Purification and recovery of invertase from potato tubers (*Solanum tuberosum*) by three phase partitioning and determination of kinetic properties of purified enzyme

[Üçlü faz ayırma yöntemi ile patates (*Solanum tuberosum*) yumrularından invertazın saflaştırılması ve geri kazanımı ve saflaştırılmış enzimin kinetik özelliklerinin belirlenmesi]

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ABSTRACT

Objective: Three phase partitioning (TPP) is a simple and economical purification and recovery technique for proteins and also enzymes. In this technique; proteins are separated in the interfacial or lower phase while contaminant parts of crude extract partitioning occur in the upper organic phase. In this study invertase was purified and recovered from potato by TPP.

Methods: The highest enzyme recovery of 121% and purification fold 5.67 was seen in the aqueous phase at the 20% ammonium sulfate saturation with 1.0: 1.0 t-butanol ratio at the first cycle of partitioning. The TPP was optimized by adding ammonium sulfate to final concentration at 25% with the constant t-butanol ratio to starting material.

Results: Purification fold and activity recovery were found as 12.8 and 156%, respectively. The Km value, optimal temperature and optimal pH of purified enzyme were also determined.

Conclusion: The present study showed that, TPP can be an attractive and effective technique for the extraction of invertase from potato.

Key Words: TPP, invertase, recovery, purification, potato tuber.

Conflict of Interest: The authors declare no conflict of interest.

ÖZET


Metod: Üçlü faz ayırımı iki döngüde gerçekleştirilmiştir. birinci döngüde enzim %121 geri kazanım ve 5.67 saflaştırma katsayısı ile %20 amonyum sülfat doygunluğunda ve 1.0: 1.0 t-butanol: ham enzim oranı ile sulu fazda görülmuştur. İkinci döngüde ise amonyum sülfat oranı %25’e çıkartılıp t-butanol ve ham enzim oranı sabit bırakılarak optimize edilmiştir.

Bulgular: Saflaştırma katsayısı ve enzim geri kazanımı sırasıyla 12.8 ve %156 olarak belirlenmiştir. Ayrıca enzimin Km değeri optimal sıcaklığı, sıcaklık stabilitesi, optimal pH’si ve pH stabilitesi belirlenmiştir.

Sonuç: Çalışma Üçlü faz ayırma yönteminin invertazın patates yumrularından alınmasını için etkili ve elverişli yöntem olduğunu göstermektedir.

Anahtar Kelimeler: Üçlü faz ayırımı, invertaz, geri kazanım, saflaştırma, patates.

Çıkar Çatışması: Yazarların çıkar çatışması yoktur.
Introduction

Invertase (β-fructofuranosidase E.C. 3.2.1.26) is a glycoenzyme that hydrolyzes the terminal non-reducing β-fructofuranoside residues in β-D-fructofuranose, like sucrose [1]. Invertase is found in many plants and it is a key enzyme for the carbohydrate mobilization as sucrose is catalyzed irreversibly by invertase into glucose and fructose residues which are main carbon source in plant metabolism [2-5]. This enzyme has many applications in cosmetics, pharmaceutical, paper but especially food and beverage industries [6-8]. Invertase was firstly isolated from yeast more than a century ago [3]. From the date until now researchers reported several purification methods that involve salt precipitation, different chromatographic techniques and ultrafiltration etc. which are led to low yields. They were also un scalable, time consuming, expensive multistep protocols [2-5,9-12]. Three phase partitioning has relatively recent use for bioseperation of enzymes and inhibitors [13-17]. It uses ammonium sulfate with certain saturation to precipitate the protein and t-butanol is added to make three phase layers and to remove lipids, phenolic compounds and some detergents [13,18]. The addition of t-butanol in the presence of ammonium sulfate pushes the protein out of the solution [13]. It is believed that the power of the separation is due to collective operation of conventional salting out, isoionic-osmolytic precipitation and hydration shifts of proteins [19]. There is no report about using TPP to separate invertase from potato; therefore this work was performed for purification and recovery of invertase by TPP. This method has an important potential use in various industries.

Material and Methods

Materials

Sucrose and t-butanol were pure grade and purchased from Merck (Darmstadt, Germany), ammonium sulfate and Coomassie Brilliant Blue R-250 were purchased from Sigma Chem. Co. (St. Louis, MO, USA), potato (Solanum tuberosum) was purchased from Kocaeli city, Akmeşe region, Turkey. All other chemicals and reagents were pure grade.

Preparation of crude extract

Fresh potato roots were washed thoroughly with distilled water and then the peels were removed. Twenty gram roots were than cut into small pieces and then blended in 40 mL, 50 mM pH 5 acetate buffer with, containing 0.8 M sodium sulfate and 0.1 mM EDTA at +4 °C for a minute. The homogenate was then filtered from five layers of cheese cloth, then centrifuged at 10000 rpm for 20 min at +4 °C [5,21]. The clear supernatant was treated as the crude enzyme extract for the further study (protein amount and specific activity was determined as 1.59 mg/mL and 0.57 U/mg, respectively).

Determination of invertase activity

800 µL substrate (50 mM sucrose in 50 mM pH 4.7 acetate buffer) and 200 µL suitably diluted enzyme solution were taken in a test tube (extinction coefficient of o-sucrose equals to 2.568 x 10^3 M^-1 cm^-1). The reaction was incubated at 37 °C for 30 min then was stopped by adding 0.1 ml of DNS (3,5-Dinitrosalicylic acid) reagent. The mixture was heated in a boiling water bath for 5 min. It was then cooled in ice bath and the amount of reducing sugars was measured spectrophotometrically at 540 nm [22]. The data presented for all invertase activity determinations are mean values of duplicate assay. One unit of invertase activity was defined as the amount of enzyme which released 1 µmole of glucose from sucrose per minute at pH 4.7 and 37 °C.

Protein concentration

Concentration of the protein was determined by Bradford method [23] using Coomassie brilliant blue G-250 dye as a reagent and bovine serum albumin (BSA) as standard, by measuring the absorbance at 595nm at 25 °C. Assays were performed in duplicate and the averages were used in calculations.

Three-phase partitioning

Effect of t-butanol

TPP experiments were carried out by employing various t-butanol ratios (crude extract:t-butanol; 1.0:0.5, 1.0:1.0, 1.0:1.5, 1.0:2.0) with a constant ammonium sulfate saturation at 20% (w/v). The mixture was mixed gently and then allowed to stand for 30 min at 37 °C. Afterwards the mixture was centrifuged at 5000 rpm for 10 min at +4 °C to facilitate the separation of phases. The lower aqueous phase and the interfacial phase were collected and dialyzed against 50 mM phosphate buffer overnight at +4 °C. Each of phases was analyzed for enzyme activity and protein content. The experimental conditions at which the highest enzyme activity observed were selected for the further experiments.

Effect of ammonium sulfate saturation

The effect of different ammonium sulfate saturations (20, 30, 40, 50, 60, 70%) (w/v) were also investigated at the best recovery activity crude enzyme:t-butanol ratio is obtained from section 2.3.1. The bottom phase and interfacial phase were collected and dialyzed against 50 mM phosphate buffer overnight at +4 °C. Each of phases was analyzed for enzyme activity and protein content. The highest enzyme recovery was chosen for the further experiments.

Optimization of TPP conditions

For the second cycle of TPP, the highest activity recovery of aqueous phase from the first cycle was used for optimization of TPP conditions. Selected phase wasn’t
dialyzed [25,26]. t-butanol in ratio of 1.0:1.0 was added, than saturated with the different ammonium sulfate to final concentrations of 25, 30, 35, 40, 45, 50%. The enzyme recovery and protein content of phases were analyzed as previously mentioned.

**Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE)**

Molecular weight of purified invertase was determined by SDS-PAGE according to the method of Laemmli [24] on a Biorad Mini Protein electrophoresis unit. Electrophoresis was carried out at 100 mV for 120 min in citrate buffer (pH4.7) and 12% polyacrylamide gel system. For each lane 20µg protein was applied. The gel was stained with Coomassie Brilliant Blue R-250 for 1 hour then destained by 40% methanol and 10% acetic acid for 2 h.

**Kinetic studies**

Purified and recovered enzyme preparation was used for kinetic studies. The pH stability of enzyme was determined by incubation of enzyme for 1 h at 37 °C at various pH, using 25 mM concentrations of the following buffer systems: glycine-HCl (pH 2.0, 3.0), sodium acetate (pH 4.0, 5.0, 6.0), sodium phosphate (pH 7.0, 8.0) and glycine-NaOH (pH 9.0, 10.0), followed by assay of the remaining activity at pH 4.5 as described section 2.2.2. The relative activities (as percentage) were expressed as the ratio of the invertase activity obtained at a certain pH to the maximum activity obtained in the given pH range. The optimal pH for activity was determined by measuring the activity of enzyme in 50 mM sucrose with the same buffers at various pH at 37°C. In order to define the optimal temperature of enzyme against sucrose, different temperatures ranging from 25 to 60 °C (25, 30, 35, 40, 45, 50, 55, 60) were assayed. After incubation periods, enzyme was assayed at standard activity conditions for determining the relative activity. The relative activity (as percentage) was expressed as the ratio of the invertase activity obtained at a certain temperature to the maximum activity obtained in the given temperature range. Michaelis-Menten constant (K_m) of enzyme was determined at the presence of varying concentrations of sucrose, (1, 2, 3, 4 and 5 mM, respectively). The data presented for all invertase activity determinations are mean values of duplicate assay.

**Result and Discussion**

**Three phase partitioning**

**First cycle of TPP**

TPP is a simple technique in which protein rich layer is interfacial phase and most of the activity of enzyme is often seen in this phase. Pigments, lipids and enzyme inhibitors are concentrated in the upper phase, proteins are precipitated in intermediate phase as a salt phase while polar components are enriched in the aqueous phase [27]. However in some cases the target protein is recovered from aqueous phase [28]. This reason requires the second cycle TPP with the use of first cycle TPP’s aqueous phase as the starting material [29]. Fig. 1 shows the enzyme recovery and the protein content of aqueous phase at the different t-butanol ratios for the constant 20% (w/v) ammonium sulfate saturation. It is observed that t-butanol gave maximum recovery of invertase in 1.0:1.0 ratios with maximum fold purification and recovery 5.67 and 121%, respectively. Interestingly, an increase of t-butanol ratio caused a fully decrease of purification and activity recovery of invertase in the aqueous phase. If the t-butanol ratio is high, the denaturation of protein is more possible [18]. This may be attributed to synergetic effects of the increase in concentration of t-butanol and decrease in saturation of ammonium sulfate [25,26]. Figure 2a depicts the best saturation of ammonium sulfate concentration for the recovery and purification fold of invertase in aqueous phase for the first cycle of TPP. Generally one starts with a minimum salt saturation of 20% (w/v) and optimizes this so as to obtain maximum purification fold and recovery [15,16,25]. In this study invertase recovery of 121% yield and 5.67 folds purification were obtained in the aqueous phase at 20% saturation of ammonium sulfate with the 1.0:1.0 crude enzyme:t-butanol ratio. For all the ammonium sulfate saturations there was no enzyme activity in the interfacial phase. This finding was led to work completely in aqueous phase to second cycle of TPP. Increasing concentrations of ammonium sulfate resulted in decreasing of invertase recovery in aqueous phase. The high concentration of salt might decrease the selectivity of partitioning thus reduce recovery of enzyme [26].

As a result, obtained 1.0:1.0 t-butanol ratio and the 20% (w/v) ammonium sulfate saturation seemed to be optimum condition for invertase partitioning in aqueous phase due to highest invertase recovery for the first recycling of TPP. However this fraction gave lower purification fold than the higher salt concentrations. For increasing the purifi-
cation fold of enzyme by this method, aqueous phase of first cycle was optimized by adding further t-butanol and ammonium sulfate.

Second cycle of TPP

For optimizing partitioning conditions, the lower phase of 1.0:1.0 t-butanol ratios and the 20% (w/v) ammonium sulfate saturation of TPP were used as starting material without dialysis. Ammonium sulfate was added to starting material until reaching final saturation of 25, 30, 35, 40, 45, and 50% (w/v). As seen in Figure 1a, 1.0:0.5 t-butanol concentration was not enough for enzyme recovery and purification in aqueous phase and the high concentrations (bigger than 1.0) resulted with completely loss enzymatic activity. Thus, t-butanol concentration was fixed to 1.0:1.0 ratios and different salt concentrations were studied. As shown in Fig. 2b, purification fold was increased as well as recovery activity at the final saturation at 25% of ammonium sulfate in aqueous phase again with 12.70 and 156%, respectively. Increasing salt concentration caused decreasing of purification fold and recovery enzyme. Although protein content was high, enzyme activity was detected as zero in the interfacial phase of the second cycle of TPP as the aqueous phase of first cycle of partitioning. This result showed us that enzyme was completely in aqueous phase in second cycle too. Özer et al. reported [20] that 50% (w/v) ammonium sulfate saturation with 1.0:1.0 ratio of t-butanol gave 8.6 fold purification and 190 % activity recovery in a single step. Sometimes the first cycle of TPP couldn’t remove the contaminant proteins efficiently. This situation was also observed in this study. Chaiwut et al. [18] reported highest enzymatic activity and proteolytic activity (253.5%) of protease in the bottom phase. Wang et al. [28] and Saxena et al. [29] reported purification of α-amylase inhibitor and amylase/protease inhibitor (81 and 80%, respectively) by TPP. They also found the most of the target protein content in aqueous phase and optimized this phase by adding of further t-butanol or salt concentrations.

Overall purification

Purification profile is given in Table 1 and result of SDS-PAGE gel electrophoresis is shown in Figure 3. As seen in Figure 3. SDS-PAGE of crude extract has major bands as 60, 28 and 18 kDa, respectively. It can be observed that...
these bands have been disappeared (lane 1) after TPP. The result indicated that aqueous phase of second cycle of TPP includes a large amount, perhaps only, invertase. In this study molecular weight of invertase was nearly estimated to be 60 kDa which agreed well with the literature [30]. From the Table 1 it is obtained that recovery of enzyme was improved by this technique. Activity recovery of enzyme during the TPP increases due to structural changes and gains a high flexibility which can lead to higher catalytically activity [19,31,32].

**Determination of optimum pH and pH stability**

Figure 4 shows that the optimum pH for potato invertase was detected at 4.5. Optimum pH for the same enzyme was detected as 4.6 by the Matsushita and Uritani [5]. For the bamboo invertase optimum pH was reported as 4.5 [2], and for barley invertase 5.0 [4]. When the enzyme was incubated in pH 2-9 buffer solution at 37°C for 30 min, the enzyme showed good stability in the pH range from 4.0 to 6.0 and nearly 60% of the activity remained at pH 7.0 (Fig. 4).

Invertase activity was assayed at 25-60 °C for the definition of the optimum temperature of purified enzyme and the highest activity was observed as 37°C (Fig. 5). On the other hand, the thermostability profile shows that the enzyme was stable at a range from 30-55°C and more than 70% of activity was remained at these temperatures (Fig. 5). Similar behavior for the invertase from the different sources was reported in the literature [2,20]. When initial enzyme activity plotted versus substrate concentration, it is observed that a curve which became asymptotic to a horizontal line that represents “maximum activity”, often referred to as Vm. Km is defined as that concentration of substrate that gives “half-maximal activity”. However, presentation of kinetic data this plot is not especially useful. A frequently used alternative presentation is the double reciprocal or Lineweaver-Burk plot. In double reciprocal plots the y-intercept represents 1/Vm while the x-intercept represents 1/Km [33].

**Characterization of purified enzyme**

**Determination of optimum temperature and thermal stability**

Km value of purified enzyme was determined by the Lineweaver-Burk reciprocal plot (Fig. 6).

As seen in Figure 6 this value was calculated as 3.5 mM. Km value for potato invertase was reported as 4.0 and 4.5 [5,21]. But in our study Km value of potato invertase was calculated as 3.5 mM. Decreasing of Km value of enzyme may be due to purification technique. In the literature, similar result was reported. X-ray studies of proteinase K showed that, increased flexibility of protein caused increased access to active site of enzyme and this caused the increased the substrate affinity of enzyme. Increasing of substrate affinity caused the decreasing of Km value [34].

**Conclusion**

The novelty of this work is that, TPP was employed firstly for purification and recovery of invertase from potato tubers. This technique is useful for purification of this enzyme. With an efficient and cheap technique, enzyme was purified with 156% and purification fold 12.8 was seen in the aqueous phase at the 25% ammonium sulfate saturat-

<table>
<thead>
<tr>
<th>Step</th>
<th>Total Activity (U)</th>
<th>Total Protein (mg)</th>
<th>Specific Activity (U/mg)</th>
<th>Purification fold</th>
<th>Recovery (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Crude extract</td>
<td>1.82</td>
<td>3.19</td>
<td>0.57</td>
<td>1</td>
<td>100</td>
</tr>
<tr>
<td>1st TPP (lower phase)</td>
<td>2.20</td>
<td>0.68</td>
<td>3.23</td>
<td>5.67</td>
<td>121</td>
</tr>
<tr>
<td>2nd TPP (lower phase)</td>
<td>2.84</td>
<td>0.39</td>
<td>7.28</td>
<td>12.8</td>
<td>156</td>
</tr>
</tbody>
</table>

**Figure 5.** Optimal temperature and thermal stability of purified invertase.

**Figure 6.** Lineweaver-Burk Diagram of invertase at the different sucrose concentrations.
There are no conflicts of interest among the authors.

Conflict of Interest

with project number 2012/057.

Acknowledgements

The present work was carried out with the financial support of Scientific Research Unit of Kocaeli University with project number 2012/057.

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