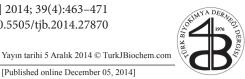
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Kinetic and structural characterization of interaction between trypsin and *Equisetum arvense* extract

Trypsin ve *Equisetum arvense* özütü arasındaki etkileşimin kinetik ve yapısal karakterizasyonu]

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ABSTRACT

Objective: In this study the inhibitory effect of E. arvense extract on trypsin activity and the effect of trypsin on E. arvense extract were studied. In addition the nature of the interaction between the extract and trypsin was investigated.

Methods: The inhibitory effect ethanol extract of E. arvense on trypsin activity was determined using trypsin enzyme assay. The structural effects of the extract-trypsin interaction for the extract were analyzed by FTIR. Finally, the HPLC analyses were carried out to analyze the individual components of the extract and the supernatant and soluble precipitate phases.

Results: E. arvense extract was found to decrease total percent activity of trypsin to 5% in 24 hour at 24 °C. FTIR analyses indicated that the interaction between trypsin and E. arvense extract caused changes in the structure and hydrogen bonding behavior and composition of the extract proteins. These interactions also caused the extract lipids to accumulate in the insoluble precipitate phase. Most of the phenolics remained in the supernatant phase enhancing the inactivation of trypsin. However, the precipitated compounds were shown to be of apolar in nature as shown in the HPLC chromatograms.

Conclusion: The methods that were used showed that the high phenolic content of *E. arvense* was the main reason for the inhibition of trypsin enzyme activity by denaturing the enzyme. Key Words: Trypsin, Equisetum arvense, fourier transform infrared spectroscopy, phenolics Conflict of Interest: Authors have no conflict of interest.

ÖZET

Amaç: E. arvense özütünün tripsin aktivitesi üzerinde ki engelleyici etkisinin ve de tripsinin özüt üzerinde ki etkisi bu çalışmada araştırılmıştır. Bunlara ek olarak ekstrakt ve tripsin arasında ki etkileşimin doğası araştırılmıştır.

Metod: E. arvense bitkisinin etanol özütünün tripsin üzerinde ki engelleyici etkisi tripsin enzim analizi ile belirlenmiştir. Özüt tripsin etkileşiminin özüt üzerinde ki yapısal etkisi FTIR ile analizlenmiştir. Son olarak özütün, süpernetantın ve çözünebilir pelletin özgün bileşenlerin analizi ise HPLC ile belirlenmiştir.

Bulgular: E. arvense özütünün total tripsin aktivitesini 24 °C sıcaklıkta ve 24 saat sonunda %5'e kadar düşürdüğü bulunmuştur. Yapılan FTIR analizleri ile tripsin ve E. arvense özütü arasındaki etkileşimin özüt proteinlerinin yapısında, kompozisyonunda ve hidrojen bağı yapma davranışında değişikliklere sebep olduğu anlaşılmıştır. Bu etkileşimler sebebiyle özütte bulunan lipidlerin çözünmez haldeki çökelti fazında toplandığı gözlemlenmiştir. Ayrıca fenolik bileşiklerin çoğunluğunun süpernetant fazında bulunması sebebiyle tripsin inaktivasyonunu arttırdığı anlaşılmıştır. Ancak çökeltide ki bileşiklerin apolar yapıda olduğu HPLC koromotogramları ile gösterilmiştir.

Sonuç: E. arvense bitkisinin sahip olduğu yüksek fenolik madde miktarının tripsin enzim aktivitesinin engellenmesinde enzim denatürasyonunun ana sebep olduğu kullanılan metodlar ile gösterilmiştir.

Anahtar Kelimeler: Tripsin, E. arvense, fourier transform infrared spektroskopisi, fenolikler Çıkar Çatışması: Yazarların çıkar çatışması yoktur.

Introduction

Phytochemicals are chemical compounds that are found in plants naturally and provide positive heath effects that are important in biotechnology [1,2,3]. Polyphenolic compounds can interact with protein molecules via weak bonds such as hydrogen bonds. They generally form soluble complexes but as the complexes grow larger they become insoluble because the polyphenols form bridges by non-covalent interactions between protein molecules and other complexes [2,4].

One of the plants used in alternative medicine is E. arvense, commonly known as field horsetail. This plant can be found in moist areas of northern hemisphere. The plant material contains silicic acid, tartaric acid, methyl esters of protocatechuic and caffeic acids as phenolic acids [5,6]. The dominant phenolic compounds are identified as isoquercitrin, apigenin, kaempferol in the plant extract [7,8]. In folk medicine horsetail tea is used for cure of edema, kidney and bladder stones, urinary tract infections, the inability to control urination and general disturbances of the kidney and bladder [9,10]. Recent investigations have shown that the plant has high antioxidant capacity [11]. Similarly other researchers have indicated that the plant material has anti-inflammatory, anti-microbial, anti-cancer, sedative and anticonvulsant effects [12-15]. Lately E. arvense is used in the bathing water for wound healing and blood coagulation in folk medicine [9,16].

Trypsin inhibitors have many benefits for the organisms. Although trypsin inhibitors have many benefits their concentration in the blood should be controlled. An increase in trypsin inhibitor concentration causes hyper secretion of pancreatic enzymes. As a result of this hyper secretion, the pancreas becomes enlarged. This will cause the diseases called hypertrophy and hyperplasia [17,18]. For this reason trypsin inhibition potential of foods should be characterized.

In this study the inhibitory effect of *E. arvense* extract on trypsin activity and the effect of trypsin on *E. arvense* extract were investigated. Trypsin enzyme assays were carried out to determine the inhibitory effect *E. arvense* extract on trypsin activity. FTIR analyses were used to study the structural effects of the extract-trypsin interaction for the extract in the both supernatant and precipitate phases formed after bringing these two materials together. Total phenolic content of the extract, the supernatant and soluble precipitate phases were also determined. Finally, the HPLC analyses were carried out to analyze the individual components of the extract and the supernatant and soluble precipitate phases.

Materials and Methods

Preparation of equisetum arvense extract

Powdered predried *E. arvense* leaves were mixed with aqueous 70% ethyl alcohol solution at a solid-liquid ratio

of 1:20. The dry leafs particles were shaked in the solution for overnight at 300 rpm at 24 °C. After the centrifugation at 2000xg for 5 minutes the supernatant was collected into 50 ml falcon tubes. The alcohol in the liquid phase was evaporated with a rotary evaporator. The aqueous extract was frozen for 24 hours and then freeze dried in a lyophilizer (Telstar-Cryodos).

Preparation of inhibition solution

E. arvense extract was dissolved in distilled water to a final concentration of 10 mg/ml. By serial dilutions of stock solution 2 mg/ml, 0.2 mg/ml and 0.02 mg/ml concentrations were prepared.

Preparation of trypsin solution

Bovine pancreatic trypsin was purchased from Fluka Analytical (cat.93610) and used without further purification. Lyophilized powder of bovine pancreatic trypsin (Fluka Analytical, USA) was dissolved in 1 mM HCl to a final concentration of 0.1 mg/ml.

Determination of enzyme kinetics

Trypsin catalyzes the hydrolysis of N-benzoyl-L-arginine ethyl ester (BAEE) to N-benzoyl-L-arginine (BA) and ethanol. The slope of the absorbance versus time graph was the production rate of BA at pH 7.6. The reaction was held in a total volume of 3.2 ml quartz cuvette at 25 °C. The unit of trypsin activity was previously determined as 1 µmol BA produced per minute at 253 nm, at pH 7.6 and 25 °C. The reaction was carried out in 100 mM Tris buffer solution. To determine absorption of blank solution, 3050 µl tris buffer and 150 µl of BAEE solution were mixed in a quartz cuvette and the absorbance was read at 253 nm. Enzyme kinetics studies were carried out in 2850 µl tris buffer, 150 µl BAEE solution and 200 µl of 0.05 mg/ml trypsin mixture solution.

Determination of enzyme inhibition

Inhibition solution was prepared by mixing various concentration of *E. arvense* extract with 0.5 mg/ml trypsin in a ratio of 1:1. Final concentrations of the compounds in the inhibition solution were 1 mg/ml, 0.1 mg/ml, 0.01 mg/ml for *E. arvense* extract and 0.05 mg/ml for trypsin. Incubation of the mixture was done at +24°C and +4°C for minimum 24 hours. In a quartz cuvette, 3050 µl tris buffer, 150 µl BAEE solutions were mixed and the absorbance values at 258 nm were read to determine the blank. For inhibition assay, 2850 µl tris buffer, 150 µl BAEE solution and 200 µl trypsin-*equisetum arvense* mixture were mixed in a quartz cuvette and the enzyme activity assay was carried out.

Processing of the precipitate

After the inhibition assay, centrifugation was performed at 10,000xg for 15 minutes. The supernatant was removed and the pellet was dissolved and kept in 1 mM HCl solution with occasional vortexing. The mixture was centrifuged at 5000xg for 10 min and the enzyme assay was carried out.

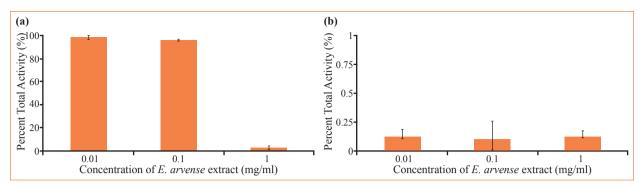


Figure 1. Percent total activity of (a) supernatant (precipitate formed after enzyme extract interaction) and (b) soluble pellet (soluble part of the precipitate formed after enzyme extract interaction) after the interaction of plant extract and trypsin at 24 $^{\circ}$ C for 60 hr.

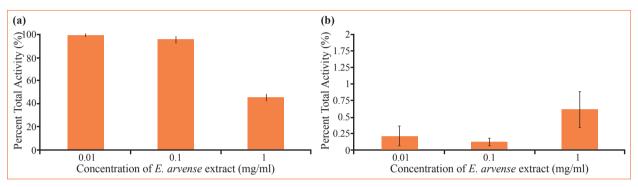


Figure 2. Percent total activity of supernatant (precipitate formed after enzyme extract interaction) (a) and soluble pellet (soluble part of the precipitate formed after enzyme extract interaction) (b) after the interaction of plant extract and trypsin at 4 °C for 60 hr.

Sample preparation for FT-IR spectroscopy

The *E. arvense*-trypsin mixture was lyophilized in a freeze drier (Telstar-Cryodos) for overnight. The powder was mixed with dry potassium bromide (KBr) (Sigma-Aldrich, USA) in a mortar (at a ratio of 1:100). The mixture was then pressed to 100 kg/cm² (1200 psi) for 5 minutes to form a disk.

FTIR spectrum accumulation and data processing

The spectral analysis was carried out by using a Perkin-Elmer spectrophotometer equipped with MIR TGS detector (Spectrum 100 Instrument, Perkin Elmer Inc., Norwalk, CT, USA). FTIR spectra of the samples were recorded between 4000 and 450 cm⁻¹ region. The background spectrum was subtracted from the spectra of the samples automatically. Spectrum 100 software (Perkin Elmer) was used for all of the data processing. From each sample, at least three different scans, which gave identical spectra, were performed. Smoothing of the spectra was done by using Savitzky-Golay algorithm. Then, the spectra were interactively baselined from two arbitrarily selected points. Finally, the spectra were normalized in specific regions for visual comparison of the samples.

Determination of phenolic content

Total phenolic content of *E. arvense* extract and trypsin *E. arvense* mixture was determined by Folin-ciocalteu

method. Folin-ciocalteu reagent was prepared by 1:10 dilution of stock solution. Sodium carbonate solution of 7% was prepared in distilled water. Gallic acid was used as standard in the calibration curve. 20 μ l of each sample was mixed with 100 μ l Folin-ciocalteu reagent and incubated for 2.5 minutes. Then 80 μ l of sodium carbonate solution was added. The mixture was kept in dark for 1 hour. Samples were subjected to photometric measurement at 725 nm. Results were expressed as mg of gallic acid equivalents (GAE)/gr dry weight extract.

HPLC analysis

High-pressure liquid chromatography (HPLC) analysis was modified from the studies of Canadanovic-Brunet et al [13]. HPLC was performed with an Agilent 1100 series device equipped with diode array detector. A reversed-phase column, Lichrospher 100- RP 18 with a 5- μ m particle size (Agilent Technologies, USA), was used at the flow rate of 0.8 mL min⁻¹. Mobile phase gradient was performed by varying the proportion of solvent A (2.5% acetic acid) to solvent B (100% acetonitrile) as follows: initial 1% B; linear gradient to 40% B in 40 minute. The injected sample volume was 20 μ L All solutions were filtered prior to injection through 0.20 μ m membrane filters (Millipore, Bedford, MA, USA). The column temperature was 35 °C. The measurements were performed at 280 nm.

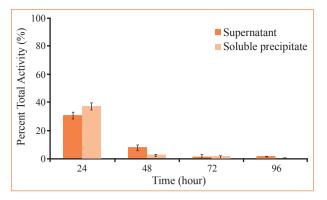


Figure 3. Percent total activity of supernatant and soluble pellet after the interaction of plant extract and trypsin at different time intervals at 24 °C. In the assays 1 mg/ml *E. arvense* extract and 0.05 mg/ml trypsin enzyme were used.

Results and Discussion

In the present study the interaction between trypsin and *E. arvense* extract was studied using enzyme assays for protein functionality and structural changes.

Trypsin stability studies

The effect of the interaction on trypsin functionality was tested using trypsin enzyme assay. The percent total activities of both the supernatant (aqueous part formed after enzyme extract interaction) and the soluble pellet (soluble part of the precipitate formed after enzyme extract interaction) at 24°C were shown in Figure 1. The solutions with various concentrations of E. arvense were mixed with trypsin solutions and incubated for 60 hours at 24°C. The final concentrations in the mixture were 0.01; 0.1 and 1 mg/ml for E. arvense extract and 0.05 mg/ml for trypsin. Trypsin solution with a concentration of 0.05 mg/ml was also incubated with distilled water at the same conditions in order to use as a control. The original trypsin total activity was taken as 100% and the other mixtures total activities were calculated compared to two this original trypsin total activity. As shown in Figure 1 the supernatant's percent total activity decreased drastically as the concentration of E. arvense extract increased in the mixture. The percent total activity of soluble pellets of mixture, on the other hand were less than 0.2.

In Figure 2 the percent total activity of both supernatant and soluble pellet at 4°C was shown. Solutions with various concentrations of *E. arvense* were mixed with trypsin solution and incubated for 60 hours at 4°C. The final concentrations in the mixture were 0.01; 0.1 and 1 mg/ml for *E. arvense* extract and 0.05 mg/ml for trypsin solution. Trypsin solution with a concentration of 0.05 mg/ml was also incubated with distilled water at the same conditions as a control. As shown in Figure 2, the supernatant percent total activity decreased slightly as the concentration of *E. arvense* extract increased in the mixture. The percent total activity of soluble pellets of mixture, on the other hand were less than 0.75%. In Figure 3 percent total activity of both supernatant and soluble pellet at 24 °C were shown. Solution with fixed concentration of E. arvense were mixed with trypsin solution and incubated for different time intervals at 24 °C. The final solution concentrations in the mixture were, 1 mg/ml for E. arvense extract and 0.05 mg/ml for trypsin. The time intervals for the incubation were 24, 48, 72 and 96 hour. Trypsin solution with a concentration of 0.05 mg/ml was also incubated with distilled water at the same conditions as a control. Trypsin total activities for control at different time intervals were taken as 100% and the other mixtures total activities were calculated depending up this control reference. As shown in Figure 3 the supernatant percent total activity decreased drastically as the incubation time of *E. arvense* and trypsin increased. The same results were seen for the soluble pellet as well.

In all of the experiments a white-colored precipitate phase formation was observed. Therefore the activity loss was studied in both the precipitate and supernatant phases. It is assumed that phenolic compounds in the extract interact with the enzyme and this interaction leads to precipitate formation. As the temperature of incubation decreased to 4 °C from the room temperature the amount of precipitate that formed due to this interaction also decreased. The results showed that the decrease in the total percent activity was relevant to the amount of precipitate that forms. The experimental result indicated that at higher incubation temperatures the enzyme inactivation kinetics was enhanced in addition to the amount of precipitate formation also increases with increasing temperature. The total percent activity decreased to 5% at room temperature where as it decreased to 40% at 4 °C. The amount of precipitate formed at room temperature was more than that at 4 °C.

The results also indicated that in the precipitate phase, almost no enzyme activity was detected. This might be due to the interaction between the enzyme and the compounds present in the extract materials. As the inactivation kinetics results indicated that the total enzyme activity in the supernatant phase decreased over time due to the irreversible loss of enzyme activity. However in the precipitate phase the enzyme activity was much less than that in the supernatant phase (at room temperature total activity of supernatant was 5% whereas at 4 °C total activity of soluble precipitate was 0.25%).

The phenolic content of *E. arvense* was reported to contain caffeic acid, syringic acid, ferulic acid, vanilic acid, rutin, apigenin, epicatechin and kaempferol [13]. Interaction of these phenolics with several proteins has been demonstrated. The interaction between caffeic acid and human serum albumin was demonstrated [19]. Investigations on rutin revealed that rutin could bind to trypsin enzyme via hydrophobic interactions and it could also bind to plasma proteins. As a result of this binding, antioxidant capacity of rutin was reported to decrease drastically [20,21]. Kaempferol was also shown to bind to bovine

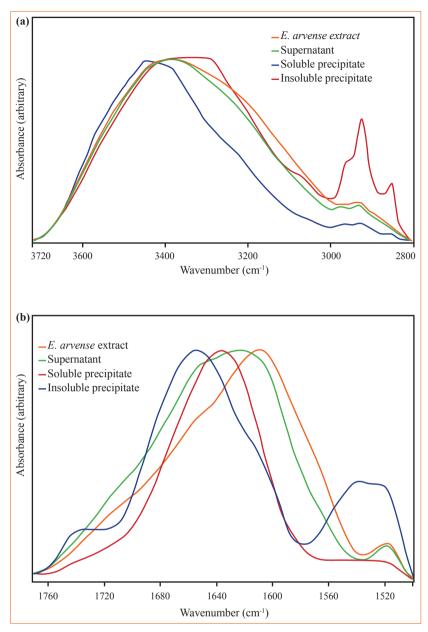


Figure 4. The FTIR analyses of the *E. arvense* extract and trypsin mixture in the wavenumber ranges of (a) 3720-2800 cm⁻¹ and (b) 1780-1500 cm⁻¹.

serum albumin strongly by hydrophobic interactions and this resulted in changes in secondary structure of protein [22]. The enhanced enzyme activity due the higher temperatures should also increase the amount of precipitate formed. Since it was known that the hydrophobic interactions increased with temperature [23], the results can be explained in the light of hydrophobic interactions between flavonoids in the plant extract and the enzyme, causing more protein inactivation and enhanced precipitate formation as well. Rawel *et al.* (2002) [24] showed that the phenolic compounds could interact with soy proteins and change their structure by increasing hydroxyl substituents. Their study also showed that these interactions could occur at temperatures as low as 4 °C. The phenolic compounds they studied were caffeic acid, kaempferol, quercetin and apigenin. They found that these substances had high affinity to proteins. As a result of this interaction these compounds were found to form precipitates [24].

The phenolic groups that were found in plant extract can form hydrogen bonds with the polar groups (i.e. amide, guanidine, peptide, amino and carboxyl groups) of protein or the hydrophobic amino acid such as proline, tyrosine and phenylalanine can interact and bind strongly to the polyphenolic substances [25,26].

It was also reported that not only hydrogen bonding but also hydrophobic interactions between enzyme and polyphenols could cause enzyme inactivation. Flavonoids especially kaempferol, were found to have inhibitory effects on trypsin as a result of their hydroxyl groups [27]. In

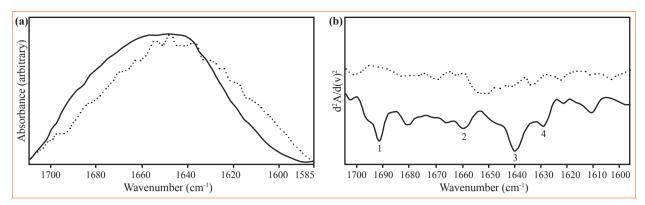


Figure 5. The average (a) absorbance and (b) second derivative spectra of original trypsin, trypsin in the supernatant of trypsin-*E. arvense* extract in 1700-1600 cm⁻¹ region (continuous line shows original trypsin, dashed line shows trypsin in trypsin-*E. arvense* extract).

2004 Maliar et al. [28] concluded that flavonoids could be used as potential trypsin inhibitors due to their hydrophobic nature. They concluded that quercetin and apigenin were the most effective phenolics for the inhibition of trypsin enzyme activity [29]. In light of these findings and the knowledge of E. arvense's rich content of flavonoids and phenolic compounds [8] especially apigenin and quercetin, it is possible that these compounds can easily interact with the trypsin enzyme and cause to decrease the total percent activity. Moreover Hirano et al. (1992) [25] showed that plant materials in general could release amino acids into surrounding solution which could have inhibitory effects on trypsin enzyme as well. In our research 70% ethanol was used for extraction, during extraction procedure polypeptides could easily be transferred to aqueous solution. These amino acids can possibly interact with trypsin enzyme and cause enzyme inactivation. The conformational change of trypsin was irreversible as the nature of the assay in this research.

Silicic acid is the major inorganic substance in the plant's chemical composition. The investigations of this acid revealed that as the pH of the environment reached to the proteins isoelectric points, proteins interact with silicic acid by possible adsorption to form complexes called amine silicate via their histidine and amino groups [30].

FTIR analyses

FTIR analyses were carried out to study the effect of the interaction between the extract and trypsin on the structure of the extract. The spectra were analyzed in 3720-2800 cm⁻¹ and 1780-1500 cm⁻¹ regions. The results are shown in Figure 4.

The FTIR studies indicated significant changes in the protein structures and lipid partitioning in the process. Since trypsin concentration kept at a lower level, the spectra obtained represent only the signals originating from the extract material. The spectra were analyzed in two different spectral frames: 3720-2800 cm⁻¹ regions and 1780-1500 cm⁻¹ region. As seen in Figure 4 a, the interaction between trypsin and the extract affects the structure-composition of the extract components. Since the spectral region between 3720 and 3000 cm⁻¹ corresponds to amide-A region stemming from proteins both the band shifting and the shapes of the amide A bands indicate both structural-compositional changes in proteins and also possible changes in the hydrogen bonding network in the proteins or between the proteins and the other extract materials. In the same figure the region between 3000 cm⁻¹ and 2800 cm⁻¹ were heavily dominated by bands originating from lipids. A striking feature of the spectra was that the lipid signals came mostly from insoluble pellets (insoluble part of the precipitate formed after enzyme extract interaction) indicating the accumulating of lipids originating from the plant material in the insoluble pellet part. The same features of the materials were also observed in Figure 4 b. The increase in the Amide II band intensity around 1550 cm⁻¹ indicates the increase in the hydrogen bond network as well in the insoluble pellet.

An FTIR analysis was carried out to understand the nature of the decrease in the enzymatic activity to be able to distinguish any possible denaturation from enzyme activity inhibition. For this purpose a spectral difference analysis was performed. Simply, we subtracted the spectrum of the supernatant samples from E. arvense extract spectrum and obtained the spectrum of trypsin in the mixture and then compared it with the original trypsin spectrum using normalization procedures. The spectra were quite different from each other as seen in Figure 5 a. The trypsin spectrum was found to lose many secondary structural components as opposed to the original trypsin spectrum (Figure 5 b). The major sub bands that showed differences were that at 1694 cm⁻¹ (1:antiparallel β -sheets), at 1658 cm⁻¹ (2: α -helix), at 1637 cm⁻¹ (3: β -sheets), and at 1627 cm⁻¹ (4:aggregated β -sheets) [31].

Therefore we concluded that the interaction between *E. arvense* extract and trypsin in the supernatant samples resulted in denaturation of trypsin.

Phenolic content studies

Literature reviews showed that phenolics had a capacity to

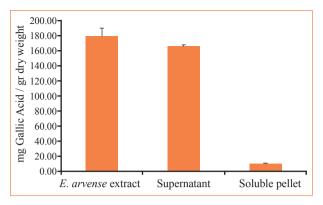


Figure 6. Total phenolic content of initial *E. arvense* extract, supernatant after 60 hr treatment of extract with trypsin, soluble pellet of extract after 60 hr treatment of trypsin.

interact and precipitate enzymes and proteins [24]. In this research a white precipitate was seen in all the samples of extracts treated with trypsin. Also the inhibition of enzymatic activity increased with the increasing concentration

of the plant extract. In order to understand whether this inhibition occurs as a result of phenolics enzyme interaction or not, the total phenolic content of extracts before and after treatment with enzyme was analyzed and the results are presented in Figure 6.

Total phenolic content of the initial *E. arvense* extract, supernatant after 60 hr treatment of extract with trypsin, and soluble pellet of the extract after 60 hr treatment of trypsin are shown in Figure 6. Both of the supernatant and *E. arvense* sample are prepared from 1 mg/ml plant extract. As a result of this the total phenolic content is given as the milligram gallic acid equivalent/g. dry weight of extract. It is seen that when the total phenolic contents of the supernatant and the soluble pellet are considered they are equal to the phenolic content of *E. arvense* extract. Most of the phenolics remain in the supernatant and some are in the soluble precipitate phase reversibly. The high amount of phenolics in the supernatant would be the reason of trypsin inhibition. The phenolics could interact with proteins and force them to precipitate. Also the presence of

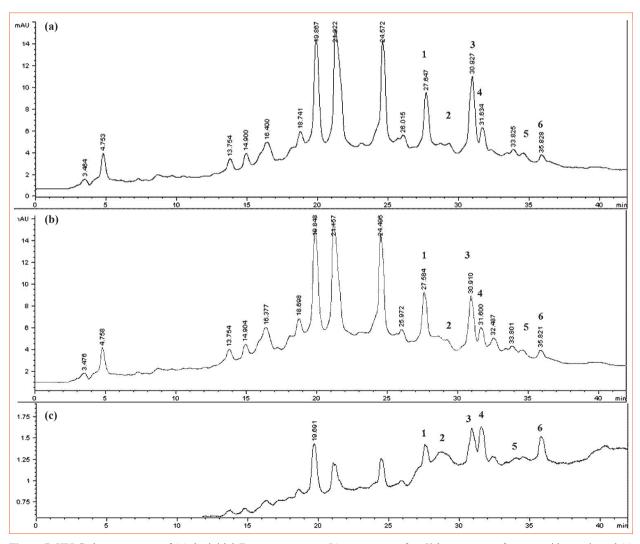


Figure 7. HPLC chromatograms of (a) the initial *E. arvense* extract, (b) supernatant after 60 hr treatment of extract with trypsin and (c) soluble pellet of extract after 60 hr treatment of trypsin at 280 nm.

Table 1.	Change in the peak areas in HPLC chromatograms of		
	initial E. arvense extract and supernatant after 60 hr		
	treatment of extract with trypsin and soluble pellet of		
	extract treated 60 hr with trypsin		

	Peak areas	
Peak number	E. arvense	Supernatant
1	546.9	517.14
2	400.78	393.38
3	515.35	400.64
4	190.42	177.55
5	148.6	122.58
6	230.53	179.56

phenolics in the soluble precipitate would be the evidence of interaction of phenolic compounds with proteins.

HPLC analysis

In order to confirm the previous results HPLC analyzes of the initial E. arvense extract, supernatant after treatment of extract with trypsin, soluble pellet of extract after treatment of trypsin is performed. The peak areas are determined and the differences are reported as shown in Figure 7.

In Figure 7, the HPLC chromatograms of initial E. arvense extract, supernatant after 60 hr treatment of extract with trypsin, soluble pellet of extract after 60 hr treatment of trypsin are shown. When the peak areas are analyzed it was seen that there is a decrease in the amounts of compounds found in the apolar region of the chromatograms upon precipitation. The amount of compounds labeled on the chromatogram from 1 to 6 changes between E. arvense extract and supernatant. The amount of decrease is shown on Table 1. The peaks of corresponding number could be seen on the chromatogram of the soluble pellet. As a result of protein phenolic interaction precipitation occurs. According to HPLC results of supernatant, it can be seen that there is a decrease in the phenolic content of *E. arvense* after treating with trypsin. Those findings are strong proofs for protein phenolic compound interaction.

Due to use of *E. arvense* as an alternative medicinal plant and its proven antioxidant, antimicrobial and anti-inflammatory effects, the effects of *E. arvense* extract on trypsin enzyme is investigated in this study. According to the enzyme kinetic assay, enzyme inhibition assay and FTIR analyses; it is seen that *E. arvense* extract has an inhibitory effect on trypsin activity. In addition, the interaction with trypin causes the extract material to precipitate out. This interaction resulted in accumulation of the extract lipids in the insoluble precipitate phase. As a result of the FTIR findings we concluded that the inhibitory effect of plant extract on trypsin enzyme was due to denaturation of enzyme. Also the enzyme kinetic analyses on both pellet form and supernatant form showed that the denaturation after interaction between *E. arvense* extract and trypsin was irreversible. In addition it was observed that the interaction between the extract and trypsin caused protein precipitation. As a conclusion, the interaction between the extract and trypsin is due to enzyme denaturation.

Antitrypsin compounds are used in agriculture as pesticides, in food industry [32,33]. Therefore *E. arvense* extract can be used as a potential antitrypsin compound.

Acknowledgements

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Conflict of Interest

There are no conflicts of interest among the authors.

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