

Extraction of Amylase from Fermentation Broth in Poly (Ethylene Glycol) Salt Aqueous Two-Phase System

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ABSTRACT

Studies were carried out on the partition of amylase from *Bacillus subtilis* in a minimal medium at 37 °C and 110 rpm. Enzyme recovery was carried out in aqueous two-phase system PEG-Phosphate salt were carried out. The best purification factor (5.4) was obtained in system PEG 1000 (16.7% w/w) with potassium phosphate (14.8% w/w), at pH 6.0, resulting in a recovery of 45.2% activity enzymatic in the salt-rich phase.

Key words: Amylases, aqueous two-phase systems, *Bacillus subtilis*, partition, poly (ethylene glycol).

INTRODUCTION

Amylases are extracellular enzymes of great industrial interest. They are involved in the hydrolysis of starch molecules (Paquet et al., 1991), decomposing them in glucose. Starch is a major source of carbohydrates in the nature, being an abundant carbon source. Every starch polysaccharide molecule possesses one reducing terminal, able to reduce dinitrosalicylic acid; therefore, measurement of reducing sugars is an useful method to determine the molar concentration of starch molecules in solution. Amylases possess important applications in the production of syrup with high glucose content,

sweetener manufacture, nourishing, detergents and ethanol. Other applications include textile and paper industries. Amongst amylolytic enzymes, α -amylases, β -amylases, glucoamylase, pullulanase and isoamylase can be found, being the former two of bigger economic interest (Pandey et al., 2000). There are several sources for the production of amylases, such as bacteria, fungi and plants. *Bacillus* spp. is considered the most important source of amylase. Bacterial α -amylase, however, is preferred over fungal amylase due to several characteristic advantages that it offers. The β -amylase has generally been obtained from plant sources. Filamentous fungi apparently constitute

the major source of glucoamylase among all microbes (Pandey et al., 2000).

In order to perform large-scale purification of enzymes and proteins an efficient downstream processing technique amenable to preserve biological activity is needed (Tanuja et al. 1997).

Extraction in aqueous two-phase system (ATPS) is a suitable method for the first step of a separation procedure removing some contaminants in a simple and economic process (Chaves et al. 2000; Hatti-Kual 1996; Kula et al. 1982). Aqueous two-phase systems are based on water-soluble polymers and salts and/or two different water-soluble polymers. Phase separation occurs over certain concentrations of the phase components. The binodial line is the boundary condition of concentrations for phase separation. Above the binodial line, the phase component separates to the upper phase and the lower phases. However, below the binodial line, one mixed phase exists (Albertsson 1986). The overall composition of phase-forming components has been exploited to influence the phase preference of proteins (Kula et al. 1982). Two-phase systems made by organic solvent and aqueous solution are being used for separation. Two-phase systems sometimes cannot be applied to the separation of active biological materials, which require mild aqueous environments. On the others hand, ATPS have low osmotic pressure, high water, buffering effects added salts and very low interfacial tension between the upper and the lower phases (Albertsson 1986). Aqueous two-phase systems have been used for separating plant and animal cells, cell organelles, membranes, enzymes, nucleotides and other biological materials (Albertsson 1986; Kohler et al. 1993). The aqueous two-phase systems have been designed for scale up of downstream processes for biomaterial separation (King 1992; Mistry et al. 1993). Partition characteristics depend on surface hydrophobicity; specific-binding sites on the surface properties of the biological materials which will be separated (Albertsson et al. 1970; Walter 1977). This separation procedure, i.e. partition, is influenced by phase system parameters, such as the molecular weight of the polymer, pH, type and concentration of phase forming components (Chaves et al. 2000). The PEG/salt system is suitable for downstream processing of amylase, due to its high capacity biocompatibility and low costs. The aim of this work was to study the recovery of amylase

through ATPS, properties such as pH and temperature stability of the crude and partially purified extract determined.

MATERIALS AND METHODS

Reagents

PEG (1000, 3350 and 8000) was obtained from Sigma (St. Louis, MO, USA). All other reagents are of analytical grade.

Microorganism

Bacillus subtilis DA 15 was obtained from the Department of Antibiotics-UFPE.

Amylase production

Bacillus subtilis was grown in fermentation medium described by Haddaoui et al. (1999) which contained the following components: minimal medium supplemented with 1% (w/v) glucose and 0.1mM CaCl₂ in a 110 rpm rotary shaker at 37°C, during six hours. The fermentation broth was centrifuged for 15 minutes at 12500 x. The supernatant was used as enzyme source.

α-Amylase activity assay

The activity of extracellular amylase was estimated by determining the amount of reducing sugars released from starch. The sugars were quantified by the method of 3,5-dinitrosalicylic acid (DNSA), according to Miller (1959). The starch solution was prepared from 1% (w/v) soluble starch in 0.1M citrate/phosphate buffer pH 6.6, at 37°C. Volumes of 0.25 ml of the enzymatic extract were added to 2.5 ml of the starch solution and the mixture incubated at 37°C in a water bath for 10 min. After 1 ml of mixture was added to 2 ml of DNSA and boiled at 100°C for 10 minutes. The amount of reducing sugars in the final mixture was determined spectrophotometrically at 570 nm. One unit of enzymatic activity is defined as the amount of enzyme that produces 1 μmol/min of glucose. The specific activity is expressed as units *per* milligrams of protein. The calibration curve for DNSA was established with several standard solutions of glucose ranging from 100μg ml⁻¹ to 500μg ml⁻¹

Protein quantification

Protein content was determined using the method described by Bradford (1976) with bovine serum albumin as standard.

Effects of pH and temperature on amylase activity

Amylase activity was measured at different pH and temperature values using starch as substrate. The enzyme activities, before and after purification, were assayed using 1% (w/v) starch in 0.1M citrate/phosphate buffer solution (pH ranging from 4.0 - 7.0). The pH optimum assays, the enzyme were incubated at 25°C, for 10 minutes, in buffers solutions of different pH values. The amylase activity was also assayed at various temperatures (40°-100°C) at pH 6.6.

Preparation of the two-phase systems

The systems (total mass 6g) were prepared from stock solutions of the polymers 50% (w/w) (PEG 1000, 3350 and 8000) in water, and 40% (w/w) salt stock solution as a mixture of potassium phosphate monobasic, potassium phosphate dibasic. All the systems, in duplicate, were prepared in graduated centrifuge tubes. The PEG-phosphate salts systems were assumed to be close the original phosphate stock solution (pH 6.0, 7.0, 8.0). The partition experiments were performed at room temperature ($25 \pm 2^\circ\text{C}$) by mixing the systems with crude extract from fermentation (1 ml). After separation of the phases by centrifugation for $236 \times g$, samples of each phase were taken with a pipette for the quantification protein and activity of enzyme. The study of enzyme partitioning at different tie lines was carried out varying the PEG and salts concentrations as show on the Table 1.

Table 1 - Composition of the two phases system

| Polymer | PEG (% w/w) and salt (% w/w) | | |
|----------|------------------------------|------------|------------|
| | Tie line 1 | Tie line 2 | Tie line 3 |
| PEG 1000 | 16.7/14.8 | 17.7/15.7 | 19.7/17.7 |
| PEG 3350 | 14.0/11.8 | 17.7/15.7 | 19.7/17.7 |
| PEG 8000 | 11.8/9.8 | 14.0/11.8 | - |

Determination of the partition coefficient (K):

The relative distribution of the enzyme in the aqueous two-phase system is characterized by the partition coefficient (K), calculated according to the following equation:

$$K = C_t / C_b$$

Where C_t stands for enzyme activity in the top phase, C_b enzyme activity in the bottom phase.

Determination of the purification factor

The purification factor (PF) is defined as the ratio of the enzymatic specific activity of the crude extract by the enzymatic specific activity after purification.

PF = Enzyme specific activity after purification / Enzyme specific activity of crude extract.

Determination of recovery enzymatic activity

The recovery enzymatic activity is defined as the ratio of the enzymatic activity after purification by enzymatic activity of crude extract multiplied by 100.

REA = Enzymatic activity after purification X 100/ enzymatic activity of crude extract.

RESULTS AND DISCUSSION

Studies of amylase partition in the polyethylene glycol/phosphate system

The development of purification procedure using aqueous two-phase systems involves the variation of several factors until a good result is achieved. The partition coefficient, recovery of enzyme activity and factor of purification were studied to characterize the distribution of amylase between the two phases of the system. The enzyme partition in the ATPS depends on the size of the biomolecule, superficial properties, molecular

load, ionic composition of the phases, polymer concentration and molecular length of the polymer (Harries et al. 1997).

Effect of polymer molecular weight

Fig. 1A, 1B and 1C indicated that the systems studied were slightly influenced by the PEG molecular mass. The partition coefficients showed no regular tendency in all the cases. This could be attributed to different two effects.

One of these effects was the increase in the upper phases hydrophobicity. In fact, as PEG chain length increased there were less hydroxyl groups for the same concentration of the polymer and hence the polymer-richer upper phase increased in hydrophobicity. On the other hand, the chain length increase also caused the increase of the excluded volume, which mean less space available for the protein in the PEG rich phase (Marcos et al. 1999). Similar behavior was observed by Marcos et al. (1999) and Almeida et al. (1998) studying the purification of penicillin acylase from *Escherichia coli* and cutinase from *E. coli* WK-6 recombinant, respectively, using poly (ethylene glycol)-sodium citrate and poly (ethylene glycol)- hydroxypropyl starch.

Effect of the tie line length

The partition coefficient (K) of α -amylase in PEG (1000, 3350 and 8000)/phosphate systems are showed in Fig. 1A, 1B and 1C, respectively. In all systems studied in this work at pH 7.0, with increasing the tie line length, K of enzyme decrease. This effect was caused by the increasing PEG concentration witch cause continuous molecular exclusion of amylase from the top to bottom phases, following continuous re-equilibration of the levels in both phases. The same didn't occur in PEG 8000 system at pH 8.0, where there was no significant difference in the K value. Study carried by Almeida et al. 1998, Furuya et al. 1995 and Chaves et al. 2002 for the extraction of cutinase in the PEG 4000/hydroxypropyl starch, hydrolytic enzyme in PEG-dextran system and extraction of Sm-13 recombinant antigen from *Escherichia coli* in the PEG (1000, 3350 and 8000)/Reppal PES 100 system, respectively, showed that the partition coefficients were significantly influenced by the PEG molecular weight. Ortin et al. (1991), studying the partition of α -lactoalbumin and β -

lactoglobulin in PEG/hydroxypropyl starch systems, showed that the increase of the tie line resulted protein displacement to the bottom phase, thus decreasing the K value. Further studies with bovine serum albumin accomplished by Christian et al. (1998) also showed the decrease of K with the increase of the length tie line in the PEG 3350/arabinogalactan system. Cunha et al. (2003) studied the extraction of recombinant and wild cutinase observing that wild cutinase had a tendency for salt rich phase, similar results was obtained in present work.

Table 2, 3 and 4 show the effects of the tie line length in different PEG-phosphate system.

In the PEG 3350 and 8000/phosphate system at pH 6.0, a decrease of recovery of enzyme activity (REA) and purification factor (PF) was observed with the increase of the tie line length in both phases. Similar was noted for PEG 3350 and 1000/salts systems at pH 7.0 (Table 2 and 3, respectively).

Bottom phases, following continuous re-equilibration of the levels in both phases. The same didn't occur in PEG 8000 system at pH 8.0, where there was no significant difference in the K value. Study carried by Almeida et al. (1998), Furuya et al. (1995) and Chaves et al. (2002) for the extraction of cutinase in the PEG 4000/hydroxypropyl starch, hydrolytic enzyme in PEG-dextran system and extraction of Sm-13 recombinant antigen from *Escherichia coli* in the PEG (1000, 3350 and 8000)/Reppal PES 100 system, respectively, showed that the partition coefficients were significantly influenced by the PEG molecular weight. Ortin et al. (1991), studying the partition of α -lactoalbumin and β -lactoglobulin in PEG/hydroxypropyl starch systems, showed that the increase of the tie line resulted protein displacement to the bottom phase, thus decreasing the K value. Further studies with bovine serum albumin accomplished by Christian et al. (1998) also showed the decrease of K with the increase of the length tie line in the PEG 3350/arabinogalactan system. Cunha et al. (2003) studied the extraction of recombinant and wild cutinase observing that wild cutinase had a tendency for salt rich phase, similar results was obtained in present work.

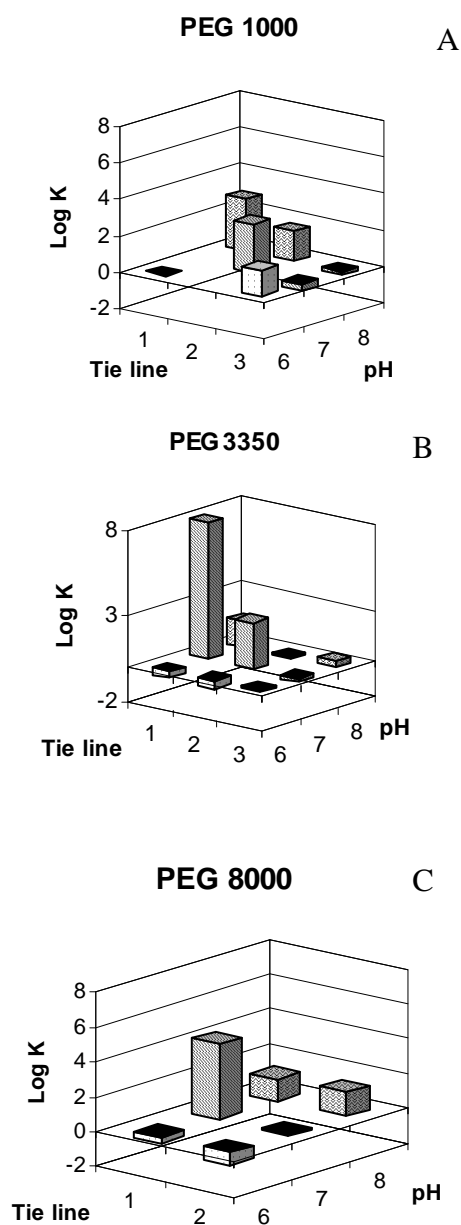


Figure 1 - Effect of tie line length and pH in log K for PEG 1000 (A); PEG 3350 (B); and PEG 8000 (C).

Table 2 - Purification factor and recovery of amylase activity in the PEG 1000/salt systems

| Tie line | | 1 | | | 2 | | | 3 | | |
|----------|--------------|------|---|------|---|------|------|------|------|------|
| | | 6 | 7 | 8 | 6 | 7 | 8 | 6 | 7 | 8 |
| REA (%) | Top phase | 33.8 | * | 22.7 | * | 43.8 | 14.3 | 44.8 | 10.4 | 22.8 |
| | Bottom phase | 45.2 | * | 25.2 | * | 41.0 | 19.4 | 28.3 | 28.6 | 47.9 |
| PF | Top phase | 0.1 | * | 0.2 | * | 0.2 | 0.05 | 0.2 | 0.01 | 0.04 |
| | Bottom phase | 5.4 | * | 0.3 | * | 0.1 | 0.03 | 0.1 | 0.03 | 0.2 |

REA - recovery of enzyme activity; PF-purification factor;

* - Formation of phases was not observed

Table 3 - Purification factor and recovery of amylase activity in the PEG 3350/salt systems

| Tie line | | 1 | | | 2 | | | 3 | | |
|----------|--------------|------|------|------|------|------|------|------|------|------|
| pH | | 6 | 7 | 8 | 6 | 7 | 8 | 6 | 7 | 8 |
| REA (%) | Top phase | 18.7 | 19.1 | 32.0 | 43.0 | 18.5 | 13.0 | 7.8 | 14.0 | 36.3 |
| | Bottom phase | 55.3 | 39.2 | 18.3 | 16.0 | 32.4 | 30.0 | 42.8 | 25.9 | 30.0 |
| PF | Top phase | 1.8 | 0.4 | 0.5 | 1.0 | 0.8 | 0.2 | 0.8 | 0.6 | 1.0 |
| | Bottom phase | 3.8 | 0.5 | 0.7 | 1.9 | 0.3 | 1.3 | 3.5 | 3.0 | 2.8 |

REA- recovery of enzyme activity; PF- purification factor.

Table 4 - Purification factor and recovery of amylase activity in the PEG 8000/salt systems

| Tie line | | 1 | | | 2 | | |
|----------|--------------|------|------|------|------|------|------|
| pH | | 6 | 7 | 8 | 6 | 7 | 8 |
| REA (%) | Top phase | 22.4 | 12.4 | 40.5 | 18.9 | 27.1 | 40.1 |
| | Bottom phase | 46.1 | 64.8 | 41.9 | 30.2 | 23.7 | 26.6 |
| PF | Top phase | 1.1 | 0.3 | 0.8 | 0.3 | 1.3 | 1.8 |
| | Bottom phase | 1.5 | 2.3 | 0.8 | 0.9 | 0.3 | 1.3 |

REA - recovery of enzyme activity; PF- purification factor.

* - Formation of phases was not observed

Table 2, 3 and 4 show the effects of the tie line length in different PEG-phosphate system.

In the PEG 3350 and 8000/phosphate system at pH 6.0, a decrease of recovery of enzyme activity (REA) and purification factor (PF) was observed with the increase of the tie line length in both phases. Similar was noted for PEG 3350 and 1000/salts systems at pH 7.0 (Table 2 and 3, respectively).

Effect of pH

As general rule, negatively charged proteins prefer the upper PEG-rich phase and positively charged proteins partition to the lower phase (Schmidt et al. 1994). As the pH increases above the isoelectric point (pI) of a protein, it becomes negatively charged, its interaction with PEG becomes stronger (Huddleston et al. 1991), and the partition coefficient increases. In the system PEG 1000/ phosphate (Fig. 1A) it was observed that it did not have a regular trend. However, in systems with the biggest molecular weight, an increase of the coefficient of partition occurred with the increase of pH (Figs. 1B and 1C). Same results were obtained by Xu et al. (2002) who studied the extraction of glucose-6-phosphate dehydrogenase and hexokinase in PEG/PES systems. Wu et al. (2001), working with two consecutive aqueous two phase extraction steps in PEG 8000/polyvinyl-alcohol system, followed in PEG 8000/(NH₄)₂SO₄ system, observed strong partitioning of endo-

polygalacturonase into the bottom phase with the increase of pH. The results presented in PEG (3350 and 8000)/ phosphate systems (Table 3 and 4, respectively), tie line 1, demonstrated that PF decreased with increasing pH. Increase of pH resulted a decrease of PF. Similar behavior was obtained in the PEG 1000 and 3350 at tie line 2 (Table 2 and 3). However, except in the PEG 8000 at pH 8.0, tie line 1, no observed a significant difference in the recovery of the enzymatic activity in the bottom phase (41.9%) and in the top phase (40%) were no observed (Table 4). In this system was observed recovery maximum of enzymatic activity (64.8%) occurred in the bottom phase, tie line 1, at pH 7.0. In present work highest purification factor (5.4) was obtained at pH 6.0, tie line 1 in the salt-rich phase of PEG 1000 (Table 2).

Effect of pH and temperature in amylase activity in crude extract and partially purified extract

The profile of pH and temperature over the enzymatic activity of amylase were evaluated before and after purification. Fig. 2A shows the optimum pH of the crude extract before and after purification in the PEG/salt system (PEG 1000, tie line 1, pH 6.0). An optimal pH was obtained at pH 5.0 for both the samples (crude extract and partially purified extract). As shown in Fig. 2B, the optimal temperature was 60°C, in both the samples. In general, these results were in

agreement with studies carried out by Tigue et al. (1994); Dercová et al. (1992) and Swamy et al. (1995) working with *B. flathermus*, *B. subtilis* and *Clostridium thermosulfurogenes*, respectively, on

the production amylase. The results demonstrated that after the purification with PEG 1000/phosphate system there was no change in the optimal pH and temperature of the amylase.

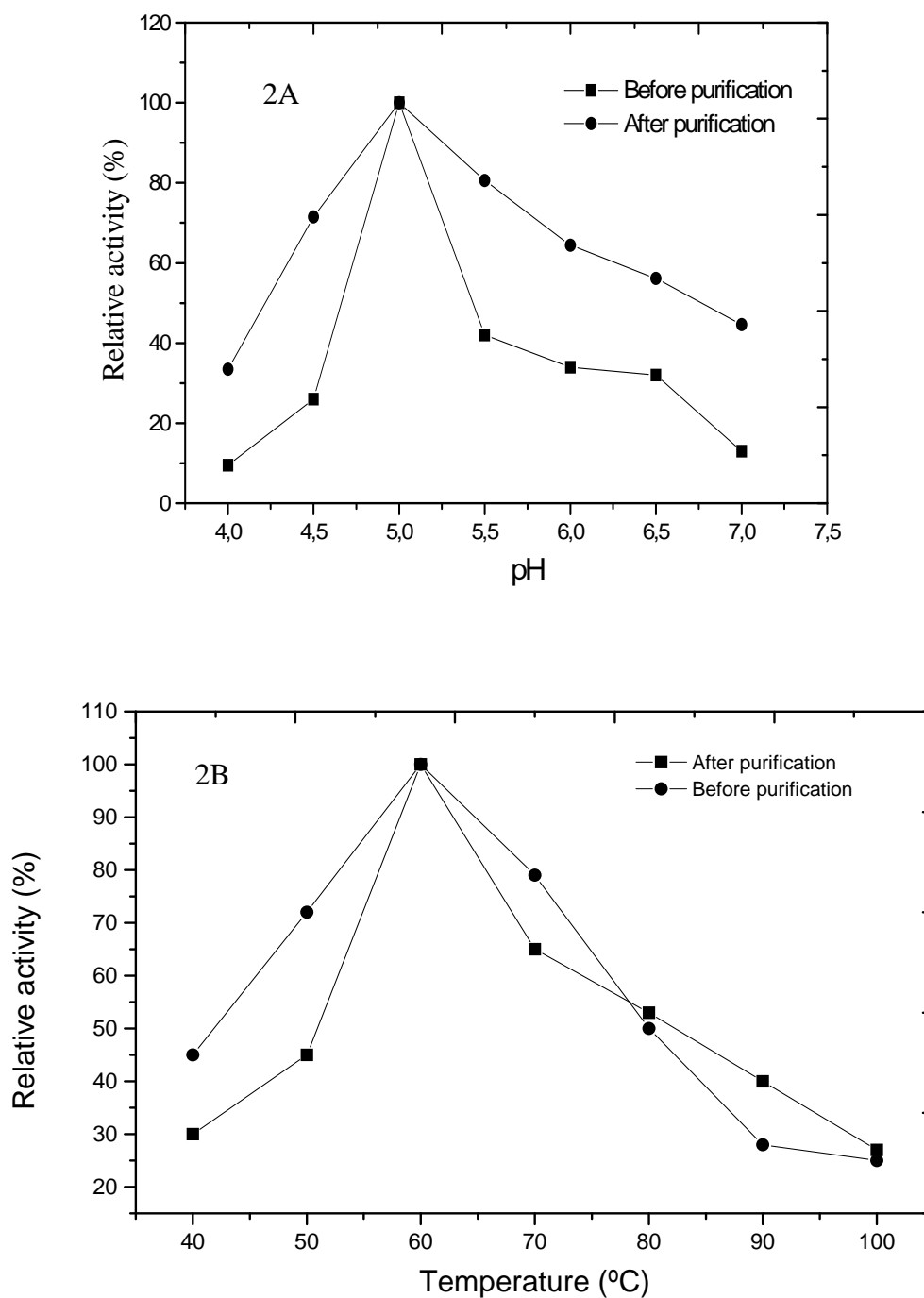


Figure 2 - Effect of pH (A) and temperature (B) on amylase activity before and after purification in the PEG 1000/phosphate system.

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RESUMO

Enzimas amilolíticas têm sido amplamente investigadas com a finalidade de melhorar os processos industriais para a degradação do amido. Foi determinado que a extração da enzima em sistema bifásico aquosos é um método aplicável para separação e purificação de biomoléculas em misturas. Vários sistemas compostos de soluções aquosas de polietilenoglicol e fosfato foram avaliados. Estudos de produção em meio mínimo suplementado, à 37°C, com uma velocidade de agitação de 110rpm e recuperação da amilase a partir do *Bacillus subtilis* em sistema bifásico aquoso PEG-fosfato foram avaliados. O melhor fator de purificação (5.4) foi obtido no sistema PEG 1000 (16.7% w/w) com fosfato de potássio (14.8% w/w), a pH 6.0, resultando na recuperação da atividade enzimática de 45.2% na fase rica em sal.

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