~{\rm s}^{-1}$, even at e = 0.04.

3.4. Conditions for the Best Convergence

Besides of the noise level of P(t), the quality of analysis depends on the absolute value of the initial substrate concentration S_0 . To demonstrate the correlation between

noise-level and initial substrate concentration, the relative deviation of estimated K_M and k_{cat} values as a function of S_0 is depicted in **Figure 4(a)** for three different levels of noise.

The absolute values of the relative deviations of the estimated parameters K_M and $k_{\rm cat}$ became larger with decreasing S_0 and increasing noise amplitude e. The dependence of $|\delta K_{M,\rm est}|/K_M$ and $|\delta k_{\rm cat,est}|/k_{\rm cat}$ on S_0 can be referred to large rounding errors of the nonlinear regression at small values of initial substrate concentrations, especially when $S_0 << K_M$ (here at about $S_0 = 10~\mu{\rm M}$), and at larger level of noise. Since the parameter s in Equation (8) interrelates S_0 to K_M , we estimated for a given parameter s the ranges of the apparent rate constant s0, which condition a correct determination of s1 and s2 kcat. Reliable results of the fit we obtain for s3 = 0.01 and s3 between 0.02 and 1.2 s⁻¹, for s3 = 0.1 and s4 between 0.024 and 1.9 s⁻¹ as well as for s3 = 1.0 and s4 between 0.07 and 1.2 s⁻¹. In that range of the parameters the

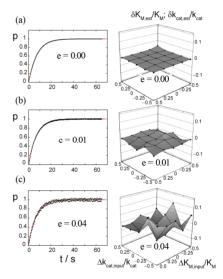


Figure 3. Left panels: Progress curves simulated with Equation (8) with parameters as indicated in Figure 1, now with random noise of the amplitudes (a) e=0, (b) e=0.01 and (c) e=0.04. The red lines represent fits with Equation (8) for the calculation of the parameters $K_{M,\rm est}$ and $k_{\rm cat,est}$. Right panels: Black spheres connected by eye-help lines and surfaces represent the relative deviation $\delta K_{M,\rm est}/K_M$ and $\delta k_{\rm cat,est}/k_{\rm cat}$ of the estimated parameters from the actual values as function of the relative deviations of the input parameters $\Delta K_{M,\rm input}/K_M$ and $\Delta k_{\rm cat,input}/k_{\rm cat}$.

Table 1. Estimates of K_M and $k_{\rm cat}$ averaged over 25 input pairs $K_{M,\rm input}$ and $k_{\rm cat,input}$ at three different noise levels e. K_M and $k_{\rm cat}$ are computed by single trace regression with Equation (8).

e	$K_M/\mu M$	$k_{\rm cat}/{ m s}^{-1}$
0	220	650
0.01	220 ± 3	649 ± 9
0.04	220 ± 11	651 ± 31

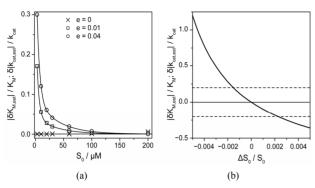


Figure 4. (a) Absolute values of relative deviations of the estimated parameters $K_{M,\rm est}$ and $k_{\rm cat,est}$ from the actual values as functions of the initial substrate concentration S_0 at the three different levels of noise: \times , e=0.0, \square , e=0.01, and \Diamond , e=0.04. The nonlinear regression method was applied to $p_{\rm crr}(t)$, see Equation (13), at the input parameters $K_{M,\rm input}=275~\mu M$, $k_{\rm cat,input}=487.5~{\rm s}^{-1}$, corresponding to $\Delta K_{M,\rm input}/K_M=0.25$ and $\Delta k_{\rm cat,input}/k_{\rm cat}=-0.25$. (b) The impact of the relative error $\Delta S_0/S_0$ in initial substrate concentration S_0 on the relative deviations $|\delta K_{M,\rm est}|/K_M$ and $\delta |k_{\rm cat,est}|/k_{\rm cat}$ of the estimated parameters from the actual values. The solid line represents results of nonlinear regression for the errorfree function p(t); other parameters as in (a). Broken horizontal lines indicate a 20% relative error in K_M and $k_{\rm cat}$.

computation algorithm for Lambert-W function and nonlinear regression implemented in Mathcad 2001 Professional yield exact values of $K_{M,est}$ and $k_{cat,est}$ (data are not shown). The precision of the value of S_0 appears to be very important for a correct analysis of progress curves, because of the direct dependence of s on S_0 . Already the relative deviation $\Delta S_0/S_0 = (S_{0,\text{meas}} - S_0)/S_0$ of the measured value $S_{0,\text{meas}}$ from the exact value S_0 in the range \pm 0.5% causes a hundredfold larger relative error in the estimated values of $K_{M,est}$ and $k_{cat,est}$, Figure 4(b). In order to keep the relative error in $K_{M,est}$ and $k_{cat,est}$ within \pm 20% error margin, the relative deviation $\Delta S_0/S_0$ should be in the range between -0.18 and +0.21%. Therefore, the single trace regression requires very precise values of the starting substrate concentrations. Generally, the demanded precision of S_0 should be better than about $\pm 0.1\%$, which is nowadays available in most laboratories.

3.5. Testing *v*₀-Plots and Single Trace Regression with Simulated Data

We tested the limits and quality of the new single trace regression as well as commonly used v_0 -plots using computer generated data with pseudo-random noise. One source of errors in v_0 -plots is an inaccurate experimental determination of v_0 . In our simulations, we determine the realistic error in v_0 using a computer simulated trace p(t) with a known level of pseudo-random noise, as opposed to previous computer tests, in which a fixed error was added to v_0 [5]. Three classical v_0 -plots (Lineweaver-

Burk, Eadie-Hofstee and nonlinear) as well as the single trace regression method were applied to the same data set. The results are presented in **Table 2** (see also **Tables 4** and **5** in Appendix).

Table 2 clearly shows that the single trace regression with Equation (8) yields the most precise results for K_M and k_{cat} when error prone simulated data are analyzed.

The nonlinear v_0 -plot yields good estimates coinciding with actual values within their standard deviations. The Eadie-Hofstee plot yields less accurate results, the estimate does not compare to the true value k_{cat} , even if within the error margin. The Lineweaver-Burk plot yields unrealistic negative values of K_M and k_{cat} . Presumably, an uneven distribution of error over S_0 -values leads to the unrealistic results. This could be to a certain degree corrected by disregarding data points for smaller S_0 -values, a strategy which is frequently employed by experimentalists exploiting the Lineweaver-Burk plot. Our analysis adheres strictly to all values of the plots to ensure the full comparability of all the methods. We also did not perform a weighting of the data points corresponding to their precision, which additionally may have led to improved results. Another reason for the failure of the Lineweaver-Burk plot is the narrow width of the analyzed S_0 range between 3 and 100 μ M, compared to the K_M -value of 220 μ M. Generally, a wider range of S_0 values used for analysis increases precision of results of all v_0 -plots considerably [5]. Regarding this fact, our new single trace regression method could be of great benefit, especially in cases, where a wide range of S_0 values cannot be addressed experimentally. As expected, results for a wider range of staring concentrations S_0 are more precise, than those for the narrow range of S_0 -values (see Figure 7 and Table 4 of Appendix). The reason for the notably more precise results of the single trace regression compared to the three v_0 -plots in **Table 2** could be the fact that the initial rates v_0 are usually underestimated in their experimental determination (cf. Appendix Figure 6). The effect of the underestimation of v_0 on the results of the v_0 -plots is actually hardly predictable. Dependent on the actual noise level, the estimated values of K_M and k_{cat} can be systematically shifted either to lower or to higher values. In that case, the small error margins of K_M and $k_{\rm cat}$ are misleading, as demonstrated for the $k_{\rm cat}$ value determined by the Eadie-Hofstee plot in Table 2. Differently to the v_0 -plots, the results of the new single trace method do not show any systematic deviation from the actual K_M and k_{cat} values at all three levels of noise studied here, see **Figure 3**. In principle, the assessment of K_M and k_{cat} can be improved by repeating the single trace experiments and averaging, as opposed to the v_0 -plots using systematically underestimated values of $v_{0,exp}$. Therefore, we may conclude that the new method has a potential to yield more precise results than the conventional v_0 -plots.

single trace regression. nonlinear v_0 -plot. Eadie-Hofstee plot. Lineweaver-Burk plot. Equation (8) Equation (17) Equation (15) Equation (14) $k_{\rm cat}/{\rm s}^ K_M/\mu M$ $k_{\rm cat}/{\rm s}^{-1}$ $K_M/\mu M$ $K_M/\mu M$ $k_{\rm cat}/{\rm s}^{-1}$ $k_{\rm cat}/{\rm s}^{-1}$ $K_M/\mu M$ -160 ± 270 217 ± 7^{b} 240 ± 22 650 ± 43 643 ± 18 210 ± 30 565 ± 60 -310 ± 650

Table 2. Comparison of different methods to determine kinetic constants from simulated enzymatic reactions with noise.

^aReactions are simulated at seven S_0 varying between 3 and 100 μM at $K_M = 220$ μM, $k_{\text{cat}} = 650$ s⁻¹, $E_0 = 50$ nM, including pseudo-random noise of the amplitude 0.5 μM; v_0 -values are determined by linear regression. ^bStandard deviation derived from five independent simulations of p(t) for the single trace regression with Equation (8) at $S_0 = 50$ mM and fivefold repetition for other methods.

3.6. Testing Single Trace Regression with Experimental Data

To gauge the conventional methods to the single trace regression analysis under experimental conditions of enzymatic catalysis we have chosen the reaction of CcO with ferrous cytochrome c, because at high salt conditions and surplus of oxygen it meets the single substrate Michaelis-Menten model and guarantees full irreversibility of the enzymatic reaction due to formation of water from oxygen during the catalysis as well as the exclusion of product inhibition [18]. The high ionic strengths conditions lead further to slower kinetics of this enzymatic reaction compared to the optimum rate at low ionic strength; see, e.g., review by Cooper [18].

Initial reaction rates $v_{0,\text{exp}}$ are measured spectrophotometrically at different starting concentrations S_0 of the substrate, ferrous cytochrome c. The corresponding Lineweaver-Burk, Eadie-Hofstee and nonlinear v_0 -plots are presented in **Figures 5(a)-(c)**.

The resulting kinetic parameters K_M and k_{cat} of the plots are summarized in **Table 3**.

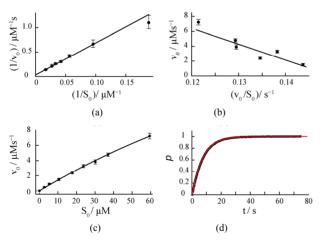


Figure 5. Four different plots of the kinetic analysis of the reaction of cytochrome c oxidase (E_0) with ferrous cytochrome c (S_0) . Experiments are performed at constant $E_0=50$ nM in a 50 mM phosphate buffer, pH = 6.5 and 100 mM KCl. Lineweaver-Burk (a), Eadie-Hofstee (b) plots and nonlinear v_0 -regression (c) for various cytochrome c initial concentrations S_0 . Normalized kinetics of the product formation at $S_0=30.08$ μ M (d). Black circles represent experimental data, the red curve represents Equation (8) with $K_M=218$ μ M and $k_{\rm cat}=645$ s⁻¹.

A quantitative comparison of the simulated product curves p(t) at different levels of noise with the experimental p(t) by Equation (9) shows that the experimental level of noise (compared to Table 1) allows for a definite determination of K_M and k_{cat} by all three v_0 -plots studied here. A representative result of a fit of Equation (8) to a single trace of an enzymatic reaction is shown in Figure **5(d)**. The sevenfold repetition of the single trace regression at seven different S₀ values (see Appendix, **Table 5**) enables computation of standard deviations of K_M and k_{cat} in **Table 3**. It is apparent from the **Table 3** that the Lineweaver-Burk plot yields the most imprecise results, what is in line with the conclusion of Hofstee [5]. The nonlinear v_0 -plot provides the smallest standard deviations of $k_{\rm cat}$ compared with single trace regression or Lineweaver-Burk and Eadie-Hofstee plots. Within the relative large error margin of K_M and k_{cat} , the results of the single trace regression are at least consistent with the results of the v_0 -plots. The variations of the kinetic parameters K_M and k_{cat} determined by single trace regression at different substrate concentrations S_0 reflect the limits of the method caused by experimental errors; see **Table 5** in Appendix. However, the experimental result itself does not suggest any systematic deviations of K_M and k_{cat} with S_0 .

In our experiment it is not clearly evident whether the single trace regression or the v_0 -plots are generally more precise in determination of k_{cat} and K_M . At least we demonstrate here that single trace regression leads to realistic estimates, which are in good agreement with the values of the v_0 -plots.

4. Conclusions

The analytical integration of the Michaelis-Menten Equation (2) in terms of the transformed Lambert-W function, Equation (8), provides a reliable tool to determine the Michaelis-Menten K_M and turnover k_{cat} constants from the analysis of a single reaction trace. Supported by our advanced nonlinear regression algorithm, this new method allows determining K_M and k_{cat} much quicker than by conventional linearization or by nonlinear plots using a set of $(v_{0,\text{exp}}, S_0)$ pairs.

The method can be easily implemented as a time and sample saving tool in the online characterization of enzymatic activity. The kinetic parameters $K_M = 219 \pm 4$

	ce regression, ation (8)		on (17)		ofstee plot, on (15)		r-Burk plot, on (14)
$K_{ m M}/\mu{ m M}$	$k_{\rm cat}/{ m s}^{-1}$	$K_M/\mu M$	$k_{\rm cat}/{\rm s}^{-1}$	$K_M/\mu M$	$k_{\rm cat}/{ m s}^{-1}$	$K_M/\mu M$	$k_{\rm cat}/{\rm s}^{-1}$
219 ± 4^a	790 ± 120	258 ± 22	770 ± 43	213 ± 30	640 ± 60	220 ± 270	670 ± 650

Table 3. Kinetic parameters for the reaction of CcO with ferrous cytochrome c.

μM and $k_{cat} = 790 \pm 120 \text{ s}^{-1}$ determined with Equation (8) for the CcO reaction are quite conform to results of conventional v_0 -plots. The fact that no systematic errors occur in K_M and k_{cat} in the large range of S_0 tested here suggests that application of the single trace method at $S_0 < K_M$ can complement, if not replace, the conventional methods requiring measurements in the larger range of S_0 (0 < $S_0 < 2 K_M$, Hofstee (1959), [5]). Note that for some enzymatic reactions, like oxidation of cytochrome c by CcO studied here, the region $S_0 > 0.5 K_M$ is experimentally not accessible because the absorption of cytochrome c becomes too strong for correct spectrophotometric determination.

We can therefore expect that the new single trace non-linear regression will apply to all single substrate irreversible enzymatic reactions whenever a reliable, precise and fast assessment of the kinetic parameters K_M and k_{cat} is demanded. The single trace regression with the transformed Lambert-W function can be adopted to analyze reversible enzymatic reactions [21,22], as well as to enzyme kinetics of multiple alternative substrates [23]. In any case, the method will provide a new tool for biotechnology that saves sample and time by reducing the number of experiments at various S_0 mandatory for the determination of K_M and k_{cat} by v_0 -plots.

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^aStandard deviations are determined as described in the text.

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Appendix

1. Comparison of the Approaches to Assess K_M and k_{cat}

1.1. Linearizations of the Michaelis-Menten Rate Equation

Enzyme kinetics is most commonly analyzed by measuring the initial reaction rate v_0 at different substrate concentrations S_0 . A number of linearization procedures are available for the evaluation of such datasets [3]. Hofstee [5] has pointed out that there is no real basis for the continued use of inverted linearization, like the $1/v_0$ versus $1/S_0$, the so-called Lineweaver-Burk plot [7]:

$$\frac{1}{v_0} = \frac{K_M}{k_{\text{cat}} E_0} \frac{1}{S_0} + \frac{K_M}{k_{\text{cat}} E_0}$$
 (14)

because the plot is sensitive to errors for low and high values of v_0 or S_0 . Eadie [8] and Hofstee [5], proposed to use the v_0 versus v_0/S_0 plot, now commonly known as Eadie-Hofstee plot:

$$v_0 = -K_M \frac{v_0}{S_0} + k_{\text{cat}} E_0 \tag{15}$$

The maximum rate of the product formation is conventionally defined by

$$V_{\text{max}} = k_{\text{cat}} E_0 \tag{16}$$

1.2. Non-Linearized Michaelis-Menten Equation

Beginning with the wide availability of powerful computers and nonlinear regression software, the Eadie-Hofstee and other linear plots have been partly superseded by nonlinear regression method that should be significantly more accurate and no longer computationally inaccessible. The nonlinear regression method uses the Michaelis-Menten Equation (2) at $t \rightarrow 0$ in the nonlinear form

$$v_0 = k_{\text{cat}} E_0 \frac{S_0}{K_M + S_0} \tag{17}$$

1.3. Error in the Initial Reaction Rate v_0

Practical determination of v_0 always leads to underestimated values of the initial reaction rate $v_{0,\rm exp}$ because of the essential non-linearity of the product function P(t) (**Figure 6**). For instance, if the idealistic curve P(t) is simulated with Equation (3) in N=425 equidistant points t_i from the time region $0 \le t/s \le 65$, the sampling time interval is $\Delta t = t_{i+1} - t_i = 0.153$ s. If the two first points are used, the "experimental" value $v_{0,\rm exp} = (P(t_1) - P(t_0))/\Delta t = 3.866 ~\mu \text{M/s}$ is by -0.9% smaller than the correct value $v_0 = 3.90 ~\mu \text{M/s}$ calculated by Equation (17). It is clear, the shorter the interval $\Delta t = t_1 - t_0$, the smaller the difference between $v_{0,\rm exp}$ and v_0 . Since in experiment the

first time point t_1 cannot be set arbitrarily close to $t_0 = 0$, the experimental value $v_{0,exp}$ is necessarily smaller than v_0 . In a more realistic case, when ten initial points are used to calculate $v_{0,exp}$ by linear regression, the underestimation of $v_{0 \text{ exp}}$ is by -8.2%. It is important to realize, that the initial part of P(t) is an essentially nonlinear function, see insert in Figure 6. Therefore, linear regression does not apply to determine v_0 . Alternatively, $v_{0,exp}$ can be obtained by fitting a second degree polynomial equation $P(t) = at^2 + bt + c$ to the data by nonlinear regression [4]. Extrapolation of the first derivative of the equation to $t \rightarrow$ 0 yields $v_{0 \text{ exp}} = b$. The nonlinear regression can improve precision of the assessment of $v_{0,exp}$ considerably. Still, the results on $v_{0,exp}$ are very prone to errors in initial estimates of the coefficients a, b and c, as well as to the number of data points and time range taken for the fit. For instance, when ten initial points are used to calculate $v_{0 \text{ exp}}$ Figure 6, at the initial parameters $a = -0.2 \text{ s}^{-2}$, $b = 3 \text{ s}^{-1}$ and c = 0, the underestimation of $v_{0,exp}$ is only -0.14%, at a = -0.5 s^{-2} , b = 1 s⁻¹ and c = 0 the underestimation is -0.16%. When twenty initial points are used to calculate $v_{0,exp}$, the underestimations are -0.56% and -0.58%, respectively.

1.4. Testing v₀-Plots and Single Trace Regression with Simulated Data

For a computational testing of commonly used v_0 -plots and the single-trace regression we used narrow and wide ranges of starting substrate concentrations S_0 (*c.f.* 2.3 and 2.4). A broader S_0 -range, up to $\sim 2~K_M$, is generally favorable for an analysis based on v_0 -plots compared to the analysis in a narrow range of S_0 used for the results presented in **Table 2** [5]. In **Figure 7** the Eadie-Hofstee (a,d), Lineweaver-Burk (b,e) and nonlinear regression

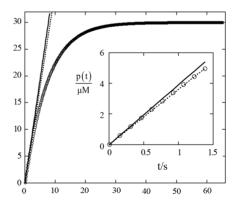


Figure 6. Product curve (circles) simulated by Equation (3) at $K_M = 220 \, \mu\text{M}$, $k_{\text{cat}} = 650 \, \text{s}^{-1}$, $E_0 = 50 \, \text{nM}$ and $S_0 = 30 \, \mu\text{M}$ for N = 425 equidistant time points with the time interval $\Delta t = 0.153 \, \text{s}$. The solid straight line refers to the linear term $v_0 \cdot t$, where the initial rate v_0 was calculated by Equation (17). The dotted line represents $v_{0, \exp} \cdot t$, where the initial "experimental" rate $v_{0, \exp}$ was calculated with the first ten points by linear regression; see inset.

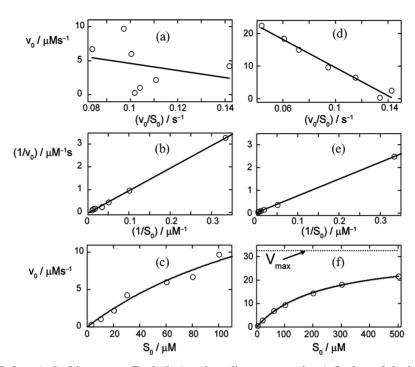


Figure 7. The Eadie-Hofstee (a,d), Lineweaver-Burk (b,e) and nonlinear regression (c,f) plots of the initial rates $v_{0,\rm exp}$, determined with the first ten points of the simulated relaxation P(t), Equation (3), in the range of S_0 from 3 to 100 μ M (left column) and of 3 to 500 μ M (right column) respectively; see Figure 6 for details. The noise was added to the function P(t) in form of the pseudo-random error of magnitude 2 μ M. The solid lines present fits with Equation (14), (a,d), Equation (15), (b,e), and Equation (17), (c,f). The resulting parameters K_M and $k_{\rm cat}$ are summarized in Table 3 and in Table 4 of Appendix. The dotted line in (f) indicates the value of the maximum rate $V_{\rm max} = 32.5 \ \mu$ M/s.

Table 4. Kinetic parameters determined by different methods from simulated data^a with pseudo-random noise for a wide range of S_0^b .

		$K_{M}/\mu M$			$k_{\rm cat}/{ m s}^{-1}$	
Noise/μM	0	≤0.5	≤2	0	≤0.5	≤2
Lineweaver-Burk	260	430 ± 100	110 ± 320	695	1000 ± 200	290 ± 600
Eadie-Hofstee	252	279 ± 11	239 ± 24	677	709 ± 12	612 ± 30
Nonlinear regression	251	261 ± 18	256 ± 25	675	684 ± 24	660 ± 30
Single trace regression ^c	220	220.4 ± 0.3^{d}	220 ± 1	650	650 ± 1	651 ± 1

^aGenerated with Equation (3) at $K_M = 220$ μM, $k_{cat} = 650$ s⁻¹, $E_0 = 50$ nM. The pseudo-random noise of the magnitude 0.5 μM or 2 μM was added to P(t), Equation (3). ${}^{b}S_0/\mu$ M = 3, 20, 60, 100, 200, 300, 500. Regression with Equation (8) at $S_0 = 250$ μM. Errors are indicated by the standard deviation determined from five different sets of noise.

 v_0 -plots (c,f) are shown for the S_0 -range from 0 to 100 μ M (*left side*) and from 0 to 500 μ M (*right side*), respectively. Notably, the K_M -value for the simulations was set to 220 μ M. The diagrams are based on data generated by the simulation of the enzymatic product curve P(t) with Equation (3) and a noise magnitude of 2 μ M.

It is evident in **Figure 7** that the data points obtained in the broad range of S_0 (d,e,f) deviate less from the regression curves compared to the respective plots obtained for the narrower S_0 -range (a,b,c). This observation is consistent with results of the corresponding plots presented in **Table 4**. Standard deviations of the K_M and k_{cat}

values in **Table 4** are significantly smaller compared to the corresponding deviations in **Table 2**. Additionally to **Table 2**, in **Table 4** the results of simulations at the noise-magnitudes of 0 and 2 μ M are presented too. However, the general features of the impact of the S_0 -range on quality of the analysis are throughout the same as in **Table 2**. The Lineweaver-Burk plot does not yield unphysical, negative results, yet provides the most imprecise results compared to other v_0 -plots. The Eadie-Hofstee and the nonlinear regression plot yield considerably more precise results than the Lineweaver-Burk plot. However, all v_0 -plots are not able to correctly determine the true

values of K_M and k_{cat} in the range of their standard deviations. At low noise-magnitudes, the results of the ν_0 -plots are generally overestimated. With increasing noise-level, this effect may be (over-) compensated by imprecision of results.

As in the case of the narrow S_0 -range, the single trace regression method yields reasonable results for all investigated noise-magnitudes and shows only nonsystematic errors of the determined K_M and k_{cat} values (see **Table 4**).

1.5. Testing *v*₀-Plots and Single Trace Regression with Experimental Data on C*c*O

The single trace regression method is able to determine K_M and $k_{\rm cat}$ for every single time course of an enzymatic reaction at a single S_0 -value, contrary to commonly used ν_0 -plots. In our case, the K_M and $k_{\rm cat}$ parameters are determined for time traces of the enzymatic reaction of C_cO with ferrous cytochrome c at seven different S_0 -values ranging from 2.68 to 59.35 μ M. The results of the new regression method are presented in **Table 5**; the corresponding mean values and standard deviations of K_M and $k_{\rm cat}$ are shown in **Table 3**. For the experimental curves no systematic deviations of the resulting K_M and $k_{\rm cat}$ are observed, as for the simulated data.

Table 5. Results of single trace regression^a of the enzymatic reaction of CcO with cytochrome c at different initial substrate concentrations S_0 .

$S_0/\mu M$	$K_{M,\mathrm{est}}/\mu\mathrm{M}$	$k_{\mathrm{cat,est}}/\mathrm{s}^{-1}$	
2.68	213.7	959.4	
5.33	221.9	914.6	
10.44	225.0	847.1	
23.65	222.4	740.9	
30.08	217.9	645.4	
37.20	214.4	724.5	
59.35	218.8	696.7	

^aWith Equation (8).

1.6. Mathcad Code for Michaelis-Menten-Lambert Analysis of a Single Progress Curve of Enzyme Kinetics

The complete Mathcad code for analysis of single progress curves of enzymatic reactions with the Michaelis-Menten-Lambert equation can be found on the homepage of the corresponding author:

http://www.uni-bielefeld.de/chemie/arbeitsbereiche/pc3-hellweg/work/sergej/MathcadCode.pdf

The Mathcad code begins with a generation of the Lambert-W function W(x). For this purpose we use the standard Mathcad function "wurzel" to solve the equation $ye^y - x = 0$ with an initial value of y = 0.1 and tolerance $= 10^{-15}$.

- a) The second part of the code starts with an input of the total substrate concentration S_0 as well as the total enzyme concentration E_0 , and the upload of an experimental product progress curve P(t).
- b) An initial value of the apparent rate constant m is estimated from the experimental normalized product curve $p(t) = P(t)/S_0$. Here we use the exponential approximation $p_{\text{exp}}(t) = 1 \exp(-mt)$ and the 2/3-amplitude criterion to estimate the characteristic relaxation time $\tau = 1/\text{m}$ of p(t). To obtain a more precise estimate of m we use the initial estimate of m further as an input parameter for a nonlinear fit of $p_{\text{exp}}(t)$ to the experimental product curve p(t). Therefore, the standard function "Minfehl" implemented in Mathcad is applied.
- c) An initial reaction rate v_0 of the product curve p(t) is determined by the standard function "neigung" using the first 11 points of p(t).
- d) An estimate of the dimensionless input parameter s is calculated by $s = |m/v_0 1|$.
- e) Exact values of the Michaelis-Menten parameters K_M and k_{cat} are determined by the "Minfehl" function using the experimental p(t)-curve and the estimates of m and s as input parameters.

In the third part, the calculated Michaelis-Menten parameters K_M , $k_{\rm cat}$ and $V_{\rm max}$ are summarized together with results of the fit of the integrated Michaelis-Menten Equation (8) to the experimental p(t)-curve.

Abbreviations

CcO: cytochrome c oxidase;

 E_0 : Total enzyme concentration;

 S_0 : Starting substrate concentration;

 k_{cat} : Catalytic constant, a key parameter in enzymatic analysis;

 K_M : Michaelis-Menten constant, a key parameter in enzymatic analysis;

 $K_{M,\text{input}}$, $k_{\text{cat,input}}$: Input parameters for K_M and k_{cat} for their estimation by nonlinear regression;

 $K_{M,\text{est}}$, $k_{\text{cat,est}}$: Estimated values of K_M and k_{cat} by nonlinear regression;

m: The apparent rate constant, defined as $m = k_{\text{cat}} \cdot E_0 / K_M$;

s: The dimensionless parameter, defined as $s = S_0/K_M$;

P(t): Progress curve of the product concentration of an enzymatic reaction;

p(t): Normalized product concentration, defined as $p(t) = P(t)/S_0$, ranging from 0 to 1;

 v_0 : Initial reaction rate of the progress curve;

 v_0 -plots: Methods for analyzing enzyme kinetics based on a variety of initial rates v_0 at different S_0 .

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