

CHEMICAL KINETICS  
AND CATALYSIS

Study of the Kinetic Parameters for Synthesis and Hydrolysis  
of Pharmacologically Active Salicin Isomer Catalyzed  
by Baker's Yeast Maltase<sup>1</sup>

D. V. Veličković<sup>a</sup>, A. S. Dimitrijević<sup>b</sup>, F. J. Bihelović<sup>b</sup>, R. M. Jankov<sup>b</sup>, and N. Milosavić<sup>b</sup>

<sup>a</sup> Innovation Center of the Faculty of Chemistry, University of Belgrade, Studentski trg 12, Belgrade, Serbia

<sup>b</sup> Faculty of Chemistry, University of Belgrade, Studentski trg 12, Belgrade, Serbia

e-mail: nenadmil@chem.bg.ac.rs

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**Abstract**—One of the key elements for understanding enzyme reactions is determination of its kinetic parameters. Since transglucosylation is kinetically controlled reaction, besides the reaction of synthesis, very important is the reaction of enzymatic hydrolysis of created product. Therefore, in this study, kinetic parameters for synthesis and secondary hydrolysis of pharmacologically active  $\alpha$  isosalicin by baker's yeast maltase were calculated, and it was shown that specificity of maltase for hydrolysis is approximately 150 times higher than for synthesis.

**Keywords:** synthesis and hydrolysis of salicin isomer, catalysis by maltase, kinetic parameters.

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INTRODUCTION

Many glycoside hydrolases exhibit transglycosylation activity, and transglycosylation reactions are well known and widely used methods for glycoside synthesis [1]. Transglycosylation is a mechanism for glycosidic bond formation in which a glycosyl moiety from a donor compound is transferred to the hydroxyl group of an acceptor molecule other than water [2] (Fig. 1). Transglucosylation is subtype of transglycosylation, where enzyme transfers glucose moiety to acceptor molecule.

Principally, all glycosidases can be divided in two classes, depending whether hydrolysis occurs with retention or with inversion of configuration at the anomeric center. Maltase (EC 3.2.1.20;  $\alpha$ -glucosidase) belongs to the family of glycosidases which hydrolyze substrate and form products with the same configuration at the anomer atom like substrate [3]. It is one of the most abundant glycosyl hydrolases present in baker's yeast (*Saccharomyces cerevisiae*), and has been used for synthesis of various pharmacologically active glycosides [4]. Sometimes, the glycosidic residue is crucial for glucosides activities; in other cases glycosylation improves pharmacokinetic parameters [5]. We used maltase for synthesis of  $\alpha$ -isosalicin [2-hydroxybenzyl- $\alpha$ -D-glucopyranoside] with *ortho*-hydroxybenzyl alcohol as glucose acceptor and maltose as glucose donor in reaction of transglucosylation (Fig. 1). Salicin, [2-(hydroxymethyl)phenyl- $\beta$ -D-glucopyra-

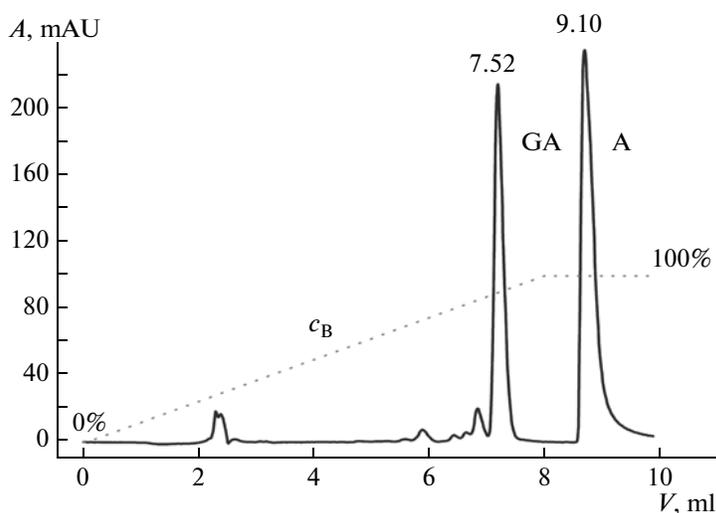
noside] and its structural isomers exhibit pronounced anticoagulant activity [5], tyrosinase inhibitory activity [6], and are also used as anti-inflammatory, analgesic and antipyretic prodrugs [7, 8]. Moreover,  $\alpha$ -glucoside of *ortho*-hydroxybenzyl alcohol ( $\alpha$ -isosalicin) is more potent anticoagulant than naturally occurring salicin [5].

Alpha-glucoside of *ortho*-hydroxybenzyl alcohol, like all other glycosides, is formed in a kinetically controlled reaction that shows optimum in time, when disaccharide is used as donor [9]. Because the reaction is controlled kinetically, it is possible to overshoot the equilibrium conversion of reactant into product. As the reactant is consumed the concentration of the product will reach the peak when rates for its synthesis and debenzilation become equal. At this point kinetic control is lost and the reaction should be stopped before thermodynamic control takes over and the product undergoes enzymatic hydrolysis (secondary hydrolysis in Fig. 1) [3]. Hence, the yield of glucoside is determined by a delicate balance between the rates of [E-G + A] synthesis ( $K_1, k_3$ ) and hydrolysis ( $k_4, k_{-3}, K_7, k_2$ ) on the one hand, and product hydrolysis ( $k_6, k_5, k_{-5}$ ), on the other.

Determination of kinetic parameters that lies in  $K_1$  is earlier described for ( $\beta$ -galactosidase [2], as well as reaction driven with  $K_7$  for  $\beta$ -glucosidase (substrate transglycosylation) [10], although this reaction does not have influence in transglucosylation reaction catalyzed by maltase from baker's yeast. Reaction of inhi-

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**Fig. 2.** One of the HPLC chromatograms for transglucosylation reaction with *ortho*-hydroxybenzyl alcohol (A, retention volume 9.10 ml) as glucose acceptor and  $\alpha$ -isosalicin (GA, retention volume 7.52 ml) as product of reaction. Yield of reaction was calculated from peak area ratio at 268 nm between GA and A. This chromatogram is obtained when initial concentration of *ortho*-hydroxybenzyl alcohol was 30 mM and reaction was stopped after 30 min. Conditions for HPLC chromatography are described in EXPERIMENTAL section.

activity of enzyme and represent number of molecules of substrate that one molecule of enzyme transforms in one second. Knowing total concentration of enzyme ( $E_0$ ) and  $V_{\max}$  is easy to calculate  $k_{\text{cat}}$  (Eq. 2). Values for specific “coef” and “const” are represented below:

$$\begin{aligned} \text{const} &= k_2(k_6 + k_{-5})(k_4 + k_{-3}), \\ (\text{coefA}) &= k_3k_4(k_{-5} + k_6), \\ (\text{coefGA}) &= k_5(k_4 + k_{-3})(k_6 + k_2), \\ (\text{coefM}) &= K_1(k_6 + k_{-5})(k_4 + k_{-3}), \\ (\text{coefGA} \cdot \text{A}) &= k_3k_5(k_4 + k_6), \\ (\text{coefA} \cdot \text{M}) &= k_3K_1(k_{-5} + k_6); \end{aligned}$$

[A] is concentration of *ortho*-hydroxybenzyl alcohol; [GA] is concentration of  $\alpha$ -isosalicin; [M] is concentration of maltose.

The product formation rate ( $d[\text{GA}]/dt$ ) for each substrate concentration was calculated from HPLC chromatograms (Fig. 2) Michaelis constant and  $k_{\text{cat}}$  are determined from Lineweaver–Burk (LB) plot, derived from Michaelis–Menten (MM) plot (Fig. 3).

**Determination of kinetic parameters for hydrolysis of  $\alpha$ -isosalicin.** Like in case of synthesis, using King and Altman modified Michaelis–Menten equation based on Fig. 1, following kinetic parameters are given for secondary hydrolysis of created product as:

$$K_{m_h} = \frac{\text{const} + (\text{coefA})[\text{A}] + (\text{coefM})[\text{M}] + (\text{coefA} \cdot \text{M})[\text{A}][\text{M}]}{(\text{coefGA}) + (\text{coefGA} \cdot \text{A})[\text{A}]}, \quad (3)$$

where  $K_{m_h}$  represent Michaelis constant for [E + GA] analog that for the synthesis reaction,

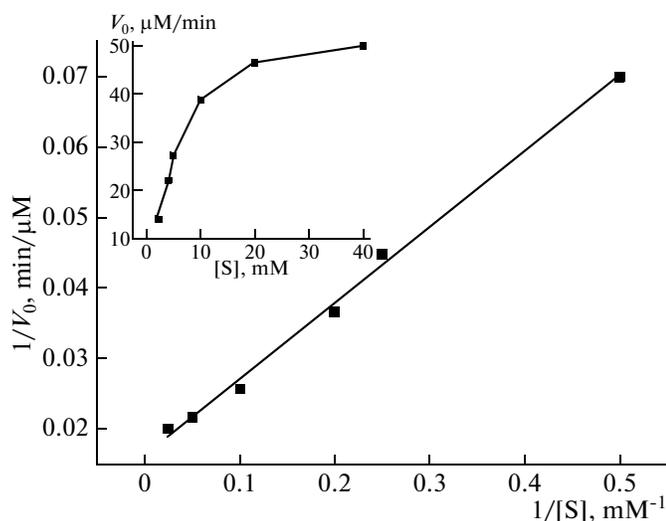
$$k_{\text{cat}_h} = \frac{V_{\max_h}}{E_0} = \frac{k_5k_6(k_3k_4[\text{A}] + k_2k_4 + k_{-3}k_2)}{(\text{coefGA}) + (\text{coefGA})[\text{A}]}, \quad (4)$$

where  $k_{\text{cat}_h}$  is turnover number for [E + GA] and also has analog meaning as in case of synthesis.

The specific “Coef” and “const” are the same as in explanation of synthetic reaction.

Michaelis constant was determined using one of integrated forms of Michaelis–Menten equation [14]. The traditional technique for determination of enzyme kinetic parameters is to perform numerous

experiments measuring the reaction rate,  $V$ , as a function of the substrate starting concentration. In a well-known manner, which we used to described synthesis.  $K_m$  and  $V_{\max}$  can be obtained by plotting  $1/V$  in a function of  $1/[S]$  (the LB plot). To determine the reaction rate by graphical differentiation of the substrate concentration–time curve, at least several experiments must be performed. This procedure is not only tedious but also inaccurate, especially if a too high substrate conversion during the measurement causes a flat curve instead of a straight line in the substrate concentration–time course [14–17]. The integrated form of Michaelis–Menten method for enzyme kinetic parameter determination is direct and more convenient. This method enables establishment of  $K_m$  and



**Fig. 3.** LB and MM plot for synthesis of  $\alpha$ -isosalicin;  $[M] = 1.2 \text{ M}$ ;  $K_m = 6.62 \text{ mM}$ ,  $V_{\max} = 0.616 \text{ } \mu\text{M}/(\text{min mg})$ .

$V_{\max}$ , by only one experiment, in which the change of substrate concentration in time is monitored:

$$\frac{\tau}{U} = \frac{K_m}{V_{\max}} \left( \frac{1}{U} \ln \frac{1}{1-U} - 1 \right) + \frac{c_0 + K_m}{V_{\max}}, \quad (5)$$

where  $\tau$  is the time point of the reaction,  $U$  is substrate conversion, and  $c_0$  is initial substrate concentration [14].

Results obtained using this form of integrated equation do not depend on the concentration of sub-

strate used, and therefore allows the use of concentration that are far below  $K_m$ . In this way it is possible to use only one, an arbitrary, concentration of substrate, thus saving the total consumption of high cost substrate, like  $\alpha$ -isosalicin is, in determination of the constants.

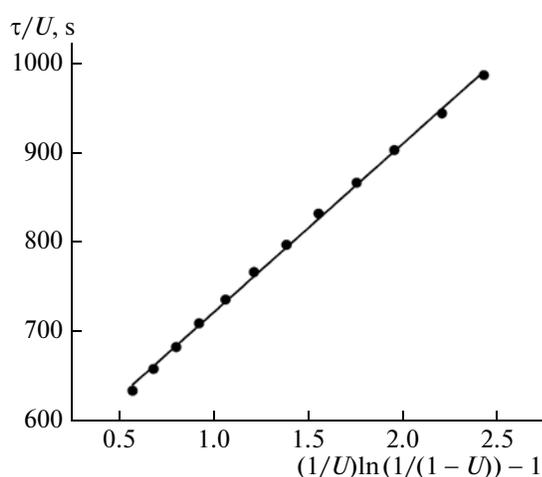
As the maltase is “the most active” toward 4-nitrophenyl- $\alpha$ -D-glucopyranoside [18], so any other substrate can be considered as competitive “inhibitor” for hydrolysis 4-nitrophenyl- $\alpha$ -D-glucopyranoside [19]. Applying Eq. (5) to calculate  $K_m$  and  $V_{\max}$  for 4-nitrophenyl- $\alpha$ -D-glucopyranoside, values obtained matched literature values [18] (Fig. 4).

The results obtained for 4-nitrophenyl- $\alpha$ -D-glucopyranoside in presence of  $\alpha$ -isosalicin are shown in Fig. 5.

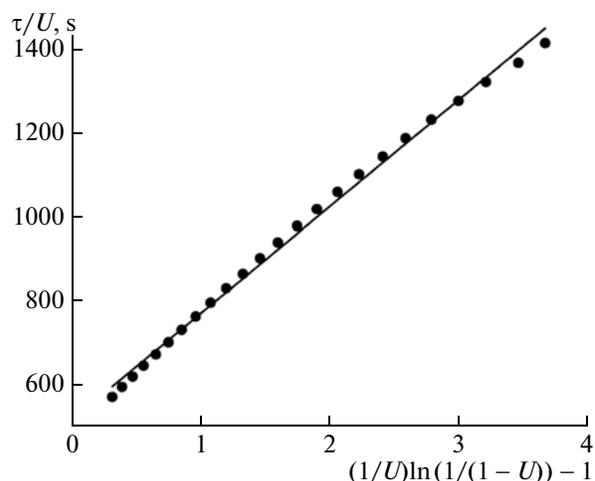
As it was expected,  $K_m$  for hydrolysis 4-nitrophenyl- $\alpha$ -D-glucopyranoside increased (for factor  $\alpha$ ) and the  $V_{\max}$  remained the same, suggesting that it is a competitive inhibition [11]. In the case of the competitive inhibition

$$\alpha = \frac{K_m^{\text{app}}}{K_m} = \frac{1 + [I]}{K_i}, \quad (6)$$

where  $[I]$  is concentration of inhibitor, i.e. in our case  $\alpha$ -isosalicin, and  $K_i$  is inhibition constant, i.e. “decomposition constant” for enzyme inhibitor complex, or on Fig. 1, “decomposition constant” for  $[E + \text{GA}]$  complex. Because “decomposition constant” for  $[E + \text{GA}]$  complex actually represents  $K_{m_h}$  for hydrolysis  $\alpha$ -isosalicin,  $K_{m_h}$  and  $K_i$  can be equated. Using Eq. (6) we calculated that  $K_{m_h}$  is 0.61 mM.



**Fig. 4.** Plot of integral form of Michaelis–Menten equation for 0.45 mM 4-nitrophenyl- $\alpha$ -D-glucopyranoside as substrate without inhibitor;  $K_m = 0.27 \text{ mM}$ ,  $V_{\max} = 25 \text{ } \mu\text{M}/(\text{min mg})$ . Concentration of enzyme used was 75 U/ml.



**Fig. 5.** Plot of integral form for Michaelis–Menten equation for 0.45 mM 4-nitrophenyl- $\alpha$ -D-glucopyranoside as substrate in the presence of 0.35 mM  $\alpha$ -isosalicin as the second substrate (i.e. inhibitor),  $K_m^{\text{app}} = 0.45 \text{ mM}$ ,  $V_{\max}^{\text{app}} = 25 \text{ } \mu\text{M}/(\text{min mg})$ . Concentration of enzyme used was 100 U/ml.

Kinetic constants for transglucosylation reaction catalyzed by maltase

A isosalicin		
Constant	synthesis	hydrolysis
$K_m$ , mM	6.62	0.61
$V_{max}$ , $\mu\text{mol}/(\text{min mg})$	0.616	8.30
$k_{cat}$ , $\text{s}^{-1}$	2.5	33.67
$K_{spec}$ , $\text{s}^{-1}/\text{mM}^{-1}$	0.38	55.2

Note:  $K_{spec}$  represents real measure of affinity of some enzyme toward its substrate and it is equal to the ratio between  $k_{cat}$  and  $K_m$ .

Because  $V_{max}$  is the rate of the reaction when all enzyme molecules are in complex with its substrate, it is possible to determine  $V_{max}$  in the high excess of substrate ( $[S] \geq 10 K_m$ ). Therefore  $V_{max_h}$  for this reaction was determined from concentration of 6 mM  $\alpha$ -isosalicin ( $10 K_m$ ), and calculated to be  $V_{max} = 8.3 \mu\text{M}/(\text{min mg})$ .

Results for hydrolysis and synthesis of  $\alpha$  glucoside of salicyl alcohol are summarized in table.

## CONCLUSION

To the best of authors' knowledge, this is the first time that detailed kinetic study of transglucosylation activity of a glucosidase from *Saccharomyces cerevisiae* was examined. Also this is the first time ever that by using kinetic parameters, secondary hydrolysis of created product by  $\alpha$ -glucosidase was described, although it is crucial for maximal yield of transglucosylation. Constants calculated in our study are derivative from our experimental condition, but it is obvious (Eqs. (1)–(4)) that in other concentrations of maltose, or maltose and alcohol in case of hydrolysis, these constants are different.

In order for enzyme to work with maximal rate, minimal alcohol concentration needed for the synthesis of  $\alpha$ -isosalicin is about 10 mM, as shown in table. It would be wrong to conclude that when concentration of the product reaches near 1 mM and concentration of alcohol is 10 times smaller than product, secondary hydrolysis starts. It should be emphasized that large amount of maltose is present in the reaction mixture (1.2 M), and as  $K_m$  for maltose is 80 mM [18] enzyme practically "does not feel" the  $\alpha$ -isosalicin. Secondary hydrolysis starts when concentration of

maltose drops to value comparable to  $\alpha$ -isosalicin. At this point the reaction should be stopped before the system enters thermodynamic equilibrium.

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