MODELING AND OPTIMIZATION OF MASHING PROCESS IN BEER PRODUCTION WITH RICE ADJUNCT

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ABSTRACT
Adjuncts like rice, wheat and sorghum are used by beer manufacturers worldwide to reduce the cost of production by replacing malt as a starch source. Rice is the most widely used adjunct in Asian countries. Understanding the enzyme kinetics in mashing process is vitally important to maximize sugar yield at a minimum cost. In this research, a semi-empirical model was developed for the mashing process, based on enzyme kinetic equations and experimental results; and this model was used to optimize the operating conditions when enzymes are not added externally. As predicted by the model, when 30% (w/w) of rice was used as an adjunct the maximum sugar yield can be obtained at 56°C and 6.5 pH, and the optimum temperature for mashing process increases with acidity. Since the acidity of solution increases during the mashing process due to the formation of organic acids, use of an increasing temperature profile is recommended to get the maximum output from the mashing process.

Keywords: mashing, wort, rice adjunct, enzyme kinetics, model, optimization

INTRODUCTION
Beer is one of the most consumed beverages in the global market, with a global production of 1.93 billion hectoliters in 2015 (“Beer production worldwide from 2008 to 2015, by region”),. Two main stages in beer production are mashing and fermentation. Mashing is the conversion of starch from malt or other cereals into simple fermentable sugars via enzymatic reactions (Szwed Łukasz, Tomaszewska-Ciosk, & Błażewicz, 2014). The product from the mashing process is a solution with high sugar content called wort. During fermentation, the fermentable sugars in wort are converted to ethanol. Maximizing the conversion of starch to fermentable sugars is essential to maximize the efficiency of beer production, making mashing is a process of crucial importance. While amylase is the main enzyme required for the mashing process, other enzymes like proteases and peptidases are also involved in mashing (Owuma, 1997). Activity of amylase enzyme depends on a number of factors including temperature, pH and composition of solution, and a maximum starch conversion can be obtained through optimization of these parameters. Two types of amylase are involved in the mashing process; α-amylase, which hydrolyses long chain starch molecules into shorter chains, and β-amylase, which further hydrolyses those short chains into simple sugars. α-amylase is reported to perform optimally around 6.5-7.0 pH and 55°C while β-amylase shows highest activity around 40.5-5.5 pH and 65°C temperature (Blazus et al., 2009; Sundararam & Murthy, 2014).

Barley malt is the main raw material used in beer production, as it contains both starch and the amylase enzyme required for starch hydrolysis. Amylase is contained in the embryo, endosperm and the aleurone layer of the malt grain (Briggs, 1964). In countries where barley is not grown as an agricultural crop, brewing beer with 100% malt is not economically feasible. Therefore, other less expensive grains like rice, wheat and sorghum are added as adjuncts during beer production in commercial breweries, thus reducing the production expenses. (Lloyd, 1986).

The percentage of adjuncts used in industrial breweries ranges from 10 to 25% in Europe and 35 to 45% in USA, while in some African countries it is as high as 50-70% (Ogebeide, 2011). When deciding the amount of adjuncts to be used for brewing, several factors have to be considered including required beer quality, cost of adjunct, starch content in adjunct, available mashing technology, fermentation time and whether enzymes are added externally. An added advantage of some adjuncts is the improvement and stabilization of the flavour of beer. One of the main limitations in adjuncts is the low nitrogen content. As most types of adjuncts have a lower nitrogen content than malt, use of too much adjuncts can lead to a reduced Free Amino Nitrogen (FAN) content in wort.

Materials

Malt used in all these experiments were provided free of charge by the Asia-Pacific Brewery, Sri Lanka, and Sri Lankan white rice (oryza sativa) was...
purchased from the local market. White rice was specifically selected due to high starch content, market availability and low price (Williams et al., 1958). All experiments were conducted using same rice and malt stocks.

**Determination of starch content**

The starch content in malt and rice was determined using Fehling’s method. The starch sample was digested with 20 ml of concentrated H₂SO₄ and neutralized with 0.1 N NaOH. This solution was volumed up to 250 ml and titrated with Fehling’s A and B solutions to determine the sugar content. The starch concentration was calculated in the sample based on the sugar content.

**Mashing process**

The mashing process was based on previous research (Mallawarachchi, Bandara, Dilshan, Gunawardena, & Ariyadasa, 2016). Mass percentages of malt and rice were varied while keeping the total weight of malt and rice constant at 20 g. Initially malt and rice were ground separately for 5s at 14000 rpm using Jaipan IS-4250 grinder. Malt was ground without removing the husk while the husk of rice had been removed beforehand.

The rice was gelatinized by cooking for 10 minutes at 100° C. Gelatinized rice was mixed with malt and 100 ml of water in conical flasks by thorough shaking. pH of the solution was adjusted by adding 0.5 M HCl or 0.5 M NaOH. The solutions were incubated in a water bath at selected temperature for 2 hours for the reactions to take place while stirring the solutions continuously. 1 ml of each sample was extracted using a pipette and diluted 500 fold using serial dilution method. 1 ml of each sample was extracted using a pipette and diluted 500 fold using serial dilution method. 1 ml of each sample was extracted using a pipette and diluted 500 fold using serial dilution method. The rice: malt ratio was kept constant at 30:70 as it gave the highest sugar yield, and the mashing curves were obtained at temperatures of 60° C, 65° C and 70° C and the pH values of 4.0 and 5.5. These temperature and pH ranges were selected to enclose the optimum conditions for both α and β amylase.

**Analysis of results**

Initially a statistical model was developed based on experimental results to select the optimum temperature and composition ranges. The data from these experiments were analyzed using the Surface Fitting Toolbox in Matlab® by Mathworks. The effect of temperature and pH were further analyzed using the semi-empirical model based on Michaelis-Menten kinetics. The rice: malt ratio was kept constant at 30:70 as it gave the highest sugar yield, and the mashing curves were obtained at temperatures of 60° C, 65° C and 70° C and the pH values of 4.0 and 5.5. These temperature and pH ranges were selected to enclose the optimum conditions for both α and β amylase.

**Kinetic Model**

According to Michaelis-Menten enzyme kinetics, an enzymatic reaction occurs in two steps: the reversible bonding of enzyme to the substrate forming enzyme-substrate intermediate, and conversion of enzyme-substrate intermediate into products. The second step is considered as the rate limiting step.

\[
E + S \overset{k_1}{\underset{k_2}{\rightleftharpoons}} ES \overset{k_3}{\rightarrow} E + P
\]

The rate of an enzymatic reaction is expressed by Michaelis-Menten equation.

\[
r = \frac{d[P]}{dt} = \frac{r_{max}}{K_m + [S]},\quad \text{where}\quad r_{max} = k_3[E]_0\quad \text{and}\quad K_m = \frac{k_2}{k_1}\quad \text{(Schuler & Kargi, 2002)}
\]

However, the values of \(r_{max}\) and \(K_m\) are dependent on temperature and pH. α-amylase contains two acidic groups, Asp231 and Glu261 which contributed to catalytic activity. Glu261 acts as a proton donor and protonates the glycosidic oxygen in starch molecules, while Asp231 acts as a nucleophile and attacks the terminal carbon atom. According to literature, at low pH values the nucleophilic group is protonated, thus preventing its catalytic activity, and at high pH values, the proton donor (Glu261) is deprotonated, making it impossible to initiate the hydrolysis of starch (Nielsen, Borchart, & Vriend, 2001). While other amylases such as β-amylase and glucoamylase also can contribute to starch hydrolysis, α-amylase is usually a dominant enzyme present in barley malt. The deprotonation of proton donor and the protonation of nucleophilic group can be expressed by the following equations.

\[
EH \rightarrow E + H^+ \\
EH + H^+ \overset{k_1}{\rightarrow} EH^+
\]

The effect of these two reactions on the reaction rate can be expressed by

\[
K = K_m (1 + \frac{K_a}{[H^+]})
\]

The effect of temperature on enzymatic performance is twofold. As the temperature increases, the kinetic energy of molecules will increase, thus increasing reaction rate. This is expressed by Arrhenius equation

\[
K = Ae^{-\frac{E_a}{RT}}
\]

On the other hand, high temperatures will lead to denaturation of enzymes, thus lowering the reaction rate. Kinetics for the thermal denaturation of enzymes are given by

\[
\frac{d[E]}{dt} = -k_d[E]
\]

Where \(k_d\) also varies with temperature according to Arrhenius equation (Schuler & Kargi, 2002).

Since the same malt stocks are used for all experiments, the initial Amylase concentration is taken as proportional to the mass of malt. \(E \text{m}\) as the solution is stirred continuously during the mashing process, it is considered that all the starch is present either in dissolved form or in suspension, without precipitating. Thus, all starch molecules in the solution are free to react with the enzyme. The substrate concentration remaining after a certain time can be written as a function of initial mass of rice, starch fraction by weight in rice, initial mass of starch, starch fraction by weight in malt, volume of sample and product concentration. Since the mashing process spans over a considerable period of time, the diffusion of substrate molecules is not considered as a rate limiting factor. Therefore, the substrate concentration can be expressed as

\[
[S] = \frac{m_{cm} + m_{cm}k_m}{V} - [P]
\]

This model was simulated using the Simulink tool in Matlab®. The Simulink representation of the model is shown in Figure 1.

**RESULTS AND DISCUSSION**

**Kinetic Model**

According to Michaelis-Menten enzyme kinetics, an enzymatic reaction occurs in two steps: the reversible bonding of enzyme to the substrate forming enzyme-substrate intermediate, and conversion of enzyme-substrate intermediate into products. The second step is considered as the rate limiting step.

\[
E + S \overset{k_1}{\underset{k_2}{\rightleftharpoons}} ES \overset{k_3}{\rightarrow} E + P
\]
impact of temperature and rice content on sugar yield was analyzed using the surface fitting tool in Matlab® as shown in Figure 2. The temperature range for the experiments was selected as 50–70 °C and pH range was selected as 4–7 so as to enclose the optimum temperature and pH ranges given in literature. 95.3% confidence intervals were constructed for each parameter based on the standard deviations in experimental results. Experimentally determined model coefficients at selected temperature and pH values are given in Table 1.

Based on the Arrhenius equation and the experimentally determined model coefficients, activation energies of each reaction were determined. Confidence limits for the activation energies were determined by calculating the activation energies corresponding to the highest and lowest values for the model coefficients. Formation of the product required an activation energy of 17848±627 kJ/kmol and the thermal denaturation needed an activation energy of 111848±5057 kJ/kmol. The activation energies for protonation of Asp231 group

Table 1 Experimentally determined model coefficients with 95.3% confidence intervals

<table>
<thead>
<tr>
<th>Temp</th>
<th>pH</th>
<th>$r_{\text{max}}$</th>
<th>$K_a$</th>
<th>$K_n$</th>
</tr>
</thead>
<tbody>
<tr>
<td>50</td>
<td>4</td>
<td>0.1±0.002</td>
<td>0.003±0.0005</td>
<td>60.5±1.5</td>
</tr>
<tr>
<td>50</td>
<td>5.5</td>
<td>0.1±0.002</td>
<td>0.003±0.0005</td>
<td>16.5±2</td>
</tr>
<tr>
<td>50</td>
<td>7</td>
<td>0.1±0.002</td>
<td>0.003±0.0005</td>
<td>7±1</td>
</tr>
<tr>
<td>60</td>
<td>4</td>
<td>0.125±0.004</td>
<td>0.004±0.001</td>
<td>42.5±2</td>
</tr>
<tr>
<td>60</td>
<td>5.5</td>
<td>0.125±0.004</td>
<td>0.004±0.001</td>
<td>9±2</td>
</tr>
<tr>
<td>60</td>
<td>7</td>
<td>0.125±0.004</td>
<td>0.004±0.001</td>
<td>4±2</td>
</tr>
<tr>
<td>65</td>
<td>4</td>
<td>0.14±0.002</td>
<td>0.006±0.0005</td>
<td>46±1.5</td>
</tr>
<tr>
<td>65</td>
<td>5.5</td>
<td>0.14±0.002</td>
<td>0.006±0.0005</td>
<td>13±0.5</td>
</tr>
<tr>
<td>65</td>
<td>7</td>
<td>0.14±0.002</td>
<td>0.006±0.0005</td>
<td>10±1</td>
</tr>
<tr>
<td>70</td>
<td>4</td>
<td>0.145±0.006</td>
<td>0.013±0.001</td>
<td>35±5</td>
</tr>
<tr>
<td>70</td>
<td>5.5</td>
<td>0.145±0.006</td>
<td>0.013±0.001</td>
<td>13±1.5</td>
</tr>
<tr>
<td>70</td>
<td>7</td>
<td>0.145±0.006</td>
<td>0.013±0.001</td>
<td>17.5±1</td>
</tr>
</tbody>
</table>
in Amylase and deprotonation of Glu 261 group in Amylase were 6796±13715 kJ/kmol and 160770±15992 kJ/kmol respectively. All these activation energies were expressed as 95.3% confidence intervals.

These activation energy values suggested that the rate limiting step was less affected by temperature compared to the protonation and denaturation of amylase. This agreed with the observation of rapid reduction in enzyme activity at higher temperatures as shown in Figure 2, which suggested that the effect of denaturation has overtaken the effect of increase in kinetic energy when the temperature exceeded 65°C. According to literature the activation energy for denaturation is 77.6 kJ/mol for Bacillus subtilis α-amylase, which is a non-thermophilic α-amylase type with an optimum temperature of 40°C, and 316 kJ/mol for Pyrococcus furiosus α-amylase, which is a hyperthermophilic type of amylase (Brown, Dafforn, Fryer, & Cox, 2013; Ludikhuyze, Van den Broeck, Weemaes, Hendrickx, & Tobback, 1997; Raul, Biswas, Mukhopadhyay, Kumar Das, & Gupta, 2014). The experimentally determined activation energy for denaturation of amylase was close to that of Bacillus subtilis α-amylase, which agreed with the rapid reduction in enzyme activity above 65°C.

In order to further analyse the variation of Model coefficients with temperature and pH, the model coefficients at selected conditions were calculated based on expected values of activation energies. Those results are given in Table 2.

### Table 2 Coefficients of the semi-empirical model calculated using activation energies

<table>
<thead>
<tr>
<th>Temp</th>
<th>pH</th>
<th>K_a</th>
<th>r_max</th>
<th>K_m</th>
<th>K_m⁰</th>
<th>K_a⁰</th>
</tr>
</thead>
<tbody>
<tr>
<td>50</td>
<td>4</td>
<td>0.002568</td>
<td>0.1013</td>
<td>1.07E-09</td>
<td>0.00000910</td>
<td>6.983</td>
</tr>
<tr>
<td>50</td>
<td>5.5</td>
<td>0.002568</td>
<td>0.1013</td>
<td>1.07E-09</td>
<td>0.00000910</td>
<td>6.983</td>
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<td>1.07E-09</td>
<td>0.00000910</td>
<td>6.983</td>
</tr>
<tr>
<td>60</td>
<td>4</td>
<td>0.005184</td>
<td>0.1236</td>
<td>6.92E-09</td>
<td>0.00002545</td>
<td>9.457</td>
</tr>
<tr>
<td>60</td>
<td>5.5</td>
<td>0.005184</td>
<td>0.1236</td>
<td>6.92E-09</td>
<td>0.00002545</td>
<td>9.457</td>
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<td>60</td>
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<td>0.005184</td>
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<td>6.92E-09</td>
<td>0.00002545</td>
<td>9.457</td>
</tr>
<tr>
<td>65</td>
<td>4</td>
<td>0.007252</td>
<td>0.136</td>
<td>1.685E-08</td>
<td>0.00003658</td>
<td>10.973</td>
</tr>
<tr>
<td>65</td>
<td>5.5</td>
<td>0.007252</td>
<td>0.136</td>
<td>1.685E-08</td>
<td>0.00003658</td>
<td>10.973</td>
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<td>65</td>
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<td>0.00003658</td>
<td>10.973</td>
</tr>
<tr>
<td>70</td>
<td>4</td>
<td>0.01217</td>
<td>0.1469</td>
<td>0.00000004</td>
<td>0.00005203</td>
<td>12.677</td>
</tr>
<tr>
<td>70</td>
<td>5.5</td>
<td>0.01217</td>
<td>0.1469</td>
<td>0.00000004</td>
<td>0.00005203</td>
<td>12.677</td>
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<tr>
<td>70</td>
<td>7</td>
<td>0.01217</td>
<td>0.1469</td>
<td>0.00000004</td>
<td>0.00005203</td>
<td>12.677</td>
</tr>
</tbody>
</table>

Comparing Table 2 and Table 1, the model coefficients calculated using expected values of activation energies lied within the 95.3% confidence intervals of experimentally determined coefficients. This suggested that the calculated activation energies could be used to predict mashing behaviour accurately.

In previous researches the Michaelis-Menten coefficients for Termamyl α-amylase and Liquozyme α-amylase, which are used in commercial breweries to catalyze the mashing process, have been analyzed at different conditions. At 65°C and 5.5 pH, Michaelis Menten coefficient varies within the range of 6.13-10.85 for Termamyl α-amylase and 6.03-11.09 for Liquozyme α-amylase (Presecki, Blazevic, & Vasic, 2013). According to the semi-empirical model, the value of K_m at similar conditions was 11.980, which was very close to the literature values. These results suggest that the amylase enzyme naturally present in malt shows similar Michaelis Menten coefficients as commercial enzymes.

In order to evaluate the ability of the model to predict mashing behaviour, mashing curves at different conditions were predicted based on expected values of model coefficients and the predicted mashing curves were compared with actual mashing curves. Comparison between predicted and actual mashing curves is shown in Figures 3a-c.
The experimental mashing curves showed that the sugar yield increased with the mashing time. When the mashing time reached 2 hours, the rate of sugar production became very low, which could be explained by the denaturation of enzymes. Considering the effect of temperature, the sugar yield at 50° C and 70° C was considerably low compared to other temperatures at any pH. At 4 pH, the lowest sugar yield was observed at 50° C and at 5.5 pH and 7 pH. Lowest sugar yield is observed at 70° C. pH values of 5.5 and 7 had yielded high sugar concentrations compared to 4 pH.

As shown in Fig. 4, the model had predicted the mashing behaviour fairly accurately at conditions considered in this work except at 50° C 5.5 pH, 65° C 7 pH and 70° C 7 pH where there was slight deviation from the experimental mashing curves. This deviation may have resulted from either and experimental error or a factor such as product or substrate inhibition, which was not considered in this model.

These mashing curves justify the difference between experimentally determined activation energies for protonation and deprotonation of amylase. The activation energy of deprotonation of amylase is much higher than the activation energy for protonation. Thus, the deprotonation of Glu261, which inhibits the reaction at high pH values, is more likely to happen at higher temperatures compared to the protonation of Asp231. This behaviour could be seen in Fig. 4, where at 70° C, the sugar yield at 5.5 pH and 7 pH had decreased more drastically compared to that at 4 pH.

The accuracy of the models at selected conditions were evaluated based on SSE, R² and Average absolute percentage error. Those results are shown in Table 3.

### Table 3: Correlation between actual results and predicted results

<table>
<thead>
<tr>
<th>Temp/PH</th>
<th>SSE</th>
<th>Avg Error (%)</th>
<th>R²</th>
</tr>
</thead>
<tbody>
<tr>
<td>50° C 4.0 pH</td>
<td>16.225</td>
<td>5.08</td>
<td>0.9971</td>
</tr>
<tr>
<td>50° C 5.5 pH</td>
<td>210.267</td>
<td>9.98</td>
<td>0.9989</td>
</tr>
<tr>
<td>50° C 7.0 pH</td>
<td>12.954</td>
<td>1.64</td>
<td>0.9998</td>
</tr>
<tr>
<td>60° C 4.0 pH</td>
<td>181.517</td>
<td>9.20</td>
<td>0.9993</td>
</tr>
<tr>
<td>60° C 5.5 pH</td>
<td>125.804</td>
<td>5.52</td>
<td>0.9991</td>
</tr>
<tr>
<td>60° C 7.0 pH</td>
<td>300.261</td>
<td>8.78</td>
<td>0.9972</td>
</tr>
<tr>
<td>65° C 4.0 pH</td>
<td>82.183</td>
<td>5.42</td>
<td>0.9908</td>
</tr>
<tr>
<td>65° C 5.5 pH</td>
<td>226.717</td>
<td>4.84</td>
<td>0.9968</td>
</tr>
<tr>
<td>65° C 7.0 pH</td>
<td>343.989</td>
<td>6.94</td>
<td>0.9987</td>
</tr>
<tr>
<td>70° C 4.0 pH</td>
<td>16.540</td>
<td>3.24</td>
<td>0.9980</td>
</tr>
<tr>
<td>70° C 5.5 pH</td>
<td>97.496</td>
<td>4.55</td>
<td>0.9848</td>
</tr>
<tr>
<td>70° C 7.0 pH</td>
<td>223.569</td>
<td>8.93</td>
<td>0.9690</td>
</tr>
</tbody>
</table>

According to Table 3 at all conditions, the average error of this model was less than 10%, and the R² has been greater than 0.95, which showed that this is a fairly accurate model to predict mashing behaviour.

### Prediction of optimum mashing conditions

This model was used to predict the sugar yield after 2 hours at different temperatures and pH values. The results predicted by the model are given in Table 4 and graphically interpreted in Figure 4. According to the predicted results, the maximum sugar yield can be obtained at 54-56 °C and 6.5 pH. This is slightly different from the optimum conditions of 50° C and 5.5 pH for amylase in barley malt as mentioned in literature (Greenwood & MacGregor, 1965). However, as shown in Fig. 5, the optimum temperature is heavily dependent on pH value, and the optimum temperature is high at acidic pH values. When the pH value reaches 4, the optimum temperature reaches 66° C.

### Table 4: Predictions for sugar yield by semi-empirical model

<table>
<thead>
<tr>
<th>Temp</th>
<th>PH</th>
<th>4</th>
<th>4.5</th>
<th>5</th>
<th>5.5</th>
<th>6</th>
<th>6.5</th>
<th>7</th>
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<tr>
<td>50</td>
<td>75.871</td>
<td>101.378</td>
<td>115.920</td>
<td>122.190</td>
<td>124.429</td>
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<td>52</td>
<td>81.896</td>
<td>105.623</td>
<td>118.299</td>
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<td>115.300</td>
<td>115.696</td>
<td>115.048</td>
<td>112.528</td>
<td></td>
</tr>
<tr>
<td>70</td>
<td>88.704</td>
<td>100.636</td>
<td>105.5491</td>
<td>107.17672</td>
<td>107.371</td>
<td>106.377</td>
<td>102.988</td>
<td></td>
</tr>
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</table>
These results justify the use of temperature profiles used in the industry. During the mashing process a number of organic acids including mono-, di- and tri hydroxyoctadecenoic acids are formed, causing a reduction in pH (Kobayashi, Kaneda, Kano, & Koshino, 1993; Kobayashi et al., 2000). As the optimum temperature is higher at acidic conditions, increasing the operating temperature with time is recommended to get maximum sugar yield. While the exact temperature profile used for brewing depends on a variety of factors including type and source of amylase used and quantity of adjuncts, most industries use temperature profiles in the range of 35-75° C. According to literature, an optimized mashing profile consisting of 15 min at 35° C, 15 min at 45° C, 40 min at 65° C, 30 min at 72° C, 10 min at 78° C has yielded more fermentable sugars than constant temperature mashing (Wijngaard & Arendt, 2006). Therefore, the results of this experiment agree with the use of increasing temperature profiles used in brewing industry.

CONCLUSION

According to the semi-empirical model of the mashing process, sugar yield can be maximized at 56° C and 6.5 pH without external enzyme addition, and the optimum temperature increases as the operating pH decreases. Since the pH is not kept constant during industrial mashing process and the wort becomes more acidic during the mashing process due to the formation of organic acids, use of a temperature profile where the temperature is gradually increased within the range of 50-70° C in industrial fermentation can be justified. Approaching the effect of temperature and pH on the mashing process through these results will be useful for Asian beer manufacturers in order to maximize sugar yield using the adjunct rice which is the main agricultural crop in Asian countries. Besides, since this model is based on a well established theory, this can be easily adjusted to predict the mashing behaviour for different raw materials and different sources of amylase. Possibilities of further research includes incorporating complex factors affecting the mashing process, including aggregation of enzymes, diffusion of substrate and enzyme and substrate and product inhibition.

Nomenclature

A – Arrhenius constant
E – Activation energy, kJ/mol
(E) – Enzyme concentration, g/l
(Eb) – Initial enzyme concentration, g/l
Kd – Rate constant for the bonding of enzyme to the substrate
Kk – Rate constant for the detachment of enzyme from substrate
Kc – Rate constant for conversion of enzyme-substrate complex into product
Kb – Rate constant for bonding of Amylase with H+ ions
Km – Thermal Denaturation coefficient, \( s^1 \)
Km – Michaelis-Menten coefficient
m – Mass of malt, g
m – Mass of rice, g
(P) – Product concentration, g/l
r – Rate of glucose production, g/l.s

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