

# ISOLATION AND PURIFICATION OF $\alpha_{s_2}$ -CN LOCAL GOAT MILK AND STUDYING ITS HYDROLYSATES INHIBITION EFFICACY TOWARDS ACE1.

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#### Abstract

Hypertension is a universally common chronic disease and is called Silent Killer Current trend is to discover natural alternatives instead of chemical medications to treat this disease. In this study hydrolysates from goat milk casein have been used as natural materials to inhibit activity of Angiotensin Converting Enzyme1 (ACE1) the enzyme which is responsible on hypertension.  $\alpha_{s2}$ -Casein from acidic local goat milk casein has been purified utilizing anion exchanger DEAE-Cellulose then by gel chromatography using sephadex G-75, electrophoresis technique SDS-PAGE Proven existence one pure type of  $\alpha_s$ -CN which has been appeared as one band and has mw 25.620 KD.

Purified  $\alpha_{s2}$ -CN has been hydrolyzed by pepsin, trypsin and mix them by 1:1. The hydrolysates appeared dissimilar ACE1 inhibition activity but highest rate 64.019 % and highest degree of hydrolysis (DH) 61.924 with hydrolysate produced by mix of tow enzymes after 8 hours of hydrolysis  $\alpha$ H8M compared with inhibition activity by pepsin  $\alpha$ H8P and trypsin  $\alpha$ H8T after 8 hour too which were 54.146, 50.033 % and DH 48.293, 37.753 respectively. Inhibition activity towards ACE1 is directly proportional with DH value and hydrolysis period.

*Key words*:  $\alpha_{s_2}$ -CN, Goat milk, ACE1

#### Introduction

According to last statistics of World Health Organization WHO about 17.1 million person around the world are infected with Hypertension yearly. There are many medications and chemical therapies treat this disease most of them focus on inhibit producing the responsible article of hypertension referred to Angiotensin2 (Ang2) which converted from Ang1 by activity of ACE1 in humans and experimental animals (Sica, 2005).

Chemical angiotensin converting enzyme1 inhibitors  $(ACE_1I_s)$  have side effects in some body organs like kidney, lungs and liver like Symptoms of dry cough caused by tightness of the Pulmonary vesicles and angioedema (Guyer and Banerji, 2015). At recent days the world turned to natural materials to treat many diseases including high blood pressure. The world has turned to the use of natural materials to treat a number of diseases, including high blood pressure, most of which focus on inhibiting the effectiveness of ACE1 within Renin-Angiotensin-system (RAS) (AL-khafaji, 2008; Laragh, 2001).

ACE (EC3.4.15:1) is a dicarboxypeptidase type and metalloenzyme it's active side containing zinc ion and there is tow forms of this enzyme ACE1 and ACE2, the difference between them in stimulation sites. The main ACE1 function is cutting dipeptide (HIS-LEU) from carboxylic side of peptidyl hormone Ang1 consisting of 10 amino acids to convert it to Ang2 which is consider vasoconstrictor hormone, it presents in all blood vessels and it affects in secretion aldosterone hormone in kidney which causes hypertension so that referred to this regulatory role system "Renin- Angiotensin- Aldosterone System" (RAAS), ACE1 found in several human body organs but its highest concentration in pulmonary tissue (Aluko, 2018).

In several studies it was noted that in 1960 the English scientist Sir John Vane discovered a peptide from the Brazilian arrowhead viper, this peptide has the ability to rapidly lower blood pressure. It was later concluded that this peptide inhibits the ACE and this inhibition is fiercely competitive, this led to make an inhibitors similar to but non-peptide synthetic inhibitor and the first medication was CAPTOPRIL which contains sulfhydryl group and many other medications but all of them have side effects (Brown and Hall, 2005; Wong *et al.*, 2004).

The peptide research field is one of the most important research fields of preventive medical chemistry since there have been medical evidence that the lack of peptide leads to hypotension and the effectiveness of reduced blood pressure, this led to the important question that the effect comes from inhibition of ACE1 only as chemical drugs inhibit or inhibition enzymes in and the answer was "Mostly with a direct and obvious effect in ACE1" Globally, there are many products containing hypotensive peptides in the commercial markets and are also included in the pharmaceutical formulations, mainly sesame peptides, seaweed peptides, peanuts, and sour milk milk. Most of these products, if not all, inhibit the effectiveness of ACE, other useful functions of the human and animal organs, most notably antioxidant and antimicrobial peptides, these properties are often found in the bioactive peptides (Shi et al., 2014; Aluko, 2012).

Biologically and physically, the bioactive peptides are a sequence of amino acids which are affiliated within the protein and has positive functional and therapeutic properties within the human body but these properties appears only when it is released from the original protein by several hydrolysis ways like lipolysis enzymes of digestive tract and food fermentation by bacteria especially lactic acid bacteria (LAB) or exposing proteins or their isolates are to single or mixed proteolysis enzymes extracted from microbial, animal or plant sources (Munn, 2013; Korhonen and Pihlanto, 2007). Milk Protein one of the proteins which can be released with distinct hypotensive peptides and that peptides can have more than one bioactive function type of these peptides depend on the composition of each type of proteins and its primary structure, which determines the extent of resistance to the proteolysis enzymes of protein, especially enzymes that work in digestive tract. (Ndiaye et al., 2012). The biological effectiveness of low molecular weight molecules is based on the type and sequence of the amino acids forming the peptide sequence. Bioactive peptides usually contain a number of acids ranging from 3 to 20 amino acids per molecule (Meisel and Fitz, 2003).

Studies on the effect of peptides and fermentation factors from goats' milk that have inhibitory effect on ACE1 are rare and are limited the most are focus on fermentation processes using different strains and Evaluation their inhibition efficacy on ACE1 and reducing blood pressure. In an experiment on this subject, 28 strains of lactic acid bacteria were used to ferment goat's milk, the *in vitro* results of the experiment showed that there were 20 types of ACE1 efficacy inhibitors, but only four showed very high efficacy in ACE1 inhibition rates by probiotic *Lactobacillus* .spp were 95.92, 84.61, 82.79, and 78.57%. Another study the ACE1 inhibition activity was 91.6% by goat milk fermentation with *lactobacillus planetarium* 691 (Chen *et al.*, 2018; Chen *et al.*, 2012).

Lee *et al.*, (2005) mentioned that a Little studies can be found on ACE-inhibition from proteolytic hydrolysates derived from casein (CN) in goat's milk, his experiment showed that goat casein hydrolyzed by pepsin and trypsin show up high ACE1 inhibition efficacy which were 67.1, 61% respectively and this activity increases with the degree of hydrolysis (DH) which were 48.0 and 59.5 ¡conclusion that isolation peptides from its hydrolysates cause increasing in ACE1 inhibition activity to 97.6%. Val-Pro-Pro and Ile-Pro-Pro are the most successful ACE1 inhibition activity peptides resistant to the digestive enzymes so that can be consumed orally. They are originally derived from the hydrolysis of  $\beta$ -CN in cow's milk and proved to be found in various types of solid, semi-solid and soft cheeses (Rutella *et al.*, 2016).

Many other researches proved ACE1 inhibition activity of hydrolyzed caseins and whey protein by digestive enzymes in addition to systolic blood pressure reduction in spontaneously hypertensive rats (SHR) comparing with treatment by chemical drug (captopril) (Sun et al., 2014; Costa et al., 2005). Many researchers, including (Aslam et al., 2018; Bansal and Bhandari, 2016), suggest that the active peptides resulting from the hydrolysis of pure  $\alpha$ ,  $\beta$  and  $\kappa$ -CN and have molecular weights lower than the molecular weight of the peptides resulting from the hydrolysis of complete casein molecules when used Pepsin, trypsin and chymotrypsin with a difference of up to 85% in  $\beta$ -CN and 75% in  $\alpha_s$ -CN. This is important in the production of bioactive peptides, especially antioxidants and inhibitors of ACE1. RP- HPLC is very effective in separating peptides containing hydrophilic amino acids. The results were also observed with the hydrolysis of the whey protein compared to hydrolysis of its previously separated fractions by chromatographic methods and by the use of the same enzymes.

#### **Materials and Methods**

#### **Enzymes and reagents**

Fresh local Iraqi goat milk purchased from animal wealth directorate-ministry of Agriculture\Baghdad-Iraq defatted by using the BL-S selecta centrifugal device at 3000×g for 15 minutes and at 4°C. The separation of goat's milk acidic casein was performed according to the methods of AL-saadi, (2001). Trypsin (Porcine Source, activity 1800 u/gm, Merk company) and pepsin

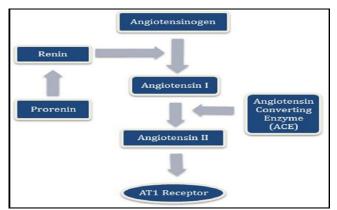


Fig. 1: Schema illustrates the role of ACE1 in Ang2 formation within RAS (White *et al.*, 2015).

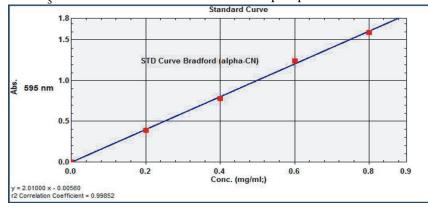
(Porcine Source, activity 2000u/gm, BDH company). Standard bovine  $\alpha_s$ -CN was from Merk Company. Sephadex G-75 Fine45 was from BDH Company. TNBS from Fluka Company. ACE1 (EC 3.4.15.1) From Rabbit Lung and Hippuric acid HA.

#### α<sub>s</sub>-CN Separation

Method of Zittle *et al.*, (1959) followed up for  $\alpha_s$ -CN separation from acidic casein with modification in acidic casein quantity which used in the beginning of the separation procedure 40gm of acidic casein previously prepared in 100 ml of urea 6.6 molar solution with stirring. Add 42.5 ml of distilled water to the solution to obtain 4.63 molar of urea solution and leave it at room temperature until forming white Precipitate which dissolved in 100ml of urea concentration to 4.63molar, the precipitate composed is  $\alpha_s$ -CN which has been kept in freeze drying.

### $\alpha_s$ - Casein Purification by Ion Exchange Chromatography

Anion exchange chromatography has been used to purify crude  $\alpha_s$ -CN which separated from caseins previously according to Whitney, (1988). 1gm of freeze dried  $\alpha_s$ -CN has been dissolve in 100ml sodium phosphate



**Fig. 2:** αs-CN Standard curve of Bradford method used to estimate protein concentrations. The absorbance on 595 nm.

buffer 0.1molar and pH 7.4 containing urea 3.3 molar and mercaptoethanol 0.010 molar then mix with DEAE-Cellulose for 15 min. Aon exchange mix poured in column with dimensions of 2×50cm. The column washed by buffer solution conjugated caseins with column and  $\alpha_s$ -CN Removed from column by using phosphate buffer which was used to dissolve sample containing NaCl in rising concentrations 0.1, 0.175, 0.2, 0.25 molar. Flow rate was 50ml/hr., 5ml/tube using fraction collector proteins solutions in tubes was read by spectrophotometer Shimadzu model 1650 PC on 280nm.

#### $\alpha_s$ - Casein Purification by Gel Filtration

Sephadex G-75 gel has been used to prepare column according to Pharmacia fine chemicals company. 0.5gm of freeze dried  $\alpha_s$ -CN which purified initially by previous step dissolved in 10ml 0.005 molar phosphate buffer. Sephadex column with dimension 1.570×1 cm balanced by the same buffer containing urea 6.6 molar and EDTA 0.004 molar. Fractions has been recovered from Sephadex column with flow rate 20ml /hr. and 3ml /tube and proteins solutions in tubes read by spectrophotometer Shimadzu model 1650 PC on 280 nm. Total volume of  $\alpha_s$ -CN was concentrated and protein concentration estimated by Bradford method.

#### Protein concentration assay

Bradford method was used to estimate  $\alpha_s$ -CN concentration in solutions by using standard  $\alpha_s$ -CN in standard curve preparation.

### Confirm Purity and Estimate the Molecular Weight of $\alpha_s$ -CN.

To confirm the purity of  $\alpha$ s-casein and determine its molecular weight, Method of electrophoresis with polyacrylamide gel was used in the presence of Sodium Dodecyl Sulphate (SDS) by using a vertical electrophoresis device comparing with standard bovine Known molecular weight  $\alpha_s$ -CN based on the method

described by Laemmli, (1970) and modified by Schägger (2006).

### Hydrolysis of $\alpha_s$ -Casein by Proteolytic enzymes.

The method of Adamson and Reynolds, (1996) was followed with the modifications of Lee *et al.*, (2005) in the amount of the protein and the number of enzymatic proteolytic units in proportion to the amount of protein obtained from the purification steps. Weight of  $0.1 \text{gm} \alpha_s$ -CN dissolved in 10 ml distilled water with pH 2 for pepsin

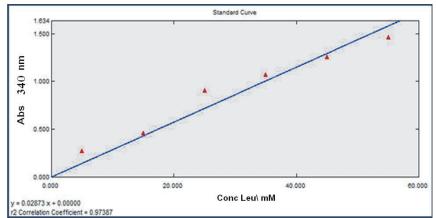


Fig. 3: Standard curve of Leucine used for DH%. The absorbance at 340 nm.

and pH 8 for trypsin and pH 2 then raising to pH 8 for mix of pepsin and trypsin in the rate of 1: 1. Changing pH by adding Hcl and NaoH. Pepsin and trypsin was added to reaction mix as 20, 18 unit respectively, reaction time was 8hr., at 37°C. Reaction time of symbiotic enzymes was 4hr., to pepsin and then 4hr., to trypsin. Samples pulled each hour to determine degree of hydrolysis DH.

#### Estimation of Degree hydrolysis (DH)

Degree of hydrolysis was estimated according to Liu and Chian (2008). Standard curve of amino acid leucine (Fig. 2) was used to calculate DH of samples. Volume of 0.250ml of samples for each time was transferred to test tube and mix with sodium phosphate buffer 0.2125 molar with pH=8.2 and 2ml of 0.1% Tri nitrobenzene sulfuric acid (TNBS) was added to reaction mix which incubated at 50°C for 1 hr. Isolated from light then reaction stopped by adding 0.1 molar Hcl and were left to cool to  $30\pm2°$ C for 30 min. The absorbance was measured at 340 nm. The degree of hydrolysis (peptides concentration or Peripheral NH3 groups) was estimated by entering the equation below Referred to Jamdar *et al.*, (2010) using UV- PROBE 2.1 program by shimadzu device SP:

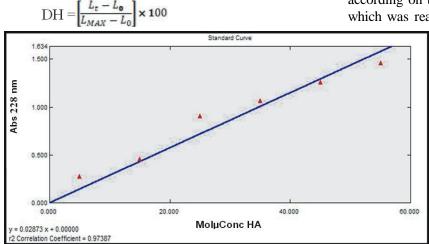


Fig. 4: Standard curve of hippuric acid used to estimate HA concentrates in reaction mixture. The absorbance at 228 nm.

So that

 $L_t$  = concentration of Peripheral NH3 groups at time (0-8 hr.).

 $L_0$  = Free amino acids (NH3 group) in original protein samples without any treatment.

#### L<sub>MAX=total</sub>

 $L_{MAX}$  = Total free amino acids resulting from acidic hydrolysis by 6 molar Hcl at 120cp for 24 hr.

### Determination of ACE1 Inhibition Activity by $\alpha$ -Casein Hydrolysates.

The efficacy of  $\alpha_{s2}$ -CN hydrolysates at each hydrolysis degree in ACE1 inhibition was estimated by spectroscopic method based on the conversion of N-Hippuryl-His-Leu hydrate peptide to Hippuric acid (Benzoylaminoacetic acid) as indicated by ACE1 according to Cushman and Cheung, (1971), modificated by (Chen *et al.*, 2018; Lee *et al.*, 2005).

Volume of 100  $\mu$ l of  $\alpha_{s2}$ -CN hydrolysate was incubated for three periods (beginning, middle and end) of enzymatic hydrolysis for pepsin and trypsin and 1:1 mixture of the two enzymes with 200 µl of Substrate HHL solution at 37°C for 5 minutes using water bath. 20 µL of ACE1 solution 0.2 unit/ ml was added and the reaction mixture incubation at the same temperature for 30 minutes and the enzymatic reaction was stopped using 0.25 ml HCl 1 molar. The amount of hippuric acid produced from the reaction was extracted by adding 3 ml of ethyl acetate at 100°C using a rotary evaporator. 3 ml of distilled water was added to the remaining volume. The absorbance of hippuric acid determined at 228 nm the quantities of HA acid released from HHL estimated according on the standard curve of hippuric acid fig. 3 which was read in the same apparatus. The amount of

HA released in the absence of proteolytic enzyme was also considered as 100% ACE1 and the equation (A-B) /  $A \times 100\%$  was used to calculate the enzymatic activity as a percentage: % inhibition = ((B-A) / A) × 100:

A = concentration of hippuric acid =  $234.036 \mu mol$ , in the absence of any factor that inhibits the effectiveness of ACE1 (100% effectiveness)

B = concentration of hippuric acid in the presence of ACE1 inhibitors, which are peptides under study.

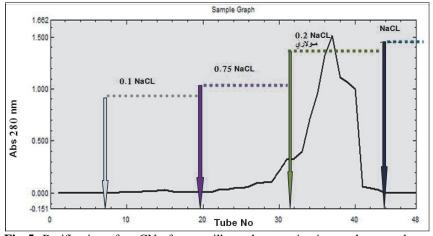


Fig. 5: Purification of  $\alpha_s$ -CN of goat milk on the negative ion exchange column DAEA-Cellulose with dimensions of 2×50 cm using a phosphate buffer 0.1 molar pH 7.4 containing urea solution 3.3 mL and 10 mmol mercaptoethanol and runoff 50 ml / h at a rate of 5 ml / tube and salty gradient from 0.1 molar to 0.25 molar sodium chloride in the buffer.

#### **Results and Discussion**

#### Acidic casein yield

The results showed that the percentage of casein obtained after acidic Sedimentation of Iraqi goat milk 2.6%. Very similar to that found by Yangilar, (2013), which set a minimum protein content in goat milk for a number of breeds (3.4%) and total casein (2.4%). The highest protein content in Spanish goat milk is 4.6%, determined by Park *et al.*, (2007). The variance in casein content due to genetic differences between species as well as other such as nutrition, the number of births and other factors that lead to changes in total protein ratios.

#### Primary $\alpha_s$ - Casein Purification

Method of Zittle et al., (1959) was followed up to

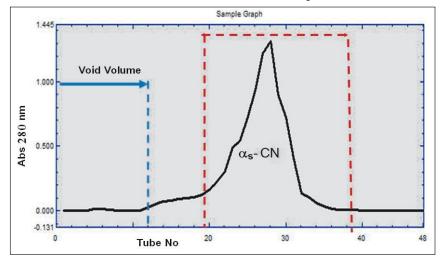


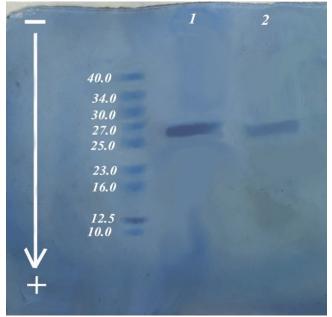
Fig. 6: Final Purification of  $\alpha_s$ -CN goat milk using sephadex column G-75 column with dimensions of 90 × 1.5 cm and phosphate buffer solution 0.005 molar with pH 7.6 containing urea 6.6 molar and EDTA 0.004 molar with 20 ml/h flow rate, adjust fraction collector to collect 3 ml volume per tube.

separate  $\alpha_s$ -CN before purification by chromatographic methods. The yield of  $\alpha_s$ -CN by using method of precipitation with salts and urea was 9.2 gm produced from 40gm of acidic casein. This quantity forms 23% and this results are compatible with the amount of  $\alpha_s$ -CN obtained by Zittle *et al.*, (1959) when using 30 g of casein to be the result of  $\alpha$ s-CN is 7gm, which constitutes a rate of 23.3% CN is possible when the amount of acidic casein increases, but the increase is not 100% straight, It's also compatible with Bramanti *et al.*, (2003) in the rate of 22.4%.

### Purification of $\alpha_s$ -Casein by Ion Exchange Method

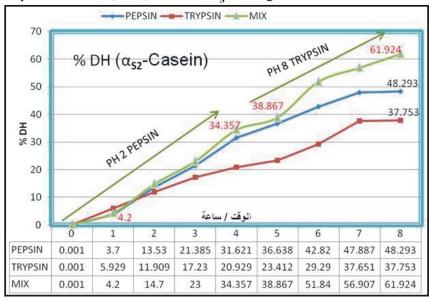
The first step of the purification process was aimed at eliminating the proteins associated with the  $\alpha_s$ -case by using Sodium chloride salt solution with different molar concentrations of 0.1, 0.175, 0.2 and 0.275 molar, depending on the and  $\alpha_{s1}$ -CN, if present in very small quantities, can be separated from  $\alpha_{s_2}$ -CN, which is present in large quantities in goat's milk, especially if  $\alpha_s$ -CN and  $\beta$ -CN is crudely used when the urea solution and mercaptoethanol which contribute to increasing the solubility of the protein and decoding bi-sulfur bonds link (Al-Saadi, 2001). The Wei and Whitney, (1985) and Whitney, (1988) methods were used for the initial purification of  $\alpha_s$ -CN from the initial separated goat milk acid casein by push through on the DEAE-Cellulose column and to recover it and disengage from this exchanger, different salt concentrations of NaCl were

> used. Total casein contains many types of case  $\alpha$ ,  $\beta$ , and  $\kappa$ -CN, each protein differs from the other parts in its negative net charge therefore recovered from DEAE-Cellulose ion exchange column by using concentrations of sodium chloride solution from 0-0.1-0.175-0.2-0.25. fig. 4 shows first peak in the tubes (6-10). It is believed to have been traced back to K-CN because it is least negative casein and has weak bonding power to the ion exchanger, another peak of  $\beta$ -CN appeared in tubes (22-26) when using NaCl 0.175 M because it has net charge stronger than  $\kappa$ -CN and weaker than  $\alpha_s$ -CN. Also the figure shows protein  $\alpha_s$ -CN in major peak that appeared in the tubes (32-42)



**Fig. 7:** Migratr of purified proteins with standard protein and standard ladder with a range of 10-40KD in 12.5% SDS-PAGE. 1: represents the standard  $\alpha_s$ -CN and No. 2: represents the  $\alpha_s$ 2-CN purified goat's

because it is carrying negative charge stronger than other protein fractions so that it recovered from the column when using NaCl 0.2 molar. These results are consistent with those obtained by Wei and Whitney, (1985) and Greppi *et al.*, (2008) which indicated the separation of  $\kappa$ -CN in concentrations of less than 0.1 molar sodium chloride due to the low charge of this protein compared with other caseins as Saadi, (2018) confirmed weakness binding of  $\kappa$ -CN protein to negative ion exchange column as the weakest casein in both goat milk and cows. Ruprichova *et al.*, (2015) noted that  $\alpha_s$ -CN in goat milk;



**Fig. 8:** Hydrolysis Percentage of  $\alpha_s^2$ -Casein induced by pepsin and trypsin and their mixture at a ratio of 1: 1 during incubation for 8 hours on 37°C.

cows and sheep can be separated by using high salt when using HPLC. The total amount of  $\alpha_s$ -CN protein recovered was dried in a freeze-dried method for subsequent purification and the weight obtained was 910, 210 mg of 1 g weight in tubes (32-42), protein concentration rate was 16.569 mg / ml.

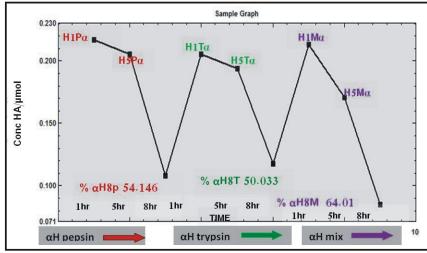
### Final Purification of $\alpha_s$ -CN by Gel Filtration Method by Using Sephadex G-75.

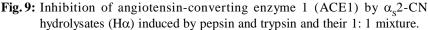
Casein contains many proteins,  $\alpha_s$ ,  $\beta$  and  $\kappa$ -CN,  $\alpha_s$ -CN is the largest of these parts. Although the ion exchange step may be enough to purify this protein, additional purification processes achieve a high degree of purity this is what Dosh, (2007) pointed out when she obtained high purity proteins using ion exchange and gel-filtration techniques, so this protein can be purified to a large extent by gel filtration. Fig. 4, shows appearance of a single major peak was in tubes (19-38) and these tubes belong to  $\alpha_s$ -CN. This result compatible with Ruprichova *et al.*, (2015) and Vincenzetti *et al.*, (2005) suggesting that the main peaks of the parts of the cows, sheep and goat milk cows are ion-exchanged and then gel-filtrated with the sephadex G-25 column or using RP-HPLC.

These results are in compatible with al-Saadi, (2001), when he obtained  $\alpha_s$ -CN by ion exchange method only without resorting to gel filtration and proved the purity of this protein later when conducting the electrophoresis of this protein in the gel acrylamide when he appeared in one band. Weights averaged 401.43 mg of 500 mg used at the beginning of the purification process and the protein concentration rate was 8.36 mg / ml.  $\alpha_s$ -CN was then concentrated, freeze dried and kept at freezing temperature.

## Confirm the Purity and Molecular Weight of $\alpha_s$ -CN by SDS-PAGE Method

Fig. 6, shows the appearance of single-purified  $\alpha_{s_2}$ -CN band, confirming that it is free from interference other casein fractions; having traveled approximately the same distance as standard proteins this confirms the purity of protein.  $\alpha_{s2}$ -CN is the predominant in goat's milk casein, as indicated by Moioli et al., (1998) and Anema et al., (1998). In contrast to cow's milk. It follows that the milk of this type of goat does not contain or contains very little  $\alpha_{s1}$ - CN protein. Compared with the molecular weight of the corresponding protein in Ladder, the molecular weight of pure  $\alpha_{s2}$ -CN was estimated about 25.620 KD.





This molecular weight corresponds to what Raak *et al.*, (2018) found when SDS-PAGE casein was migrated and found it to be less than 26.8 KD. The differences in the molecular weights of the fractions of the caseins in the milk of different species are mainly due to genetic factors.

### Hydrolysis by Enzymes Pepsin and Trypsin and Hydrolysis by Pepsin with Trypsin for $\alpha_{s_2}$ Casein.

Hydrolysis of  $\alpha_{s2}$ -CN casein was carried out using pepsin and trypsin, both of which were combined at a ratio of 1:1 for 8 hours. The results show in fig. 7 that the degree of hydrolysis DH increases with the incubation time of the two enzymes and their mixture. This is evidenced by the high absorption values along the wavelength of 340 nm this means increased release of peptides resulting from the specific hydrolysis action of the enzyme and the overall behavior of the three treatments is similar in that they reached the maximum hydrolysis after 8 hours of incubation at 37°C but in different proportions 48.293, 37.753, 61.924% with Pepsin and trypsin and their mixture respectively. These ratios are consistent with Lee et al., (2005) when treated of acidic casein from goat milk with multiple enzymes, including pepsin and trypsin indicates the availability of cyclic amino acids that form the alkaline for action of pepsin enzyme in  $\alpha_{s2}$  protein. As a result of the synergistic action of the two enzymes together on the DH of  $\alpha_{s2}$ -CN, the results indicated in fig. 7, that there is a clear increase in the values of DH was after 4 hours of incubation with pepsin enzyme is 34.357% and increased after incubation with trypsin another 4 hours to 61.924.

This result is consistent with Sun *et al.*, (2014) when using these enzymes together to hydrolyze  $\alpha_s$ -CN and concluded that synergistic action of enzyme contributes to obtain a high degree of hydrolysis as the initial hydrolysis by the action of pepsin

enzyme transforms the complex protein form into a free polypeptide due to the fact that the action of the enzyme pepsin is specialized in attacking peptide bonds due to the most Cyclic amino acids Tyr Trp and Phe as well as less hydrophobic amino acids and make the peptide chain more open which contributes to making the amino acid Arg and Lys more vulnerable and more ready to be attacked by trypsin.

### Effectiveness $ACE_1$ Inhibitory of $\alpha_{s2}$ -CN ( $\alpha$ H) Hydrolysates

Fig. 8, shows the effect of hydrolysis of three time periods resulting from the action of pepsin enzyme in  $\alpha_{s2}$ -CN resulting after 1, 5 and 8 hours of incubation with ACE1 ( $\alpha$ H1P,  $\alpha$ H5P and  $\alpha$ H8P) as well as three hydro of the same previous readings but by the enzyme trypsin in  $\alpha$ H1T,  $\alpha$ H5T and  $\alpha$ H8T and their indicated mixture  $\alpha$ H1M  $\alpha$ H5M and  $\alpha$ H8M on the effectiveness of ACE1, which is the concentration of hippuric acid HA released by the activity of ACE1 in substrate Hippuryl histidine leucine HHL. The diagram 8 shows also an inverse proportion between the HA concentration and the inhibition percentage and the inhibition rate was higher than 50% for the hydrolysates produced after 8 hours of 54.146, 50.033 and 64.019% by pepsin and trypsin at a ratio of 1:1 and their mixture respectively at a

**Table 1:** Relationship of DH ratio and values of  $\alpha_{s2}$ -CN Hydrolysis concentrations with HA concentration and ACE1 inhibition efficacy.

	H1pa	Н5ра	Н8ра	H1Ta	Н5Та	Н8Тα	H1Ma	H5Ma	H8Ma*
Hydrolysates s2-CNa	after	after	after	after	after	after	after	after	after
	1hr	5hr	8hr	1hr	5hr	8hr	1hr	5hr	8hr
Inhibition activity %	7.251	12.187	54.146	12.187	17.123	50.033	8.896	26.996	64.019*
Hippuric acid Conc	0.217	00.205	00.107	00.213	00.171	0.117	0.205	0.194	00.084
Hydrolysis degree %	3.700	36.638	48.293	05.929	23.412	37.753	4.200	38.860	61.924
Peptides Conc (group NH <sub>3</sub> )	3.023	29.794	39.267	04.835	19.044	30.700	3.66	31.600	50.340
Highest ACE1 inhibition ratio of S2-CNα hydrolysates: *									

concentration of 0.117 HA, 0.084 and 0.107 $\mu$ mol (Table 1). This means that DH led to increased inhibitory activity of the released hydrolysates, whether only peptides group or a single peptide chain, all of which were released from  $\alpha_{s2}$ -CN. From the table 1, the increase in the inhibitory activity was noticeable between 5 hours to the end of the incubation of 8 hours while the rise was between 1 hour and 5 hours less. The hydrolysates with the highest inhibitory activity was  $\alpha$ H8M of 64.019% with a concentration of peptides 50.346 mMol.

Bioactive peptides block the process of HHL conversion to HA and this theory has been confirmed by most studies conducted to demonstrate the effect of protein decomposers on the effectiveness of ACE1 one of the results of Sun et al., (2014) when hydrolyzing casein with pepsin and trypsin to obtain an inhibitory efficiency of 72% after 6 hours and the concentration of peptides 60.06 mMol. It is also consistent with Lee et al., (2005), which received a 96% inhibition in pepsin and 67% in trypsin only when the DH value of goat milk casein hydrolyzed to more than 48%. On the other hand, some researchers pointed out that the rise in the degree of hydrolysis is not the only factor leading to high efficiency inhibition of ACE1 Munn, (2013). The effect of high hydrolysis on the inhibitory efficiency (especially low inhibition at the beginning of the degradation process) is explained by the high concentration of peptides (quantitatively) and the increased chances of their diversification, especially when more than one type of enzyme is used and thus correlates with ACE1 catalytic sites and blocking the enzyme union With the substrate on which the enzyme acts in vivo (Chiang et al., 2006).

#### Conclusion

Local goat milk protein hydrolysates, especially  $\alpha_{s2}$ -CN, have the ability to inhibit ACE1 in vitro efficiently when using pepsin or trypsin and mix of them by ratio of 1:1 because these hydrolysates contain low molecular weight ACE1 inhibition peptides. Separating these peptides from their hydrolysates may increase their effectiveness. Peptides from other goat milk casein fractions hydrolysates ( $\beta$ ,  $\kappa$ -CN) may also have ACE1 inhibition effectiveness. *In vivo* studies to choose the effective hydrolysate which lowering blood pressure are important to establish the importance of eating goat milk and its products in lowering blood pressure because protein will be digested by digestive enzymes in the gut and release antihypertensive peptides.

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