

Hydrolysis Kinetics of Oil Palm Empty Fruit Bunch in Ionic Liquids and Cellulase Integrated System

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Abstract

Ionic liquids (ILs) are developing as potential solvents in lignocellulose solvation, which enables cellulase accessibility into the substrate. Nevertheless, ILs could result in enzyme deactivation because of the high polarity. Therefore, developing a system of ILs-compatible cellulase (IL-E) to promote lignocellulose conversion into sugars is a challenge in ILs applications. This study used an IL-E to attain high conversion yield of sugars from oil palm empty fruit bunch (EFB). Cellulase (*Tr-Cel*) from *Trichoderma reesei* was stable in the ILs, 1-ethyl-3-methyl imidazolium diethyl phosphate [EMIM]DEP and choline acetate [Cho]OAc. The inhibition and deactivation of cellulase were evaluated using the model substrate, carboxymethyl cellulose (CMC) and EFB as a lignocellulosic material to assess the hydrolytic activity. The enzyme kinetics revealed that [Cho]OAc acted as a noncompetitive inhibitor. Additionally, [EMIM]DEP may not be considered as an inhibitor as it increases the V_{max} and does not significantly affect the K_M . In both cases, the study proved that IL did not result in a severe loss of cellulase activity, which is a promising outcome for one-pot hydrolysis of lignocellulosic materials.

Keywords: affinity, activity, ionic liquid, inhibition, cellulase, lignocellulose, hydrolysis

1. Introduction

It is prominent that lignocellulose is one of the most beneficial and important renewable biomaterials which offers a sustainable substitute for fossil fuels resources (Varanasi et al., 2012) as it consists the desirable carbohydrates, cellulose and hemicellulose (Mamman et al., 2008). Oil palm empty fruit bunch (EFB) is the prime solid waste after oil extraction. Holocellulose contributes to 60-70% of the total weight of EFB. A biological conversion is required to utilize these components, however, as a result of the complex structure, high cost and lengthy treatment are required for the processing (Alonso, Bond, & Dumesic, 2010). Diverse approaches have been implemented in the pretreatment of lignocellulose including chemical (Pellera & Gidaracos, 2018) such as acidic (Harun & Danquah, 2011) and alkali treatment (Zainan, Alam, & Al-Khatib, 2013), physical or mechanical treatment (Barakat et al., 2014), physicochemical (Brodeur et al., 2011) and biological methods (Balat, 2011) using microbes (Alam, Kabbashi, & Hussin, 2009; Galbe & Zacchi, 2007). Nevertheless, these methods are associated with various disadvantages as they consume high energy, might be toxic, or may be pricey (Muhammad et al., 2014; Yang et al., 2010). More drawbacks include disposal challenge, cost of recovery and slow rate of reaction (Balat, 2011). In contrast to chemical hydrolysis, enzymatic hydrolysis is an eco-friendly substitute (Salihu & Alam, 2015). Though, conventional solvents could result in loss of enzyme activity and are associated with the environmental threats which directed the concern towards green solvents. In this context, ionic liquids (ILs) are good solvents for complex carbohydrates and offer many attractive properties such as stabilizing enzymes (Elgharbawy, Alam, Moniruzzaman, & Goto, 2016; Fu, Mazza, & Tamaki, 2010). Wang et al. (Wang, Radosevich, Hayes, & Labbé 2011) reported that some cellulases were stabilized in IL when examined in [EMIM]OAc (15%) in the saccharification process of yellow poplar biomass, and [EMIM]OAc (10-20%) for enzymatic hydrolysis of switchgrass (Shi et al., 2013). Additionally, many studies have reported cellulases stability in systems that

are IL-E compatible, for instance, cholinium-based ILs (Ninomiya et al., 2015). Furthermore, single-step hydrolysis is the desired process where IL pretreatment of lignocellulose is integrated with enzymatic hydrolysis for bioethanol production, as it eliminates the regeneration stage of the cellulose.

Two main paths may demonstrate the enzyme-catalyzed hydrolysis upon biomass IL-pretreatment, which include the regeneration of cellulose from the IL solution, to undergo enzymatic hydrolysis (Tan, Lee, & Mohamed, 2011; Zhao et al., 2009). The second path is regarded as a single-step process where the hydrolysis is directly performed in the IL in a water-based buffer solution by cellulase enzymes (Gunny, Arbain, Edwin Gumba, Jong, & Jamal, 2014). Various ILs showed promising results in lignocellulose structure modification and lignin removal, such as choline acetate [Cho]OAc (Asakawa, Kohara, Sasaki, Asada, & Nakamura, 2015). A group of researchers have tried to use IL-surfactant ([BMIM]Cl+ PEG-8000) to facilitate sugarcane bagasse dissolution using both cellulase and xylanase, which are generated in house from *Aspergillus assiutensis* VS34. The bagasse was pretreated at 90 °C for 2 h, followed by enzymatic hydrolysis with both enzymes. Ninety percent of the activity was maintained in the IL (Sharma, Nargotra, & Bajaj, 2019). This shows the ability of IL to cater for compatibility with some enzymes (Elgharbawy, Alam, Moniruzzaman, et al., 2016; Ibrahim, Moniruzzaman, Yusup, & Uemura, 2015). On the contrary, Cellic® Htec2 cellulase acting on CMC was deactivated in the presence of [BMIM]Cl but without surfactant, showing that [BMIM]Cl is a competitive inhibitor (Nemestóthy et al., 2017).

2. Method

2.1 Materials

Sime Darby Plantation, Malaysia provided the oil palm empty fruit bunch (EFB) raw material used in this study. Cellulase (*Tr-Cel*) from *Trichoderma reesei* ATCC 26921, lyophilized powder (≥ 1 unit/mg solid) was purchased from Sigma-Aldrich, USA. Choline acetate [Cho]OAc was synthesized as reported in the previous publication (Elgharbawy, Alam, Jamal, Kabbashi, & Moniruzzaman, 2016). 1-Ethyl-3-methyl imidazolium diethyl phosphate [EMIM]DEP was obtained from Merck, Germany.

2.2 Determination of Kinetics Parameters of *Tr-Cel* with CMC and EFB

Tr-Cel was incubated for 60 min in various concentrations of ILs (10, 20, 40, 60, 80 and 100%). After incubation, the solution was subjected to cellulase activity assay using CMC as described by Salvador et al. (Salvador, Santos, & Saraiva, 2010).

Dry EFB (500 mg) was weighed in a glass vial, and following IL addition the mixture was incubated for 60 min at 90 °C, cooled to room temperature and a buffer solution (sodium citrate buffer pH 4.8 ± 0.2 , 50 mM) was added to the tubes to attain a solution with 10% IL. Lastly, the cellulase (*Tr-Cel*) was added to start the enzymatic hydrolysis.

2.2 Determination of Kinetics Parameters of *Tr-Cel* with CMC and EFB in IL

The kinetic parameters of *Tr-Cel* were studied by monitoring the IL-E at various concentrations with 500 mg of both CMC and EFB in 50 mM sodium citrate buffer (60 min, 50 °C, pH 4.8 ± 0.2) and different loading of the enzyme (10, 20, 30, 40 and 50 units). The linear equation of Michaelis-Menten was used to calculate the parameters (V_{\max} and K_M), graphically based the velocity (V) of the enzyme-catalyzed reaction, at different substrate concentrations $[S]$. The enzyme activity was fixed at 50 units (50 FPU g^{-1}) while varying the loading of substrates 100, 300, 500, 700 and 1000 mg and then measuring the concentration of sugar.

CMC substrate was prepared at these concentrations: 1.0, 2.0, 3.0, 4.0, 5.0, 6.0, 8.0, 10.0 g/L. The initial enzyme velocity was assessed using cellulase activity assay method (DNS reagent) (Zhang, Hong, & Ye, 2009). The reciprocal of substrate concentration $1/[S]$ (X-axis) and the reciprocal of initial enzyme velocity $1/V$ as (Y-axis), were plotted to obtain the Lineweaver-Burk double reciprocal graph. Lineweaver-Burk model was used as it is a common tool to determine important terms in enzyme kinetics, such as V_{\max} and K_M .

For EFB substrate, enzymatic activity was analyzed using linear Michaelis-Menten model to determine the kinetics parameters V_{\max} and K_M . The model with the largest determination coefficient (R^2) was regarded as the best fitted to describe the enzyme kinetics.

3. Results and Discussion

3.1 Determination of Kinetics Parameters of *Tr-Cel* with CMC and EFB

The main function of an enzyme is to accelerate the rate of reaction. Kinetic parameters are essential to the understanding of enzyme activity and function. Figure 1 shows the kinetic models of *Tr-Cel* using CMC as a model substrate for the linear Michaelis-Menten model. The kinetic parameters for each linear model are summarized in Table 1.

Table 1. Kinetics parameters for *Tr-Cel* with two different substrates: EFB and CMC

Model	K_M (mg. mL ⁻¹)		V_{max} (mg. mL ⁻¹ min ⁻¹)		R^2		
	Substrate	EFB	CMC	EFB	CMC	EFB	CMC
Lineweaver-Burk		0.184	0.045	11.19	68.49	0.9592	0.9059
Hanes-Woolf		1.00	1.00	1.345	9.65	0.9961	0.9209
Eadie-Hofstee		48.38	18.18	29.63	37.43	0.6058	0.8736

The value of K_M mainly measures the enzyme affinity to a substrate. The smaller the K_M value, the greater the affinity of the enzyme for its substrate while the V_{max} parameter is an indication of the maximum velocity once the substrate occupies all the active sites of the enzyme. Usually, most of the K_M values are expressed in mM (mmol. L⁻¹) and the value of K_M is converted from mg. mL⁻¹ to mM or μ M using the molecular weight of the substrate. However, the compared substrate (EFB) does not have a well-recognized molecular weight; hence, the values were expressed in mg mL⁻¹.

Comparing the K_M and V_{max} values in case of CMC, using Hanes-Woolf and Lineweaver-Burk models, a higher value of V_{max} was obtained when the enzyme was modelled with Lineweaver-Burk (68.49 mg.mL⁻¹.min⁻¹). Likewise, K_M value was the lowest when the enzyme was modelled with Lineweaver-Burk. Eadie-Hofstee model did not show a good fit for the enzyme with either substrate. The plots can be seen in Figure 1.

In comparison, the EFB showed a good fit with both Lineweaver-Burk and Hanes-Woolf, although the values of K_M and V_{max} were more comparable with Lineweaver-Burk. It can also be seen that K_M resulted in the same values in both cases when modelling the enzyme with Hanes-Woolf; therefore, Lineweaver-Burk was used to observe the contrast trend. The values varied with EFB as the enzyme fitted into both Lineweaver-Burk and Hanes-Woolf models with high R^2 values. It can be seen that the K_M obtained was similar in the Hanes-Woolf model for both substrates although V_{max} was greater with CMC.

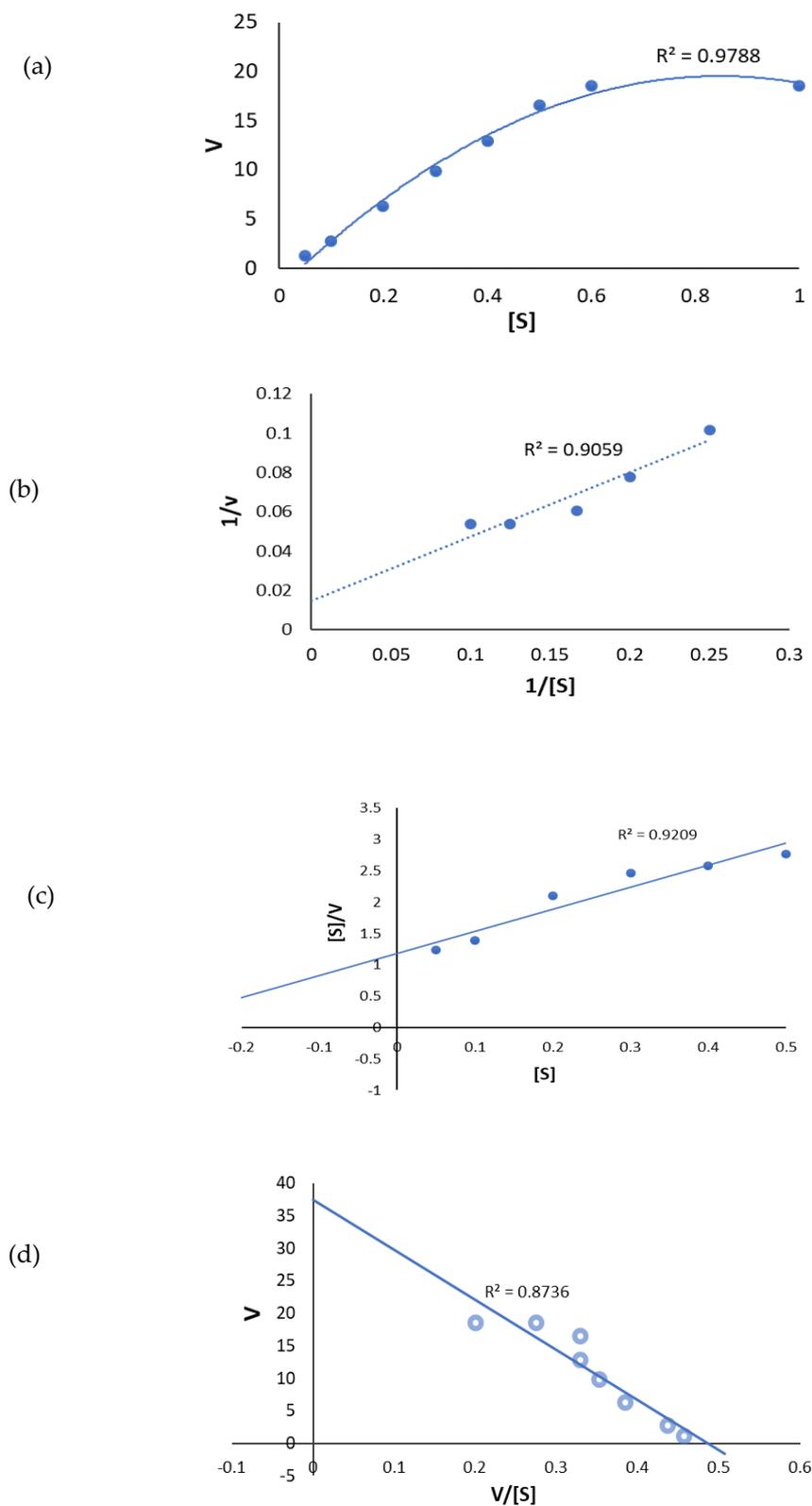


Figure 1. Kinetic model of (a) Michaelis-Menten hyperbolic plot, (b) Lineweaver-Burk plot, (c) Hanes-Woolf, (d) Eadie-Hofstee for in-house produced cellulase (*Tr-Cel*) using CMC as the substrate model, $[S]$ = substrate concentration (mg. mL^{-1}), V = reaction velocity ($\text{mg. mL}^{-1} \text{min}^{-1}$). R^2 = coefficient of determination

In this case study, it appears that the enzyme has more affinity towards CMC compared to EFB. This, to a great extent, could be explained as CMC has always been a model substrate for cellulases, and is considered as a good substrate (Wang, Post, Mayes, Frerichs, & Sindhu, 2012). The lower K_M might also be related to the complex structure of the lignocellulosic material. Adsorption of lignin to the surface of cellulase might occur in the case of EFB which occupies the active site in the absence of the IL (Noori & Karimi, 2016).

3.2 Determination of Kinetics Parameters of Tr-Cel with CMC and EFB in IL

Pretreatment with IL has been regarded as a promising approach that promotes the lignocellulose saccharification; nevertheless, the presence of IL in the medium could inhibit the function of cellulase. Hence, an investigation is required on the occurrence of inhibition by measuring both the reaction velocity and enzyme affinity to the substrate. Consequently, the kinetic parameters were assessed in the presence and in the absence of the ILs to describe the IL impact on the activity of cellulase during the one-step hydrolysis. The inhibition trend (Figure 2 and Figure 3) shows the pattern of inhibition of both [EMIM]DEP and [Cho]OAc, on the cellulase for the 60 min hydrolysis of CMC and EFB.

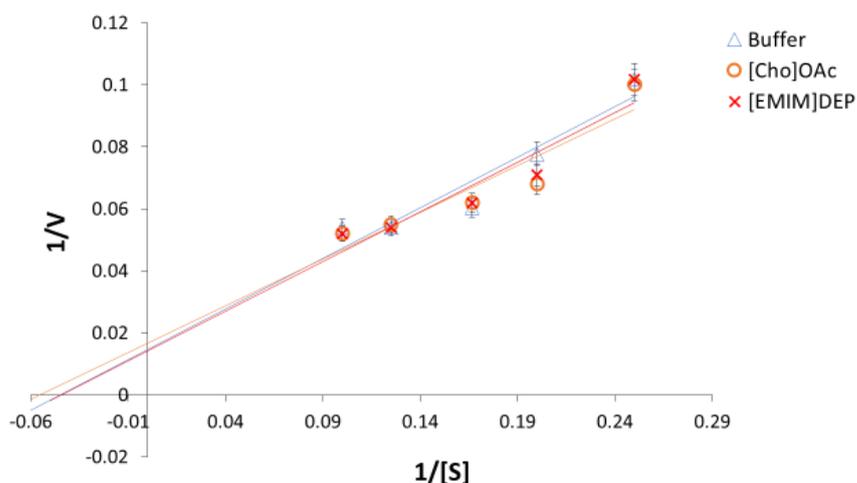


Figure 2. Lineweaver-Burk plot for the inhibition trend of cellulase in [EMIM]DEP, [Cho]OAc compared to buffer solution using carboxymethyl cellulose (CMC) as the substrate for hydrolysis. [S]= substrate concentration (mg. mL^{-1}), V = reaction velocity ($\text{mg. mL}^{-1} \text{min}^{-1}$). Conditions: pretreatment: $90\text{ }^{\circ}\text{C}$, 60 min; enzymatic hydrolysis: $\text{pH } 4.8 \pm 0.2$, 50 mM, $50\text{ }^{\circ}\text{C}$, 60 min, 50 FPU. g^{-1})

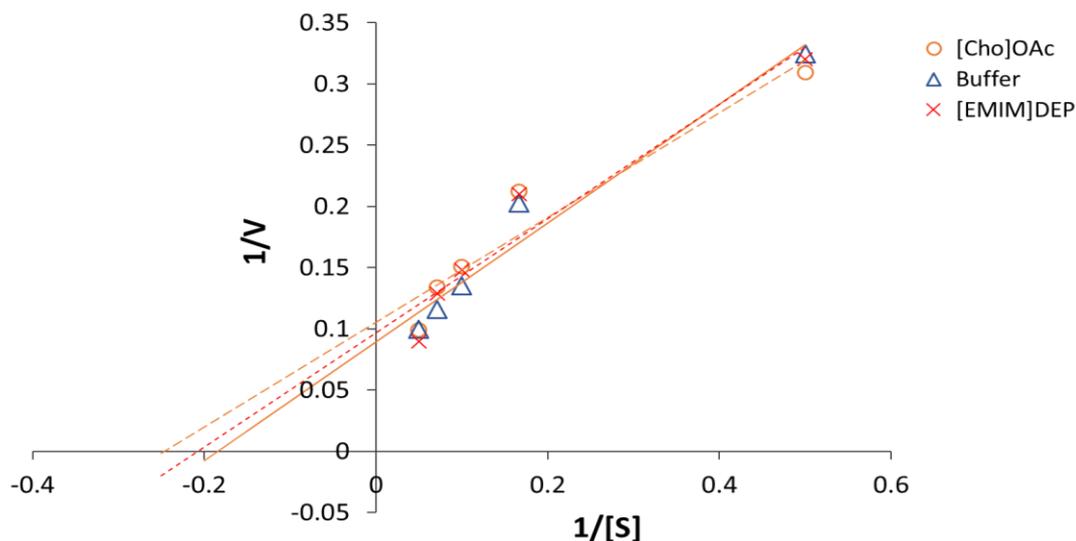


Figure 3. Lineweaver-Burk plot for the inhibition trend of cellulase in [EMIM]DEP, [Cho]OAc compared to buffer solution using EFB as the substrate for hydrolysis. $[S]$ = substrate concentration (mg mL^{-1}), V = reaction velocity ($\text{mg. mL}^{-1} \text{min}^{-1}$)

ILs resemble the mixed inhibitors where it is thought to occur when the inhibitor (IL) binds at a separate site other than the active site to either the enzyme-substrate complex or free enzyme (Figure 4).

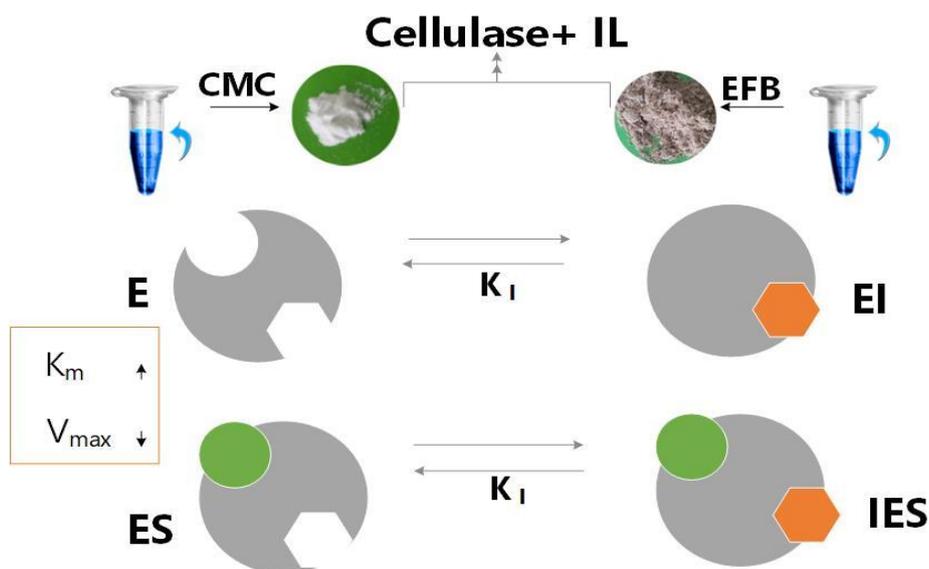
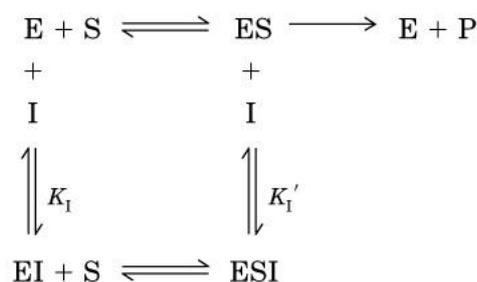


Figure 4. Illustration of mixed non-competitive inhibition in the cellulase-IL system

We concluded that the inhibitor fits reversible, mixed inhibition in which is recognized when the K_M and V_{\max} are both affected. It can be seen that V_{\max} is reduced as in noncompetitive inhibition whereas K_M increased slightly, which can be described by mixed inhibition. The maximum velocity decreased when the hydrolysis was conducted in choline acetate [Cho]OAc, however, K_M slightly increased, which points to less affinity towards the substrate. As calculated from the linear plot, K_M rose from 0.045 to 0.056 mg. mL^{-1} while V_{\max} got reduced from 68.49 to 59.88 $\text{mg. mL}^{-1} \text{min}^{-1}$. Therefore, it supports the hypothesis of mixed inhibition as both K_M and V_{\max} changed. In the presence of [EMIM]DEP, the velocity of cellulase slightly increased to 70.42 $\text{mg. mL}^{-1} \text{min}^{-1}$ when K_M was not significantly affected.

The reaction sequence can be written as in Scheme 1 (Henderson, 1972):



Scheme 1. Enzyme reaction sequence in the presence of the inhibitor (IL) (Henderson, 1972)

At equilibrium, K_M can be expressed as:

$$K_M = \frac{[\text{E}][\text{S}]}{[\text{E.S}]} \quad (1)$$

$$K_M = \frac{[\text{E.I}][\text{S}]}{[\text{E.S.I}]} \quad (2)$$

$$K_I = \frac{[\text{E}][\text{I}]}{[\text{E.I}]} \quad (3)$$

$$K_I = \frac{[\text{E.S}][\text{I}]}{[\text{E.S.I}]} \quad (4)$$

As enzyme concentration is not changing; hence, it can be stated as:

$$[\text{E}_T] = [\text{E}] + [\text{E.S}] + [\text{E.I}] + [\text{E.S.I}] \quad (5)$$

By substituting concentrations with the corresponding equilibrium terms and rearranging Equation 5:

$$[\text{E}] = \frac{[\text{E}_T]}{1 + \frac{[\text{S}]}{K_M} + \frac{[\text{I}]}{K_I} + \frac{[\text{S}][\text{I}]}{K_I K_M}} \quad (6)$$

The product formation rate $r_p = \frac{k[\text{E.S}]}{K_M}$, therefore:

$$r_p = \frac{k[\text{E}_T][\text{S}]}{K_m} \quad (7)$$

moreover, $k[\text{E}_T]$ can be expressed as V_{\max} , and the Equation (3.8) turns out to be:

$$V = \frac{V_{\max}}{(K_M + [\text{S}])\left(1 + \frac{[\text{I}]}{K_I}\right)} \quad (8)$$

The equation could be arranged in linear relation (Gonze & Kaufman, 2016):

$$\frac{1}{V} = \frac{1}{[\text{S}]} \left(\frac{K_M \left(1 + \frac{[\text{I}]}{K_I}\right)}{V_{\max}} \right) + \frac{1}{V_{\max}} \left(1 + \frac{[\text{I}]}{K_I}\right) \quad (9)$$

In this case, the mixed inhibition could be expressed through the modifications:

$$K_M' = K_M \left[1 + \frac{[\text{I}]}{K_I}\right] \text{ and } K_M = \frac{1}{K_s}$$

Where K_s is the term describing enzyme dissociation constant (from E.S or E.I).

The new values of K_M after the inhibition were 0.056 mg. mL⁻¹ in [Cho]OAc system and 0.0443 mg. mL⁻¹ in the case of [EMIM]DEP with CMC. The value of K_s was 22.22 min⁻¹, K_s' is 17.86 min⁻¹, and since the concentration of the presumed inhibitor (IL), is known throughout the reaction, K_I was calculated at 0.409 min⁻¹. The catalytic efficiency (k_{cat}/K_M) decreased from 30.71 to 24.46 min⁻¹ mg⁻¹ mL⁻¹ in the presence of [Cho]OAc and increased to 31.86 min⁻¹ mg⁻¹ mL⁻¹ in [EMIM]DEP. The maximum k_{cat}/K_M value of an enzyme is 10⁸-10⁹ M⁻¹s⁻¹, and it is an indicator of the enzyme catalytic efficiency.

It was observed in this study that the activity reduced IL concentration increases (Figure 5), which indicated that the enzyme affinity towards the substrate decreased. Nonetheless, though the [Cho]OAc resulted in slight inhibition, the K_M and V_{\max} , did not change remarkably.

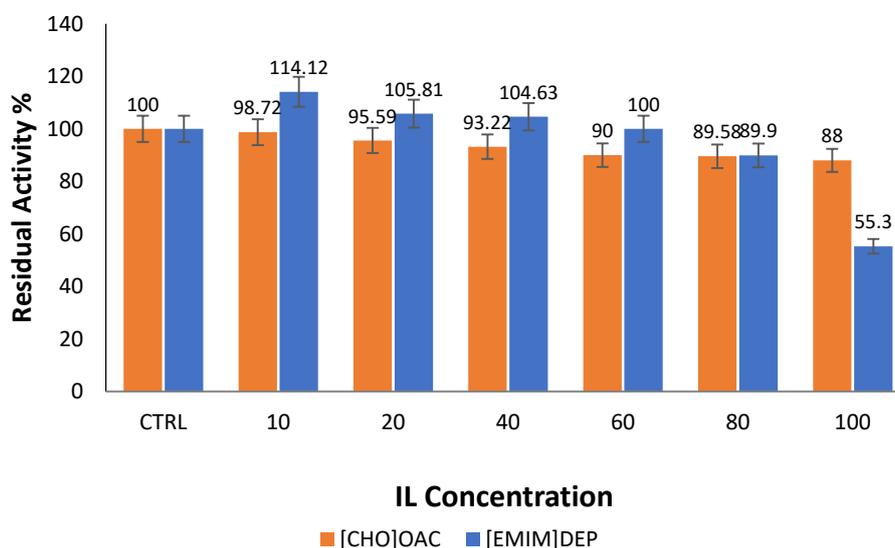


Figure 5. Effect of various concentrations of ILs [EMIM]DEP, [Cho]OAc on the cellulase activity compared to the buffer solution. CTRL= control

The changes were different in the presence of [EMIM]DEP as it slightly increased the V_{max} and did not significantly affect the affinity. It was reported that higher concentration of the substrate could promote surface accessibility and cellulases binding sites, hence, preventing inactivation effect. This could be able to explain the hydrolysis continuation for 48 h and the production of sugars regardless of the IL presence in the hydrolysis vessel (Elgharabawy et al., 2016). The slight impact of IL could also be explained by binding of IL anion to the lignin instead of the enzyme and therefore, the more free enzyme would be available which rescues the inactivation effect. Moreover, cellulase is surrounded by water molecules in the aqueous-IL system which could reduce the inhibition caused by IL or lignin (Zhao, 2016) as illustrated in Figure 6 which also explained the pattern in this study. It can be observed that [Cho]OAc has a slight inhibitory effect if compared to [EMIM]DEP. This could be a result of the composition of the IL. [Cho]OAc consists of cholinium cations and acetate anions, and the former is derived from choline chloride, which is part of the vitamin-B complex, the latter is derived from intercellular metabolites. [Cho]OAc is a completely bioderived IL, which is more biocompatible, compared to imidazolium-based IL, [EMIM]DEP. This might also explain the behavior of the enzyme when water is added in both ILs (Asakawa et al., 2015). Enzymes usually require a certain amount of water in order to function in a particular reaction (Kohno, Saita, Murata, Nakamura, & Ohno, 2011).

It was inferred by Engel et al. that the activity of *T. reesei* cellulase reduced to 15-30% in IL concentration of 10% (Engel et al., 2010). Their group have investigated the [DMIM]DMP effect on individual cellulases and showed that the β -glucosidase, cellobiohydrolase and endoglucanase retained 34%, 60%, and 63% of their relative activities in 10% [DMIM]DMP (Engel, Krull, Seiferheld, & Spiess, 2012). Furthermore, Hu et al. (2016) showed that low concentration of [EMIM]DEP has a reversible inactivation on endoglucanases, and they supported that by SDS-PAGE, however, irreversible effect took place at a high concentration of the IL (more than 40%).

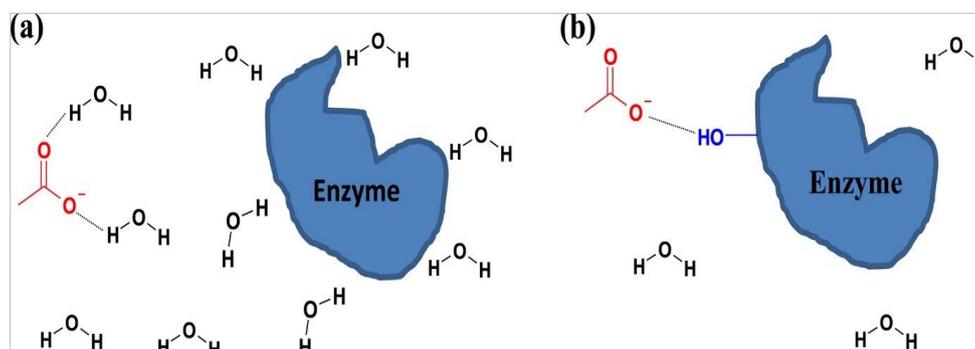


Figure 6. Illustration of interactions of the enzyme in acetate-containing IL: (a) between acetate anion with water molecules in diluted IL solution, and (b) between acetate anion and enzyme molecule in concentrated IL. Adapted from (Zhao, 2016) with permission from Wiley

Some hydrolases, such as certain lipases and chymotrypsin, retained their activity in pure IL or might show better enantioselectivity, thermal stability or selectivity, compared to traditional solvents. Nonetheless, few studies have reported tolerating cellulases that could withstand high concentrations IL (Hu et al., 2016). Furthermore, to date, the activity of commercial cellulases in [EMIM]OAc, [BMIM]Cl, [DMIM]DMP and [AMIM]Cl were reportedly used as pretreatment of cellulose.

The kinetic constants for endoglucanase using different substrates have been documented. V_{\max} values were 18, 46 and 37 $\mu\text{mol}^{-1} \cdot \text{mg protein}^{-1}$ with CMC, β -glucan and xyloglucan, respectively. The value of K_M was 13.3 $\text{mg} \cdot \text{mL}^{-1}$, 3.5 $\text{mg} \cdot \text{mL}^{-1}$ and 5.66 $\text{mg} \cdot \text{mL}^{-1}$ with the same substrates order. The value of k_{cat} was between 5.6 and 11.5 min^{-1} and k_{cat}/K_M was between 42 to 400. Reports have also revealed that endoglucanases vary in their affinity towards polysaccharides as supported by the observed values of K_M (Kaur, Oberoi, & Chadha, 2015). Glucose and [EMIM]OAc acted as uncompetitive inhibitors for β -glucosidase (BG), which in this context indicated the probability of IL binding to the E-S complex. Most of the reported BG(s) are competitively inhibited by glucose; however, a non-competitive inhibition was also reported. Cellulase enzyme from *Bacillus sp.*, MSL2 strain, from rice paddy field soil (48 kDa) retained 77% of its activity in [EMIM]OAc. The kinetic parameters of the purified enzyme revealed that the V_{\max} was 1000 $\mu\text{M} \cdot \text{min}^{-1}$, while K_M value was 0.8 $\text{mg} \cdot \text{mL}^{-1}$ (Sriariyanun, Tantayotai, Yasurin, Pornwongthong, & Cheenkachorn, 2016). The types and concentration of inhibitors released throughout the pretreatment of lignocellulose depend on the composition of biomass, reaction conditions and pretreatment method.

It can be concluded that IL systems with a small concentration of certain ions can contribute to supernatural alteration in enzyme enantioselectivity or activity. Hence, ILs may be precisely adjusted for anticipated applications in biocatalysis. For instance, chloride anions are not suitable for cellulase stability while acetate and phosphate showed good compatibility to a certain level. Selecting the appropriate IL for the enzymatic reaction is one of the most important key factors for IL-enzyme catalyzed reactions.

4. Conclusions

The IL presence in the hydrolysis vessel did not result in irreversible inhibition. Promising activities were recorded in both [EMIM] DEP and choline acetate [Cho]OAc. The kinetic study of the one-step hydrolysis showed that the ILs result in a non-competitive mixed inhibition in which both V_{\max} and K_M and are changed compared to the non-IL reaction. In both cases, the study proved that IL did not result in a severe loss of the activity of cellulase as V_{\max} was not dropped dramatically nor the affinity. The study showed that a small amount of water is necessary in order to prevent the inhibition and activate the cellulase in the reaction medium that contains IL. More studies in terms of molecular simulation and modelling are required to generate a clear understanding of the cellulase behavior in ILs.

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