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1 2	The immobilisation of proteases produced by SSF onto functionalized magnetic nanoparticles: Application in the hydrolysis of different protein sources
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## **ABSTRACT**

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Alkaline proteases produced from protein-rich waste (hair waste and soya residues) by solid state fermentation (SSF) were immobilised onto functionalized magnetic iron oxide nanoparticles (MNPs) using glutaraldehyde as a crosslinking agent. The covalent binding method had a better immobilisation yield compared to simple adsorption, retaining 93%-96% (459±106 U/mg nanoparticles, 319±34 U/mg nanoparticles) of hair waste and soya residues proteases, respectively after crosslinking with 5% glutaraldehyde for 6 h. However, the adsorption immobilisation yield was 47%-54% after 8 h for both proteases. MNPs and immobilised proteases were characterized using transmission electron microscopy (TEM), scanning electron microscopy (SEM), Fourier transform infrared spectroscopy (FT-IR) and electron diffraction. Our results indicated successful crosslinking between the proteases and amino-functionalized MNPs. The operational stability (pH and temperature) and storage stability of free and immobilised enzyme were also analysed. Despite the fact that the optimum pH of free and immobilised proteases was identical in the alkaline region, the immobilised proteases reached their optimum condition at higher temperatures (40°C - 60°C 40 °C − 60 °C). After 2 months of storage at 4°C 4 °C, the immobilised proteases showed good stability, retaining more than 85% of their initial activity. The high magnetic response of MNPs render an ease of separation and reusability, which contributes to the residual activity of both immobilised proteases on MNPs remained retaining more than 60% of their initial values after seven hydrolytic cycles. These results resulted showed the enhancement of the stability of the crosslinking interactions between the proteases and nanoparticles. The immobilised proteases were capable of hydrolysing select selected proteins (casein, oat bran protein isolate, and egg white albumin). However, differences in the degree of hydrolysis were observed, depending on the combination of the protease and type of substrate used.

Keywords: protease; protein-rich waste; solid state fermentation; hydrolysis; immobilisation

#### 1. Introduction

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Because of the proteolytic nature of alkaline proteases, they have been commercially utilised for industrial applications, explicitly in the food, pharmaceutical, textile, detergent, and leather industries [1,2]. Their specific role in protein hydrolysis has drawn worldwide attention regarding the versatility of these enzymes for biotechnological applications. Currently, the general cost of protease production is very high, considering the cost of substrates, commercial media and maintenance of cultures used for inoculation. For this reason, the need to develop novel processes with higher yields is highly recommended from a commercial point of view [3]. Different options have been considered to reduce the cost and increase the utilization of proteases. One of the most promising alternatives is solid state fermentation (SSF), which allows the procurement of value-added products using inexpensive waste as a solid substrate. SSF has been successfully applied in the production of proteases using protein-rich waste without the need to inoculate a specific microorganism [4,5]. In addition to recycling the abundant solid waste produced by industries as cheap substrates, SSF avoids the hassle of maintaining cultures since microorganism can develop mutations over time [6]. Economically, no additional cost is required, as there is no restricted sterilized environment required to produce proteases through SSF since a specific microorganism for inoculation is not involved. Apart from that, soluble proteases are susceptible to autolysis, which leads to their rapid inactivation. The instability and lack of flexibility, reusability and recovery make their use a challenge for commercialization. Therefore, the use of immobilisation enzymes offers an attractive method in which immobilised enzymes have enhanced stability that allows for their recyclability and simple recovery without contamination of the final product [7]. There are

five basic enzyme immobilisation methods, including simple adsorption, covalent binding, encapsulation, crosslinking, and entrapment [8].

Of all the methods, covalent binding provides firm binding between the enzyme and carrier and avoids enzyme, as it can be regulated by using specific functional groups (NH<sub>2</sub>, SH) to bind with the proteins. Of all the methods, covalent binding provides firm binding between the enzyme and carrier, thus avoiding enzyme leakage as it can be regulated by using specific functional groups (-NH<sub>2</sub>, -SH) [9]. This method of immobilisation provides an efficient way to increase the stability and flexibility of enzyme reusability and recovery [10]. Coupling agents, such as glutaraldehyde, maleic anhydride and genipin, have been widely used as their functional groups can interact with the functional groups of modified carriers and proteins [10,11]. Glutaraldehyde can be used either to alter enzymes after immobilisation or to activate the support for enzyme immobilisation. In addition, the use of glutaraldehyde can increase protein stability, thus avoiding protease autolysis [2,12].

In recent years, the immobilisation of enzymes onto nanomaterials, particularly iron oxide magnetic nanoparticles (MNPs), to form nanobiocatalysts has attracted much attention in some fields of research, including biolabeling, bioseparation, biosensors, biofuel cells, and environmental analysis [9,13]. The use of MNPs has been particularly attractive for immobilisation because of their special characteristics, such as their high surface area, simple manipulation and separation by the application of an external magnetic field, biocompatibility, biodegradability, and low toxicity [10,14]. There are several studies reporting the application of purified protease immobilised on magnetic supports [15–18]; however there are only a few studies that have exploited MNPs for crude protease immobilisation and further use in protein hydrolysis and synthesis [19].

In this work, the use of relatively inexpensive enzyme preparative for immobilisation onto functionalized MNPs and crosslinking with glutaraldehyde was assessed. The goal was

to test the viability of using low-cost proteases derived from animal (hair waste) and vegetable (soy fibre residues) protein-rich waste that were produced by SSF after being immobilised onto functionalized MNPs in the hydrolysis of different type of proteins. The relative differences in terms of stability and reusability between the free and immobilised enzymes were significant, exhibiting the feasibility of the immobilised enzymes produced in this work. Also, the magnetic properties of the support render a convenient separation between the substrate and the enzymes within the catalytic system.

## 2. Materials and Methods

#### 2.1 Material and reagents

Ferric chloride (FeCl<sub>3</sub>·6H<sub>2</sub>O), ferrous sulphate (FeSO<sub>4</sub>·7H<sub>2</sub>O), (3-aminopropyl)-triethoxysilane (3-APTES, 97%), glutaraldehyde solution (50%), cetyl-trimethyl-ammonium bromide (CTAB), trichloroacetic acid (TCA), Folin-Ciocalteu reagent, casein from bovine milk, and egg white albumin were obtained from Sigma-Aldrich (Spain). Oat bran containing 17.6% protein was purchased from a supermarket (Mercadona, Spain). Hair waste with a protein content of 75.6% was obtained from the local tanning industry located in Igualada, Barcelona, and soya fibre residues with a protein content of 25.1% were received from Natursoy, Spain. β-mercaptoethanol, tricine sample buffer, 10%-20% Mini-PROTEAN gels, and Coomassie 250 stain were purchased from Bio-Rad (Spain). All other chemicals were from commercial sources and were of analytical grade.

## 2.2 Preparation of the protease enzyme concentrates from protein-rich waste using SSF

Two different sources of alkaline proteases were produced from protein-rich waste, hair waste and soya fibre residues using solid state fermentation (SSF) as described elsewhere [4,5]. Briefly, hair waste and soya fibre residues were individually mixed with anaerobically

digested sludge at a weight ratio of 1:2, and then, the mixtures were added to a bulking agent (wood chips) at a volumetric ratio of 1:1. SSF was undertaken in triplicate using 10 L airtight reactors for approximately 1 to 3 weeks. Later, the proteases, Phw from hair waste and Psr from soya fibre residues, were extracted from the reactors using Tris-HCl buffer (50 mM, pH 8.1). The times for extracting proteases for different substrates were specified for each case based on the maximum activity of the protease produced during SSF. Phw at day 14<sup>th</sup> yielded 787±124 U/g DW, and Psr at day 4<sup>th</sup> yielded 634±24 U/g DW. The extracts were ultrafiltrated through Amicon Ultra-15 centrifugal filter devices (Millipore, Ireland) with a 10kDa molecular weight cut-off (MWCO) prior to lyophilisation. The initial activities after lyophilisation for Phw and Psr were 466 U/ml and 330 U/ml, respectively.

## 2.3 Preparation of oat bran protein isolate (OBPI)

The preparation of oat bran protein isolate (OBPI) was performed as described elsewhere [20] with slight modifications. Briefly, oat bran was added to 1.0 M NaCl at a ratio of 1:8 (w/v), and the pH was adjusted to 9.5 using 1.0 M NaOH. The mixture was agitated for 30 min at room temperature. Then, the supernatant was collected after centrifuged at 5,000 x g for 25 min at 4°C 4 °C. The pH was adjusted to 4 with 1.0 M HCl prior to centrifugation at 5,000 x g for 40 min at 4°C 4 °C. The supernatant was then discarded, and the protein isolate was dissolved in Milli-Q water and adjusted to pH 7 with 0.1 M NaOH. The protein isolate was lyophilised and stored at -20°C -20 °C for future use.

## 2.4 Synthesis of amino-functionalized Fe<sub>3</sub>O<sub>4</sub> magnetic nanoparticles (MNPs)

Magnetic nanoparticles (MNPs) were synthesised by co-precipitation in water phase [21,22] with slight modifications. A mixture of 25 mM ferrous sulphate and 50 mM of ferric chloride were dissolved in 100 ml of Milli-Q water with the addition of 0.1% of CTAB as a stabiliser. The mixture was stirred at 40°C 40 °C for 1 hour under a-the nitrogen atmosphere.

Then, 125 ml of deoxygenated NaOH (0.5 M) was added dropwise to the mixture, and the mixture was left for 1 hour at 40°C 40 °C to let the solution chemically precipitate. The resultant MNPs were separated magnetically and washed five times with Milli-Q water. The recovered MNPs were dried overnight at 60°C 60 °C.

The surface of the MNPs was modified using a silanization reaction. Approximately 0.61 g of MNPs was dispersed in a solution containing 3.05 ml of APTES, 0.763 ml of Milli-Q water, and 45.75 ml of methanol. The mixture was ultrasonically agitated for 30 min. Then, 10 ml of glycerol was added and heated at 90°C 90 °C for 6 h, and the mixture was stirred until separation. The surface-modified MNPs were recovered by applying a magnet, and they were washed three times with Milli-Q water.

## 2.5 Immobilisation of alkaline proteases (Phw, Psr)

For immobilisation of surface modified MNPs, 1 ml of alkaline protease from hair waste (Phw) with an initial activity 466 U/ml or alkaline protease from soya residue (Psr) with an initial activity 330 U/ml was dispersed in 9 ml of Tris HCl buffer (pH 8.1) and mixed with 100 mg of amino functionalized MNPs. The activation of the NH2 groups in the nanoparticles was carried out by adding glutaraldehyde as a crosslinking agent at various concentrations (1%, 2.5%, and 5% (v/v)). The mixture was gently agitated at 4°C 4°C for 8 h. Subsequently, the immobilised proteases were separated by a magnetic field, washed five times with Tris buffer (50 mM, pH 8.1) to remove any unbound glutaraldehyde and finally resuspended in 1 ml of Tris buffer (50 mM, pH 8.1). The immobilised proteases were stored at 4°C 4°C for future use.

To begin with immobilisation of protease onto surface-modified MNPs, 1 ml of alkaline protease from hair waste (Phw) with an initial activity 466 U/ml and alkaline protease from

protease from hair waste (Phw) with an initial activity 466 U/ml and alkaline protease from soya residue (Psr) with an initial activity 330 U/ml, respectively, was dissolved in 9 ml of Tris-HCl buffer (pH 8.1). Then, 100 mg of amino-functionalized MNPs was dispersed into

the mixture. Afterwards, the activation of the NH<sub>2</sub> groups in the nanoparticles was carried out by adding glutaraldehyde as a crosslinking agent at various concentrations (1%, 2.5%, and 5% (v/v)) in the mixtures, followed by gentle agitation at 4 °C for 8 h. Subsequently, the MNPs with the immobilised proteases (MNPs-protease) were separated by a magnetic field and washed five times with Tris buffer (50 mM, pH 8.1) to remove any unbound glutaraldehyde and enzyme. Finally, the MNPs-protease were resuspended in 1 ml of Tris buffer (50 mM, pH 8.1) and stored at 4 °C for further application.

For immobilisation via adsorption, approximately 100 mg of naked MNPs were dispersed in 9 ml of the Tris buffer solution (50 mM, pH 8.1). Then, 1 ml of free alkaline protease from hair waste (466 U/ml) or soya residue (330 U/ml) was added. The mixture was gently agitated at 4°C 4 °C for 8 h. Later, the immobilised proteases were magnetically separated and treated as previously described.

The immobilisation yield (%) and the amount of immobilised protease (Phw\_im, Psr\_im)

loading on MNPs (U/mg) were calculated using following equations (Eqs. 1, 2) [15,23]:

189 Enzyme loading 
$$(U/mg) = (U_i - U_{sp})/W$$
 (1)

190 Immobilisation yield (%) = 
$$(U_i - U_{sp})/U_i \times 100$$
 (2)

Where  $U_i$  is the initial enzyme activity (U),  $U_f$  is the enzyme activity in the supernatant after

immobilisation (U), and W is the weight of MNPs used for immobilisation (mg).

193 Furthermore, the immobilisation efficiency (%) and activity recovery (%) were calculated as

194 follows (Eqs. 3, 4) [23]:

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$$Efficiency (\%) = (U_e/U_{imm}) \times 100$$
 (3)

196 Activity recovery (%) = 
$$(U_e/U_i) \times 100$$
 (4)

Where  $U_e$  is the activity of bound enzyme that is measured in the immobilisate (U),  $U_{imm}$  is the immobilised enzyme activity determined from subtracting the remaining enzyme activity in the supernatant from the initial activity (U).

## 2.6 Characterization of the immobilised enzymes

Functionalized MNPs before and after immobilisation were characterized using high resolution transmission electron microscopy (HRTEM, JEM-2011/JEOL) and scanning electron microscopy (SEM, Zeiss Merlin). The samples were prepared by placing a drop of the sonicated solutions on a copper grid, and then, the samples were allowed to dry. Samples with the immobilised enzyme were stained with uranyl acetate (2%) prior to analysis. Functionalized nanoparticle immobilisation was then confirmed using Fourier Transform Infrared spectroscopy (FT-IR, Bruker Tenser 27) within a range of 500-4,000 cm<sup>-1</sup>.

## 2.7 Stability of immobilised protease

The storage stability was determined by maintaining the immobilised enzymes via crosslinking and simple adsorption at 4°C 4 °C for 60 days. The activity of the enzymes was measured at day 0<sup>th</sup> as the initial activity, while the activity for the 60<sup>th</sup> and 7<sup>th</sup> days were used as the final activity of the immobilised enzymes and free enzymes, respectively.

To study the operational stability, both immobilised and free enzymes were incubated for 1 h at various pH and temperature values according to the response surface of the central composite design (CCD) performed using the Design-Expert software (version 6.0.6). The CCD consisted of 13 experimental points, including five replications at the central point and four star points ( $\alpha$  = 1). The pH was adjusted using the following buffers: acetic acid-sodium acetate 1 M (pH 5), Tris-HCl 1 M (pH 8), and Na<sub>2</sub>HPO<sub>4</sub>-NaOH 0.05 M (pH 11). Analysis of variance (ANOVA) was conducted to determine the significance of the main effects.

The residual activity of each factor tested was calculated by that assuming the initial activity of the immobilised or free enzyme was 100%. The residual activity of each factor was calculated by assuming that the initial activity of the immobilised or free enzyme is 100%.

## 2.8 Reusability of immobilised protease

The reusability of immobilised proteases from hair waste (Phw) and soya fibre residue (Psr) was tested on casein as a model protein. The initial activities of the immobilised enzymes were measured and compared with seven consecutive repeated uses of immobilised enzymes under the assay conditions. After each cycle, immobilised enzymes were magnetically separated and washed with Tris-HCl buffer (50 mM, pH 8.1). Then, they were resuspended in fresh medium and incubated at 50°C 50 °C for 120 min. The activity of immobilised enzymes from the first batch was considered to be 100%.

## 2.9 Application of immobilised proteases in protein hydrolysis

Prior to hydrolysis, 4% (w/v) suspension of selected proteins (casein from bovine milk, egg white albumin, and OBPI) in Tris-HCl buffer (50 mM, pH 8.1) were incubated at 50°C 50 °C for 15 min. Then, the reaction was initiated by adding 1 ml of free enzymes (330-460 U/ml) or 1 ml immobilised enzymes suspension (319-459 U/mg NP) into 9 ml of substrate. The mixture was incubated in a water bath at 50°C 50 °C with mechanical agitation at 100 rpm. An aliquot of six 6 ml was withdrawn at 0.5, 2, 4, 6, and 24 h. The free enzyme activity was deactivated by heating the samples in boiling water for 15 min. Then, the samples were cooled by placing the samples in a cold water bath for 15 min. Afterwards, the samples were centrifuged at 5,000 x g for 15 min to separate any impurities or enzyme from the hydrolysate. The immobilised enzyme was separated from the hydrolysate by a magnetic drive. The hydrolysate was kept frozen at -80°C -80 °C prior to lyophilisation.

## 2.10 Analytical methods

## 2.10.1 Protease assay

The proteolytic activity of the free and immobilised protease was determined using casein as a substrate according to method described by Alef and Nannipieri [24] with slight modifications. Briefly, 1 ml of the enzyme extract (free enzyme) or 0.1 g of immobilised protease in 0.9 ml of Tris buffer (pH 8.1) was added to 5 ml of a 2% (w/v) casein solution and incubated at 50°C 50 °C and 100 rpm for 1 hour. The reaction was terminated by adding 5 ml of 15% (w/v) TCA. The samples were centrifuged at 10,000 rpm for 10 min at 4°C 4 °C. An aliquot of 0.5 ml of the supernatant was added to the alkaline reagent prior to the addition of 0.5 ml of 25% (v/v) Folin-Ciocalteu phenol reagent. The resulting solution was incubated at room temperature in the dark for 1 h. The absorbance was measured at 700 nm using a tyrosine standard. One unit of alkaline protease activity was defined as the liberation of 1  $\mu$ g of tyrosine per minute under the assay conditions. All activity tests were performed in triplicate.

## 2.10.2 Total protein content determination

The total protein content was determined by the Lowry method [25] using bovine serum albumin (BSA) as a standard. The absorbance was analysed at 750 nm using an UV-visible spectrophotometer (Varian Cary 50).

# 2.10.3 Degree of hydrolysis

The degree of hydrolysis was determined by quantifying the soluble protein content after precipitation with TCA [19,26]. 1ml of protein hydrolysate was mixed with 1 ml of 10% (w/v) TCA and incubated at  $37^{\circ}$ C for 30 min to allow for precipitation. This was followed by centrifugation (10,000 x g, 10 min). Then, the soluble protein content in the

supernatant was determined by the Lowry method [25], and it was expressed in milligrams.

The degree of hydrolysis (DH) was determined using the following equation (Eq.1):

DH(%) = 
$$\frac{\text{soluble protein content in 10\% TCA}}{\text{total protein content}} \times 100$$
 (5)

## 2.10.4 Electrophoresis

Tricine-SDS PAGE was used to observe the pattern of smaller proteins generated after the hydrolysis reaction. Electrophoresis was performed using 10-20% Mini-PROTEAN Tris-Tricine gels under denaturing and reducing conditions. The reduction was achieved by heating the sample at  $90^{\circ}\text{C}$  90 °C for 5 min in the presence of  $\beta$ -mercaptoethanol (2% v/v). The gel was fixed with methanol (40% v/v) and acetic acid (10% v/v) and subsequently stained with Coomassie Brilliant Blue R-250. Then, the gel was destained with a solution containing methanol, acetic acid, and water (20: 4: 26 v/v).

#### 3. Results and discussion

# 3.1 Immobilisation of Phw and Psr onto magnetic nanoparticles

The Phw and Psr enzymes from SSF were immobilised onto magnetic nanoparticles via simple adsorption and crosslinking with glutaraldehyde (GA). Both methods were carried out for 8 h with the aim of investigating the effect of time and crosslinker concentration on immobilisation (Fig. 1). The simple adsorption yielded a maximum activity recovery of 28% with an activity loading of 87±22 U/mg NP for Phw and of 33% Psr (activity loading of 70±17 U/mg NP) after 8 h of immobilisation (Fig. 1a, 1c). The immobilisation efficiency for both Phw and Psr in simple adsorption increased during 8 h with a maximum of 60-61% efficiency yield, while the maximum immobilisation yield in simple adsorption for both Phw

and Psr were 47% and 54%, respectively (Fig. 1b, 1d). The surfaces of naked MNPs likely possess high reactivity, which makes them susceptible to degradation under particular environmental conditions. This fact could involve weaker binding forces that contribute to the poor stability of the protein attachment's to the surface [9,27].

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Immobilisation via the crosslinker showed good results for both enzymes studied (Phw and Psr). The immobilisation yield increased according to the increase in the GA concentration from 1%-5% up to 6 h; then, it decreased abruptly for Phw (Fig. 1b). Only when 1% GA with Psr was used did the immobilisation yield continue to increase (Fig. 1b). Only when using Psr with 1% GA, the immobilisation yield continues increasing (Fig. 1d). Maximum activity recovery and immobilisation yields were obtained after 6 h of crosslinking time, 90% and 96% respectively, which is (equivalent to an activity load of 459±106 U/mg NP) for Phw with 5% GA (Fig 1a, 1b). Similarly in Psr with 5% GA the maximum activity recovery and immobilisation yield were 92% and 93%, respectively (equivalent to activity loading of 319±34 U/mg NP) (Fig. 1c, 1d). In addition, the immobilisation via crosslinker was superior to simple adsorption as both of the enzymes (Phw, Psr) showed good immobilisation efficiency in the range of 45% to 98% during 6 h of immobilisation time (Fig. 1b, 1d). This indicated that the crosslinking time and GA concentration play an important role during the immobilisation of enzymes in this study. As GA plays a role as a spacer arm for the carriers by providing aldehyde groups to couple to free amine groups from the enzymes, forming imines, it can also act as a denaturing agent [12]. Additionally, some studies obtained different crosslinking times (between 1 h to 4 h) and GA concentrations (from 1% to 6%), implying good biocompatibility for these specific enzymes [12,21,28,29].

## 3.2 Characterization of the functionalized nanoparticles used for immobilisation

Transmission electronic microscopy (TEM) images of MNPs before and after modification with APTES and after the enzymes immobilisation onto the activated surface

were compared (Fig. 2). The average particle size of naked MNPs slightly increased from 10.2 nm (Fig. 2a) to 16.1 nm (Fig. 2b) after surface modification with APTES. This effect has been observed previously in other studies [30,31]. After surface modification with APTES, fewer nanoparticle aggregates formed. As suggested previously [10], surface modifications of magnetic nanoparticles can improve their solubility and help avoid aggregation of particles. In Fig. 2c and Fig. 2d, a layer covering surface of MNPs upon immobilisation of the proteases (Phw and Psr) can be seen. The thickness of this layer covering the surface of MNPs was estimated to be approximately 5.1 nm for Psr and 8.4 nm for Phw, indicating an increase in the size of the particles. Based on electron diffraction analysis (Fig. 3) of the TEM images, the crystalline structure of the particles was not affected by surface modifications. Fig. 3a shows a clear loop, confirming the crystalline structure of MNPs. After surface modification by APTES or CTAB as a stabilizer, the crystalline structure was not modified; however, the size of some nanoparticles was enlarged, as observed in Fig. 3b and Fig. 3c. Once protease immobilisation was performed, the structure of the nanoparticles became an amorphous structure, confirming that the enzyme covered the surface of the nanoparticles (Fig. 3d). The surface of the naked MNPs and functionalized MNPs can be observed in SEM images (Fig. 4a and Fig. 4b). The small and spherical particles with well-defined edges are observed as in other studies [32]. In contrast, in Fig. 4c and Fig. 4d, the edge surface of nanoparticles is smooth because they are covered by the enzymes, indicating that the immobilisation of proteases onto functionalized MNPs was successful. The surface modification and immobilisation of proteases (Phw and Psr) onto nanoparticles was confirmed by a comparison of the FT-IR spectra of naked MNPs, functionalized MNPs, and Phw and Psr immobilised onto functionalized MNPs. The FT-IR

spectrum in Fig. 5A shows a strong absorption peak at 584 cm<sup>-1</sup>, which could corresponds to

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341 Fe-O, as indicated in other studies [33,34]. It has been suggested that this strong peak could be due to the stretching vibration mode associated with metal-oxygen absorption. In this 342 region, the stretching vibration peaks related to metal (ferrites in particular) in the octahedral 343 and tetrahedral sites of the oxide structure were found [31]. In Fig. 5A, the peaks at 1,662 cm<sup>-1</sup> 344 <sup>1</sup> and 3,444 cm<sup>-1</sup> were due to the bending and stretching vibration of -OH, respectively [35]. 345 After grafting with APTES, the characteristic peak of the Fe-O bond shifted from 584 cm<sup>-1</sup> to 346 638 cm<sup>-1</sup> and 640 cm<sup>-1</sup> because of the formation of the Fe-O-Si bond (Fig. 5B, 5C, 5D). The 347 shifting of the absorption peaks to high wavenumbers is due to the greater electronegativity 348 of -Si(O-) compared to H, which contributes to the bond forces for Fe-O bonds [36]. 349 Additional strong peaks at 1,039 cm<sup>-1</sup>, 1,035 cm<sup>-1</sup>, and 1,034 cm<sup>-1</sup> correspond to the Fe-O-Si 350 351 bending vibrations, indicating that alkyl silanes are successfully attached to functionalized MNPs (Fig. 5B, 5C, 5D). Additionally, the presence of silane groups was observed at 995 cm<sup>-</sup> 352 <sup>1</sup>, 893 cm<sup>-1</sup>, and 896 cm<sup>-1</sup> and were from the stretching vibrations of the Si-OH and Si-O-Si 353 groups from APTES [35,36]. Characteristic peaks of the immobilised enzymes attached via 354 the crosslinker (Fig. 5C, 5D) were observed at 1,536 cm<sup>-1</sup>, 1,544 cm<sup>-1</sup> and 1,630 cm<sup>-1</sup> because 355 of the C=N and C=O absorption from the glutaraldehyde and NH<sub>2</sub> from the enzyme [29]. 356 Small shifts in intensity from 2,928 cm<sup>-1</sup> (Fig. 5B) to 2,991 cm<sup>-1</sup> (Fig. 5C) and 2,984 cm<sup>-1</sup> 357 (Fig. 5D) correspond to the C-H stretching vibration from the methyl group [37], which 358 illustrated the effect before and after the immobilisation of the enzymes. Additionally, in Fig. 359 5C and Fig. 5D, there were broad and strong peaks at 3,325 cm<sup>-1</sup> and 3,344 cm<sup>-1</sup>, which 360 indicated the vibration modes of the O-H and -NH groups from enzymes that interact with 361 nanoparticles, which has been suggested previously [19]. 362

## 3.3 Operational stability of immobilised Phw and Psr

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The operational stability in terms of temperature and pH is an important criterion in the application of immobilised enzymes [2,38]. To study this factor (Phw\_im, Psr\_im), various

366 pH values (5-11) and temperatures  $(30^{\circ}\text{C} \cdot 70^{\circ}\text{C})$  30 °C  $-70^{\circ}\text{C}$  were tested, and the results were compared with those of free enzymes (Phw\_free, Psr\_free). The results were analysed 367 using analysis of variance (ANOVA) to indicate the significant factor influencing the stability 368 369 of both enzymes. The ANOVA results in Table 1 show that the regression coefficients had a high statistical significance (p<0.05) and show the values obtained for the coefficient of 370 determination for both Phw\_im and Psr\_im (R2 0.9730 and R2 0.9733) and Phw\_free and 371 Psr\_free (R<sup>2</sup> 0.9723 and R<sup>2</sup> 0.9712, respectively). The values indicated that the model of the 372 immobilised enzymes could not explain only 2.7% of the variables behaviour, while with the 373 374 free enzymes the value was 2.8-2.9%. For immobilised enzymes, the calculated F-value  $(\alpha=0.05, DOF=4,3)$  was 9.12 for the regression. This value was higher than the tabulated F-375 values (1.65, 1.85), indicating that the treatment differences were highly significant. 376 377 Similarly, in free enzymes, the obtained F-values (2.90, 3.94) were less than the critical Fvalue ( $F_{0.05(4.5)} = 5.19$ ), reflecting the significance of the model. The following Equations (6 – 378 9) represent the second order polynomial model of the residual activity for the experimental 379 380 data:

Residual Phw\_im(%) = 
$$-407.2 + 56.5$$
pH +  $10.1$ T -  $2.6$ pH<sup>2</sup> -  $0.1$ T<sup>2</sup> -  $0.16$ pHT (6)

Residual Psr\_im(%) = 
$$-275.9 + 41.8$$
pH +  $6.3$ T  $- 1.8$ pH<sup>2</sup>  $- 0.05$ T<sup>2</sup>  $- 0.17$ pHT (7)

Residual Phw\_free(%) = 
$$22.2 + 2.5$$
pH +  $1.5$ T -  $0.02$ 6T<sup>2</sup> (8)

Residual Psr\_free(%) = 
$$66.4 + 1.04$$
pH +  $0.54$ T -  $0.021$ T<sup>2</sup> (9)

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For free enzymes, the models were reduced by removing the interaction between pH and temperature, as it was not significant to the stability of the free enzymes.

Contour plots of the second order polynomial model were generated as a function of the independent variables of pH and temperature for immobilised and free enzymes. The contour plots of free Phw and Psr exhibited their stability under mesophilic conditions (30°C 30 °C to 40°C). Free enzymes were stable over a broad range of pH values (Fig. 6b, 6d), with no

optimum condition obtained in the range tested (pH 5-11). However, both of the immobilised enzymes had improved the stability by achieving their optimum condition in the alkaline region (pH 8 to 11) with thermophilic temperature stability ranging from 40°C 40 °C to 60°C 60 °C for immobilised Phw (Fig. 6a) and 40°C 40 °C to 55°C 55 °C for immobilised Psr (Fig. 6c).

## 3.4 Storage stability of immobilised Phw and Psr

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Storage stability plays a crucial role in the use immobilised proteases, as the shelf life determines the viability of an immobilised enzyme over time [39]. The storage stabilities of enzymes immobilised via a crosslinker (Phw\_GA and Psr\_GA) and adsorption (Phw\_adsorp and Psr\_adsorp) were tested by dispersing the immobilised enzymes in Tris buffer and maintaining them at 4°C 4 °C for 60 days. Free enzymes (Phw\_free and Psr\_free) were used as controls to monitor the durability of enzyme activity. Phw free and Psr free were not stable in solution, as their activity decreased over time. This fact could be related to the behaviour of the proteases, as they tend to autolyse themselves by nucleophilic attack on the intermediate in a presence of water [1,40]. After 7 days of storage at 4°C 4°C, the residual activity of Phw\_free and Psr\_free was less than 17% (Table 2). There was a significant decrease in the activity of the immobilised enzyme via adsorption over 60 days of storage, with a residual activity of less than 45%. The weak bonding between the enzyme and nanoparticles could induce partial desorption during the period of storage. Generally, immobilisation via adsorption involves relatively weak interactions, such as electrostatic interactions, hydrogen bonds, van der Waals forces and hydrophobic interactions, which tend to strip off enzymes from the carrier easily, thus leading to a loss of activity and contamination of the reaction media [9]. However, the enzymes immobilised via crosslinking (Phw\_GA and Psr\_GA) retained 91% and 86% of their residual activity, respectively, after 60 days of storage at 4°C (Table 2). These results show that enzymes immobilised created by crosslinking provide a distinctive advantage in stability over immobilised enzymes created by adsorption at a longer duration of storage.

## 3.5 Reusability of immobilised Phw and Psr

For the sake of the cost-effective use of enzymes, reusability is a critical factor to consider [41]. The reusability of immobilised Phw and Psr created using crosslinking was evaluated in a repeated batch process using fresh casein as a model protein in each batch cycle (Fig. 7). Both immobilised Phw and Psr retained 66% and 64% of their activity, respectively, after 7 cycles, indicating a significant enhancement of the stability of the crosslinking interaction between the proteases and nanoparticles. In this work, testing the reusability was feasible, as the immobilised enzymes were easily separated by a magnetic force.

# 3.6 Application of immobilised Phw and Psr in the hydrolysis of proteins

Casein has been used as a model protein to evaluate the degree of hydrolysis (DH) of immobilised enzymes created via crosslinking and free Phw or Psr. The results of the hydrolysis are shown in Fig. 8a. The DH of immobilised Phw (24%) was higher than that of free Phw (15%) during the first 30 min of reaction time (Fig. 8a). The maximum DH was achieved after 24 h and was 80% for immobilised Phw and 40% for free Phw. Besides, the residual activity of both immobilised Phw and Psr were stable for 24 h whereas the residual activity of free Phw and Psr had decreased pronouncedly after 24 h at 50 °C indicating an improvement of process efficiency and thermal stability of the immobilised enzymes [19]. The higher stability may be possibly due to the multipoint covalent binding of protease to the solid support that limit the flexibility and conformational mobility of the enzyme, hence inhibits unfolding or denaturation of the enzyme [42]. Likewise, the DH of immobilised Psr

(12%) showed the same rate as that of free Psr (13%) and continued to increase over time with a similar profile, reaching the maximum DH 30% in free Psr and 50% in immobilised Psr after 24 h of reaction time. Thus, the immobilised enzymes in the present study could enhance the ability of free enzymes to hydrolyse protein, as shown by the model protein, reflecting that the active enzymes were successfully immobilised.

The effect of both proteases (Phw and Psr) was also evaluated in hydrolysis of the protein source from animals (egg white albumin) and vegetables (oat bran protein isolate). The hydrolysis of oat bran protein isolate (OBPI) corroborated the efficiency of immobilised Phw and Psr, as they had a higher DH compared to free Phw and Psr over a longer duration (Fig. 8b). These results agree with those obtained using functionalized magnetite nanoparticles to hydrolyse soya protein isolates and whey protein isolates [29,43]. It seems that Psr exhibited a higher DH than (free and immobilised) Phw when using a vegetable protein source. In the hydrolysis of egg white albumin, immobilised Psr reached a maximum DH 60% after 24 h and free Psr achieved maximum DH 45% after 6 h, while Phw reached a maximum DH of 70% and 51% for immobilised and free Phw, respectively (Fig. 8c). Hence, the different source of the proteases seems to be a distinct behaviour that depends on the substrate source; in fact, it could determine the choice of enzymes according to the application, as suggested by others [2].

SDS-PAGE of the hydrolysates from selected proteins (casein, OBPI, egg white albumin) is shown in Fig. 9. The presence of several smaller peptides with low molecular weights support the idea that proteins were hydrolysed by immobilised protease derived from an animal source (Phw) and vegetable source (Psr). Similar profiles were obtained for both immobilised enzymes (Phw and Psr), showing their ability to perform the reaction after being grafted on nanoparticle surfaces created by crosslinking with glutaraldehyde.

#### 4. Conclusion

Low-cost proteases obtained through the SSF of protein-rich wastes (hair waste and soya fibre residues) were successfully immobilised onto functionalized MNPs over a relatively short time. The ease of the separation and reusability of these enzymes in comparison with free enzymes could be considered an advantage of their use in industrial processes. Additionally, stability was enhanced from mesophilic to thermophilic conditions under alkaline conditions, preventing autolysis of the enzymes and maintaining their initial activities for 2 months with only 9%-14% activity loss. Immobilised Phw and Psr were able to hydrolyse some proteins derived from plant and animal sources with a high degree of hydrolysis, indicating that they are promising for immobilised enzyme applications in a wide range of industrial processes.

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## 621 Tables

**Table 1** Analysis of variance (ANOVA) for the response surface quadratic model for immobilised (Phw\_im and Psr\_im) and free (Phw\_free and Psr\_free) enzymes.

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Protease	Source of variation	Sums of square	Degree of freedom	Mean square	F-value	Prob>F
		1010= 1		2424 #	<b>70.45</b>	0.0001
Phw_im	Regression	12107.4	5	2421.5	50.47	< 0.0001
	Residual	335.9	7	47.98		
	Pure error	150	4	37.5		
	Lack of fit	185.9	3	61.95	1.65	
	Total	12443.2	12			
Psr_im	Regression	5108.7	5	1021.7	50.99	< 0.0001
	Residual	140.3	7	20		
	Pure error	58.8	4	14.7		
	Lack of fit	81.5	3	27.2	1.85	
	Total	5248.9	12			
Phw_free	Regression	3382.4	3	1127.5	105.4	< 0.0001
	Residual	96.3	9	10.7		
	Pure error	20.8	4	5.2		
	Lack of fit	75.5	5	15.1	2.9	
	Total	3478.7	12			
Psr_free	Regression	6452.3	3	2150.8	101.1	< 0.0001
	Residual	191.5	9	21.3		
	Pure error	32.3	4	8.1		
	Lack of fit	159.2	5	31.8	3.94	
DI : D2.0	Total	6643.8	12			

Phw\_im: R<sup>2</sup> 0.9730, adj R<sup>2</sup> 0.9537, pred R<sup>2</sup> 0.8519 Psr\_im: R<sup>2</sup> 0.9733, adj R<sup>2</sup> 0.9542, pred R<sup>2</sup> 0.8260 Phw\_free: R<sup>2</sup> 0.9723, adj R<sup>2</sup> 0.9631, pred R<sup>2</sup> 0.9315 Psr\_free: R<sup>2</sup> 0.9712, adj R<sup>2</sup> 0.9616, pred R<sup>2</sup> 0.9224

**Table 2** The storage stability of free and immobilised enzymes created via crosslinking with glutaraldehyde (GA) and via adsorption (adsorp) during 60 days of storage.

Enzymes	Initial activity (U/ml)	Final activity (U/ml)	Residual activity (%)
Phw_GA	501±72	458±51	91
Psr_GA	346±69	297±84	86
Phw_adsorp	190±15	77±9	41
Psr _adsorp	152±9	46±5	30
Phw_free	537±26	91±21 <sup>a</sup>	17
Psr_free	358±29	$42\pm3^a$	12

The standard deviation was calculated from 3 replicates.

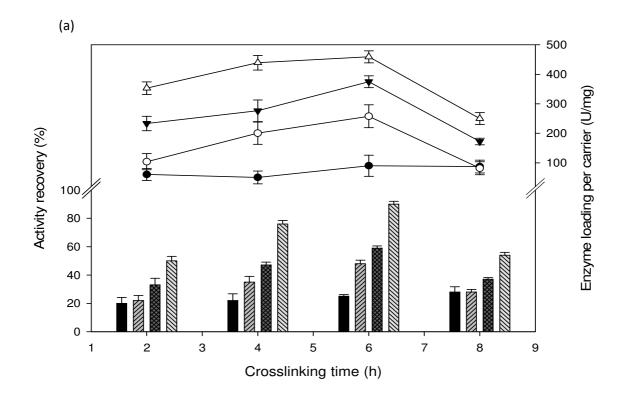
<sup>&</sup>lt;sup>a</sup> The final activity was determined after 7 days of storage.

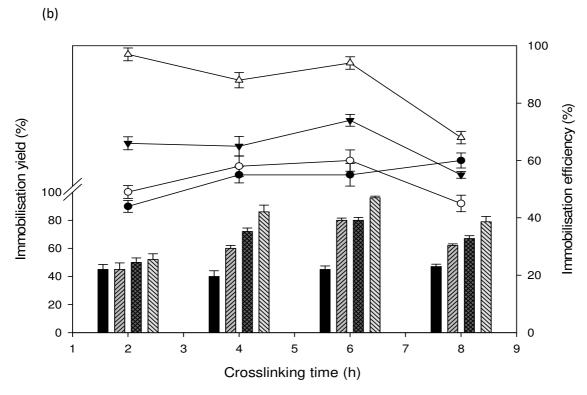
Figure captions

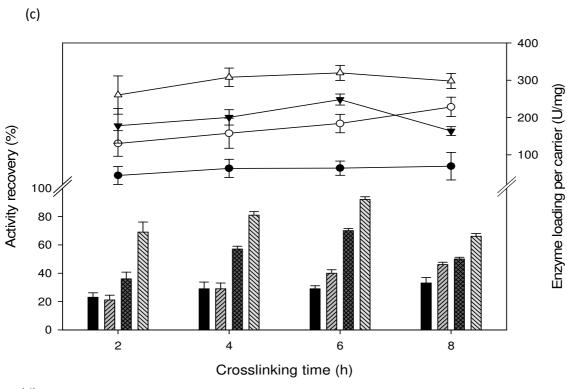
- 670 Fig. 1 The effect of the glutaraldehyde (GA) concentration and crosslinking time on the
- activity recovery of (a) Phw and (c) Psr with 0% GA (simple adsorption) ( ), 1% GA ( ),
- 672 2.5% GA ( ), and 5% GA ( ); enzyme loading per carrier of (a) Phw and (c) Psr with 0%
- GA (simple adsorption) ( $\rightarrow$ ), 1% GA ( $\rightarrow$ ), 2.5% GA ( $\rightarrow$ ), and 5% GA ( $\rightarrow$ );
- immobilisation yield of (b) Phw and (d) Psr with 0% GA (simple adsorption) ( ), 1% GA (
- 675 Z), 2.5% GA (M), and 5% GA (M); immobilisation efficiency of (b) Phw and (d) Psr
- 676 with 0% GA (simple adsorption) (→), 1% GA (→), 2.5% GA (→), and 5% GA (
- 677  $\longrightarrow$  onto amino-functionalized MNPs.
- Fig. 2 TEM images at a magnification of 30,000x of (a) naked MNPs, (b) amino-
- functionalized MNPs, (c) amino-functionalized MNPs after being immobilised with Phw, and
- 680 (d) amino-functionalized MNPs after being immobilised with Psr.
- **Fig. 3** The electron diffraction images of (a) naked MNPs, (b) MNPs after the addition of
- 682 CTAB, (c) MNPs after surface modification with APTES, and (d) MNPs after the
- 683 immobilisation of the enzymes Phw and Psr.
- Fig. 4 The SEM images at 50.00 KX magnification of (a) naked MNPs, (b) amino-
- 685 functionalized MNPs, (c) amino-functionalized MNPs after being immobilised with Phw, and
- (d) amino-functionalized MNPs after being immobilised with Psr.
- Fig. 5 The FT-IR spectra of (A) naked MNPs, (B) amino-functionalized MNPs, (C) amino-
- 688 functionalized MNPs after immobilisation of Phw, and (D) amino-functionalized MNPs after
- 689 immobilisation of Psr.
- **Fig. 6** Contour plots of the residual activity (%) of the enzymes in terms of their operation
- stability as a function of pH and temperature of (a) immobilised Phw, (b) free Phw, (c)
- 692 immobilised Psr, and (d) free Psr

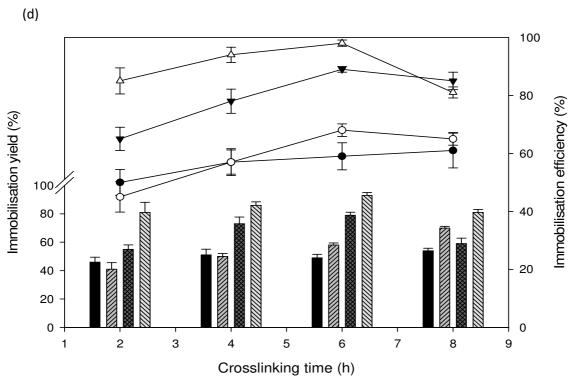
693	<b>Fig. 7</b> The reusability of immobilised Phw (■) and immobilised Psr (□) onto amino-
694	functionalized MNPs using casein as a model substrate for the hydrolysis reaction
695	Fig. 8 The degree of hydrolysis of the selected protein hydrolysates obtained from the
696	hydrolysis of immobilised Phw (—○—), free Phw (—◆—), immobilised Psr (—△—), and free Psr
697	( with (a) casein; (b) oat bran protein isolate (OBPI); and (c) egg white albumin. The
698	residual activity of immobilised Phw ( ¬¬), free Phw ( ¬¬), immobilised Psr (¬¬) and
699	free Psr ( in 50 °C during 24h.
700	Fig. 9 SDS-PAGE of the hydrolysates after reduction using 2-mercaptoethanol for selected
701	proteins before and after treatment with the immobilised proteases; (1) Molecular mass
702	marker; (2) casein; (3) egg white albumin; (4) hydrolysate from casein after treated with
703	immobilised Phw; (5) hydrolysate from egg white albumin after treatment with immobilised
704	Phw; (6) hydrolysate from OBPI after treatment with immobilised Phw; (7) hydrolysate from
705	casein after treatment with immobilised Psr; (8) hydrolysate from egg white albumin after
706	treatment with immobilised Psr; (9) hydrolysate from OBPI after treatment with Psr.
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**Fig. 1** 

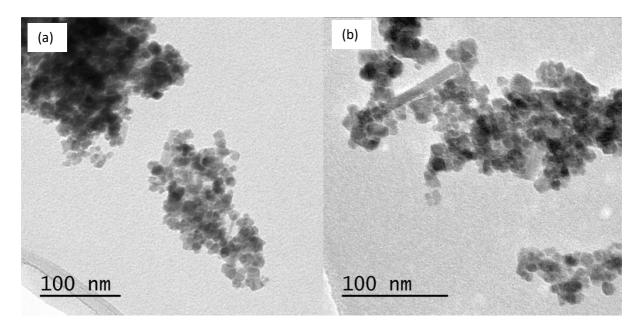


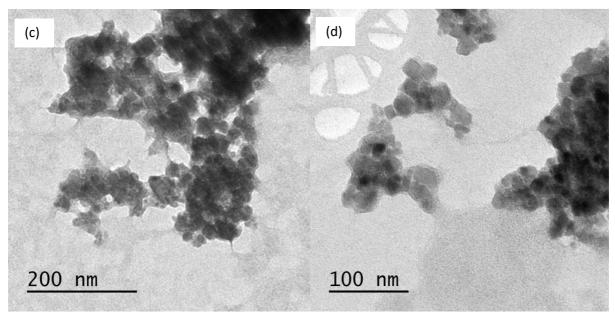




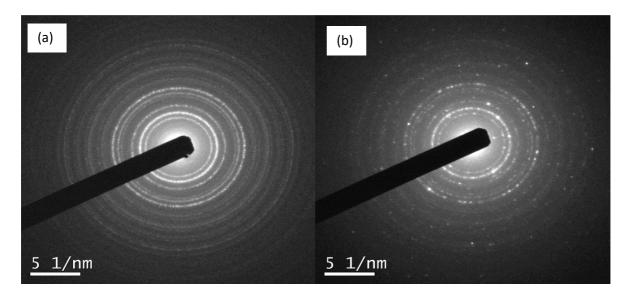


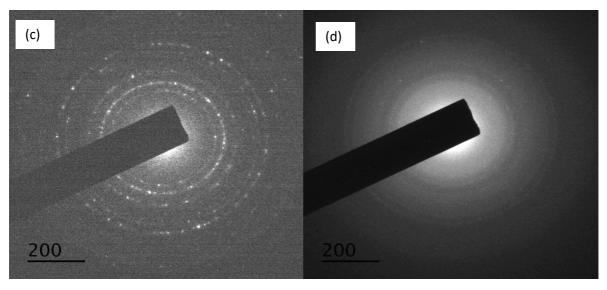
**Fig. 2** 



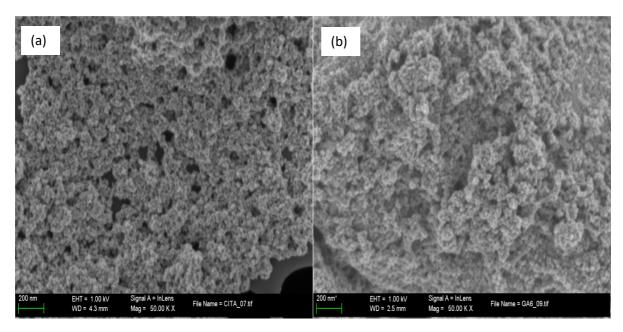


# **Fig. 3**





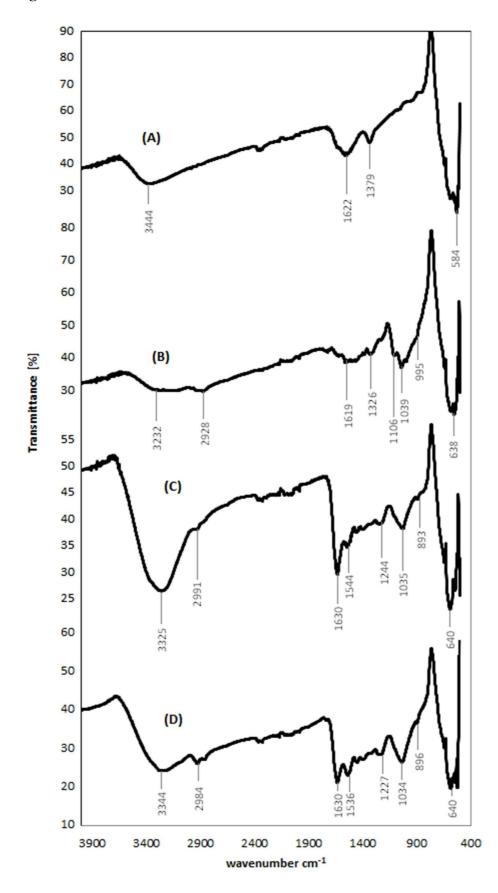
**Fig. 4** 



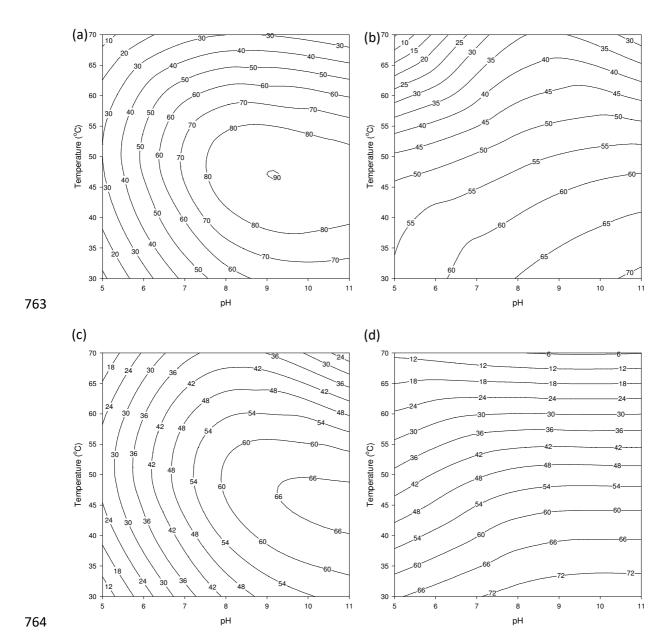
(d)

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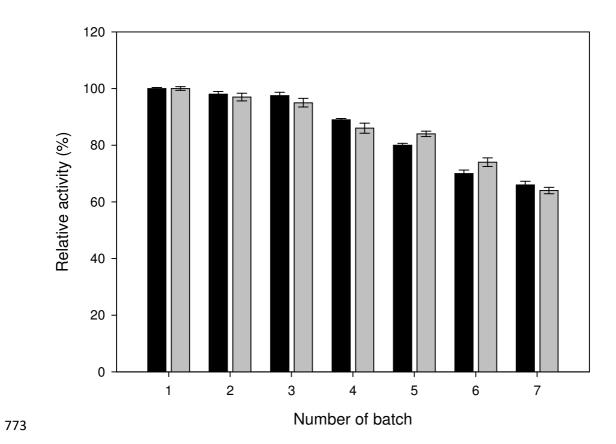
**Fig. 5** 



# **Fig. 6**

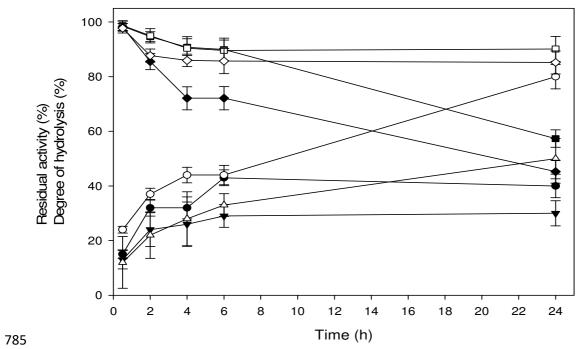


**Fig. 7** 

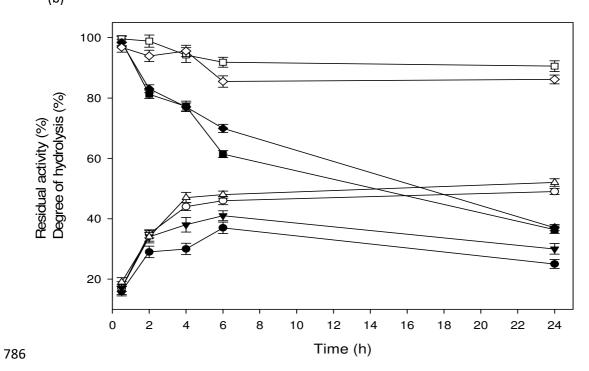


784 **Fig. 8** 

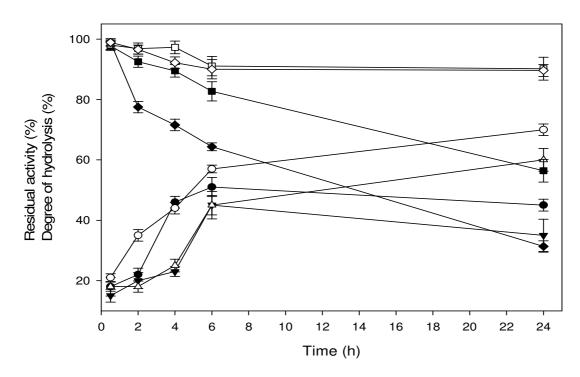
(a)



(b)



(c)



**Fig. 9** 

