Cloning and Expression of Camel Pro-Chymosin Encoding Gene in *E. coli* and Characterization of the Obtained Active Enzyme Aboulnaga, E. A. Food Science Department, Faculty of Agriculture, Mansoura University, 35516 Mansoura, Egypt.



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ABSTRACT

Chymosin is considered the main enzyme in milk industry since it is used for commercial production of cheese. Conventional animal chymosin production losses large numbers of unweaned calves which affect on the animal wealth. In order to reduce the manufacturing cost, recombinant production of chymosin is the good choice. In the current study, camel pro-chymosin gene (1101 bp) was *in vitro* synthesized and cloned into *pASG*-vector. The obtained plasmid *pASG_pro.chym* was transformed into the bacteria *E. coli* BL21(DE3). The bacterial system expressed pro-chymosin under control with tet-promoter and it was able to produce approximately 260 mg/L of recombinant enzyme under lab scale. The SDS-PAGE showed that the zymogen protein (367 amino acids residues long) has a molecular weight of 40.6 kDa, while the active form (323 amino acids residues long) has a molecular weight of 35.6 kDa. The recombinant pro-chymosin is presented in inclusion bodies and it is solubilized in 4-8 M urea. After solubilization and renaturation, recombinant pro-chymosin was subjected to a low pH and it was converted into mature active chymosin. The optimum milk clotting conditions were a pH of 5.75, temperature of 50-55 °C, and 15 mM of CaCl₂. It can be concluded that the obtained recombinant chymosin from *E. coli* is suitable for commercial cheese production.

Keywords: Camel chymosin- gene cloning- expression in E. coli- solubilization and renaturation- optimum clotting conditions.

INTRODUCTION

Chymosin (EC 3.4.23.4), an aspartyl protease, is found in mucosal cells of the fourth stomach of unweaned calves (Williams et al., 1997). The enzyme is responsible for partial proteolysis of κ -casein and it attacks the peptide bonds between the amino acids Phe₁₀₅ and Met₁₀₆ which destabilized casein micelles resulting in clotting milk (Albert et al., 1998). So, it is the main additives in cheese manufacturing. Many microorganisms such as Rhizomucor miehei (Grav et al., 1986) and R. Pusillus (Baudvs et al., 1988) produce rennet-like proteases that can replace the calf rennet. However, to produce high quality cheese, animal rennet enzyme (chymosin) is considered to be better choice than plant and microbial rennet which characterized by their un-specific hydrolysis of casein producing un-favourable flavour and taste as well as reducing the cheese yield (Kumar et al., 2010). For that, animal chymosin, especially bovine chymosin, is widely used in dairy industries for cheese production (Mohanty et al., 1999). However, traditional preparation of bovine chymosin requires slaughter of large number of unweaned calves with high price which lead to affect on national economy. Also, the extraction and purification is time consuming and its expose for deterioration agents is high (Starovoitova et al., 2006). So, biologists solve this problem by recombinant producing mammalian chymosin to provide the enzyme requirement in cheese industries. Currently, recombinant chymosin is used for one-third of the cheese production over the world (Starovoitova et al., 2006 and El-Sohaimy et al., 2010).

The calf chymosin gene was among the first mammalian genes that were cloned in different microorganism. Using recombinant DNA technology, calf chymosin encoding gene was expressed in GRAS (Generally Recognized As Safe) microorganisms as bacteria (Nishimori *et al.*, 1982; Emtage *et al.*, 1983; El-Sohaimy *et al.*, 2010; Menzella, 2011 and Noseda *et al.*, 2016), yeast (Mohanty *et al.*, 1999; Starovoitova *et al.*, 2006; Zhang *et al.*, 2009; Noseda *et al.*, 2013; Wang *et al.*, 2015 and Noseda *et al.*, 2016), filamentous fungus

(Cardoza *et al.*, 2003 and Sharma *et al.*, 2009), and plant (van Rooijen *et al.*, 2008).

First approach was made by cloning of full-length cDNA of the mRNA and their sequence has been determined (Nishimori et al., 1982). The precursor protein, pre-pro-chymosin, has a 16 amino acids signal peptide that was corresponding on transfer the zymogen pro-chymosin into stomach which is further activated by removal of the pro-peptide, the 42-amino acids from the NH₂-terminus (Emtage et al., 1983). Expression of the pro-chymosin in yeast or fungus resulting on production of the enzyme in an active form. While, expression of the pro-enzyme in E. coli exit as insoluble form which needs further solubilization and renaturation of the enzyme from the inclusion bodies (Mule et al., 2009 and El-Sohaimy et al., 2010). The major benefit of this insoluble form that it makes purification and activation could be achieved since most of E. coli proteins (about 90%) are precipitated under the acidic condition which is used to activate the pro-chymosin (Rodríguez-Carmona, 2010). Therefore, several approaches have been developed for optimizing the solubilization and renaturation process to reduce the process cost as well as increase the yield of the mature active chymosin (Koritzinsky et al., 2013). The enzymatic characteristics of E. coli recombinant chymosin are indistinguishable from those of native one (Starovoitova et al., 2006).

Recently, recombinant camel chymosin attracts great interest in cheese industries because it produces cheese with good flavour and extended cheese shelf life (Langholm Jensen *et al.*, 2013). It has the ability to clot both camel's and cow's milk. In addition, it has 70% higher clotting activity, exhibits higher level of thermostability, and exhibits sevenfold higher clotting activity/proteolytic activity than bovine chymosin (Bansal *et al.*, 2009 and Sorensen *et al.*, 2011).

Camel chymosin has been cloned and expressed firstly in *Aspergillus niger* (Kappeler *et al.*, 2006). The second group was expressed it in yeast *Pichia pastoris* with almost similar properties like the one produce by fungi (Wang *et al.*, 2015). However, expression of camel chymosin in *E. coli* has not been yet reported till now. In the current work, a full-length camel pro-chymosin gene

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was *in vitro* synthesized, introduced into *E. coli* BL21(DE3) strain via pASG-plasmid, and expressed under tet-promoter. The insoluble form of the pro-chymosin is further renaturation and activation. Also, the optimum clotting condition for the active camel chymosin was studied.

MATERIALS AND METHODS

Materials

The chemicals used in the current work were purchased from the following company, Sigma-Aldrich,

IBA-lifesciences, and Takara. Oligonucleotides were purchased from Tsingke (China). The DNA purification kit, Phusion[™] DNA polymerase, and restriction enzyme were purchased from Fermentas-GmbH (Fisher scientific, Schwerte, Germany).

Strains, plasmids and primers.

Bacterial strains, empty vector, and constructed plasmids that were used in this work are presented in table (1).

Table 1. Bacterial strains and Plasmids used in the current study:					
Strains	Source				
<i>E. coli</i> , DH5α-cells (Cloning strain)	fhuA2 Δ(argF-lacZ)U169 phoA glnV44 Φ80 Δ(lacZ)M15 gyrA96 recA1 relA1 endA1 thi-1 hsdR17	Invitrogen			
E. coli BL21(DE3)	$F^{-} ompT hsdS_{B}(r_{B} m_{B}^{-}) gal dcm rnel31(DE3)$	Invitrogen			
Original vectors					
pE/pENTRY	Cloning & compatible expression vector; Kan ^R ; ColE1 ori; lacP/Z				
pASG.wt-(blue)	std. expression vector; pseudo-wt; Amp ^R ; lacP/Z	T. Selmer			
pFusion (blue)	fusion vector; Amp ^R ; lacP/Z				
prepared plasmids	•				
pE-chym.P.1	Kan ^R ; ColE1 ori; Chym (part1)				
pE-chym.P.2	Kan ^R ; ColE1 ori; Chym (part2)				
pF-pro.chym	Amp ^R ; ColE1 ori; pro-Chym	This study			
pE-pro.chym	Kan ^R ; ColE1 ori; pro-Chym	-			
pASG-pro.chym	tetP; Pro.chym; f1ori; Amp ^R ; ColE1 ori				
Amp ^R , Ampicillin resistance gene; Kan ^R , Kanamycin resistance gene					

Construction of the chymosin gene.

Camelus dromedaries pro-chymosin protein (367 amino acid, with calculated molecular weight of 40.6 KD) was converted into its coding gene sequence (1101bp) taken in consideration codon optimization for enhancing

optimum protein expression in *E. coli*. The pro-chymosin gene (1101 bp) was artificial synthesized by overlapping PCR technique (Bryksin and Matsumura, 2010) in KanRvector (pEntry) using the oligonucleotides presented in table (2).

Table 2. List of all oligonucleotides used throughout this study.

	Name	Sequence	bP
Oligonucleotides to synthesize the first part of chymosin gene	P1.F1	CATGATGGACCGCCACCTGGTGGCGCGCGCGACCTGTTCAGCGTGGAGACGATCCAAAGGC	59
	P1.F2	ACCCGTCCCTGGCCAGCGAGTACTCGGTGCCGGTGTTCGACAACATGATGGACCGCCAC	59
	P1.F3	GGTGTTCACCTACTCGGAGTTCGACGGCATCCTGGGCCTGGCCTACCCGTCCCTGGCCA	59
	P1.F4	CGAACCAGACCGTGGGCCTGAGCACGGAGCAGCCGGGCGAGGTGTTCACCTACTCGG	57
	P1.F5	CTTCCTGGGCTACGACACGGTGACCGTGAGCAACATCGTCGACCCGAACCAGACCGTGG	59
	P1.F6	CAAGCCGCTGAGCATCCATTACGGCACCGGCAGCATGGAGGGCTTCCTGGGCTACGAC	55
	P1.F7	AAGCTCTTCATCCCCGCAAGAGCTCGACGTTCCGCAACCTGGGCAAGCCGCTGAGCATC	59
	P1.R1	CTTGTGCAGCGGGATGCGGGGGGGGGGGCCGAGGCCATTGGAGACGTGGCAC	50
	P1.R2	CAGGCCGCGTTCCTTCAGCGCCTTGCGCAGGGTCTTGCCCTTGTGCAGCGGGATG	55
	P1.R3	GCTGCTCACGGCGTACTGCTGGCGCTGCAGGAAGTCCTCCAGCAGGCCGCGTTCCTTC	58
	P1.R4	TGGTCAGCGGTTCGCGGGCCACCTTGCCCAGCGAGCTGTACTTGCTGCTCACGGCGTA	58
	P1.R5	GGTGCCGATGTAGATCTTGCCGAAGTACTGGCTGTCCAGGTAGCTGGTCAGCGGTTCGC	59
	P1.R6	GCTACCGGTGTCGAACACGACCGTGAACTCCTGCGGGGGGGG	58
	P1.R7	CACGTTGCTCTTGCAGTAGATGCTCGGGACCCACAGGTCCGAGCTACCGGTGTCGAAC	58
	P1.R8	AAGCTCTTCTGGATCGAAGCGATGGTGGTTCTTGCACACGTTGCTCTTGCAGTAG	55
št.	P2.F1	ACCGCGCGAACAATCGCGTGGGCCTGGCGAAGGCGATCTAATAGGGAGGAGACGATCCA	59
firs	P2.F2	CCTGGGCGACGTCTTCATCCGCGAGTACTATAGCGTGTTCGACCGCGCGAACAATCG	57
the	P2.F3	GCACCAGCGGTTTCCAGGGCGACAACAATAGCGAGCTGTGGATCCTGGGCGACGTCTTC	59
Oligonucleotides to synthesize t part of chymosin gene	P2.F4	TCCCCTGAGCCCGAGCGCGTACACCAGCAAGGACCAGGGCTTCTGCACCAGCGGTTTCC	59
	P2.F5	CGCTCCATGCCCACCGTGGTGTTCGAGATCAACGGCCGCGACTATCCCCTGAGCCCGAG	59
	P2.F6	CCGAGAACCGCTACGGCGAGTTCGACGTGAACTGCGGCAGCCTGCGCTCCATGCCCACC	59
	P2.F7	AAGCTCTTCATCGGTGCCACCGAGAACCGCTACGG	35
	P2.R1	GTCAGCATGCTGCCCTGGCCGTTGCGGTCCATGTACACGCGGAGACGTGGCACTTTTCG	59
	P2.R2	CAGGCTACCGGTGTAGTACGACGGGTCGATCGCGCCCAGGGTCAGCATGCTGCCC	55
	P2.R3	CGGTGAACTGCCAGTACTGCTGGACGGTCACCGGCACCCAGTGCAGGCTACCGGTGTAG	59
	P2.R4	GACGGCCACACCGTTGATCGTCACGCTGTCCACGGTGAACTGCCAGTAC	49
	P2.R5	GCCGGTGTCCAGGATCGCCTGGCAACCACCGACGCAGGCGACGGCCACACCGTTG	55
	P2.R6	CTTCAGGATGTCGCTGCTCGGGCCGAACAGCACGCTCGTGCCGGTGTCCAGGATC	55
<u> </u>	P2.R7	AAGCTCTTCTCGATCGCCATCTGGATCTTCAGGATGTCGCTGC	43

Restriction site for the *LguI* restriction enzyme is: GCTCTTC,

The gene was divided into two fragments (Chym.P.1 and chyms.P.2) for accurate assembly. After assembly, the PCR products was cut with *lguI* and ligated with *T4-ligase* then it was transferred into *E. coli* DH5 α with the general protocol (Chung *et al.*, 1989). The correct two plasmids (pE-chym.P.1 and pE-chym.P.2) were extracted and identified by cutting with restriction enzyme before sending for sequencing. Afterwards, the two plasmids were cut with *Esp31* restriction enzyme and the two fragments were fused together and inserted in AmpR vector (pFusion). Then, the assembled fragment (1101 bp) was cloned in pEntry vector to produce *pE-pro.chym* and then sub-cloned into the expression vector (pASG.wt). The expression plasmid (*pASG-pro.chym*) was used for protein production.

Bacteria cultivation:

Escherichia coli DH5 α and *E. coli* BL21 (DE3) cells were cultivated at 37°C on lysogeny broth (LB) media supplemented with the appropriate antibiotics according to the demand (ampicillin 100 µg/ml or kanamycin 50 µg/ml). The required plasmids were transferred into chemically competent *E. coli* BL21 (DE3) or DH5 α cells following the standard protocol (Chung *et al.*, 1989).

Gene expression and biomass production:

Pro-chymosin gene was expressed in *E. coli* BL21(DE3). Freshly transformed cells were plated on LB-agar medium supplemented with (100 μ g/mL Ampicillin) and one correct colony was expanded step-wise to yield final culture volume of 500 mL in LB-Medium (100 μ g/mL Ampicillin). When the cells reached an OD₅₇₈ nm of about 0.4, the gene was expressed by adding the inducer (anhydrotetracycline, 200 ng/mL) for 2 hr at 37 °C. The induced cells were harvested by centrifugation at 6,000 g for 15 min, washed with Tris-buffer (50 mM Tris-HCl, 100 mM NaCl, 1 mM EDTA, pH 8.0) and stored at -80°C until used (Aboulnaga *et al.*, 2013).

Inclusion bodies purification:

Inclusion bodies purification was started with about one gram of wet cells. Cells were resuspended in 20 ml of lysis buffer (50 mM Tris-HCl, 100 mM NaCl, 1 mM EDTA, 1 mg/ml lysozyme, pH 8.0) and incubated at 37 °C for 30 min. Then, the suspension was sonicated in an icewater bath using a Branson Sonifier with a duty cycle of 50% at 60% power for 10 min. The inclusion bodies were isolated from the mixture by centrifugation at 10,000 g for 20 min at room temperature. Clear supernatant was discarded and the pellet was washed twice with 50 ml of 10 mM EDTA, once with 0.5% triton X-100, and once with 20 mM phosphate buffer pH 7.5 (Rodríguez-Carmona, 2010).

Solubilization and renaturation of recombinant prochymosin:

Solubilization and renaturation has been done according to Menzella (2011) protocol with the following modification. Cleaned inclusion bodies were dissolved in solubilization buffer (50 mM KH₂PO₄, 8 M urea, pH 10.5). The solution was incubated at 30°C for 2 hr with continuous stirring, and then centrifuged at 10,000 g for 10 min to remove the insoluble molecules. The clear preparation was diluted with the renaturation buffer (50 mM KH₂PO₄, 0.5 M arginine, 10 μ M CuSO₄, pH 10.5) to

have a final protein concentration of 1 mg/ml (dilution factor between 20-30 times) and incubated for 12 hr at 4 °C. The correctly folding pro-chymosin was activated by acidifying the solution to pH 2.0 with 1.0 M HCl and incubated for 6-8 hr at 20 °C for digesting the pro-peptide. Then, the pH was adjusted to 6.3 by adding 1 M NaOH and it was incubated for second time for an additional 2 hr and the active chymosin was stored at 4 °C.

Enzyme activity measurements:

The milk-clotting activity of the chymosin was determined based on the end-point dilution methods (Emtage *et al.*, 1983). The substrate was prepared by reconstitute 10 g skim milk in 100 ml phosphate buffer (20 mM KH₂PO₄, 10 mM CaCl₂, pH 5.5) at room temperature (25 °C) and then the skim milk solution was preincubated at 35 °C for 30 min before measurement. The international milk clotting units (IMCU) was calculated by using commercial calf chymosin (1500 IMCU) as standard. One unit was defined as the amount of enzyme necessary to clot 1.0 ml milk at 35 °C in 40 min.

Determination of optimal pH for recombinant chymosin activity:

To identify the optimum pH for recombinant chymosin activity, skim milk powder was reconstitute in either 50 mM acetate buffer (pH 5.0- 5.5) or 50 mM potassium phosphate (pH 5.5–6.5) buffer. The clotting activity was measured as mentioned above and the relative activity was plotted against pH.

Determination of optimal temperature for recombinant chymosin activity:

The optimum temperature was determined after incubating both enzyme and substrate solutions (skim milk powder 10% in 20 mM KH₂PO₄, 10 mM CaCl₂, pH 5.5) at different temperature ranging from 40 °C to 65 °C for 10 min. Afterwards, the enzyme solution was added and gently mixed with substrate and incubated at the selected temperature to measure the clotting activity.

SDS-PAGE analysis:

SDS-PAGE was performed with 12% polyacrylamide gel to separate proteins of the whole cell extract, cell free extract, and inclusion bodies in the presence of pre-stained marker (10- 250 kDa). To visualize the protein bands, Coomassie brilliant blue (G-250) was used to stain the gels (Laemmli, 1970).

RESULTS AND DISCUSSION

Construction of the camel pro-chymosin gene and expression plasmid:

The amino acids sequence of camel pro-chymosin protein (Fig.1) was obtained from the online database. The encoding codon for this protein was optimized to facilitate maximum expression in *E. coli*. The optimum codon encoding for camel pro-chymosin gene (pro.chym, 1101 base pair) was *in vitro* synthesized and inserted in a bacterial cloning vector (*pE-Entry*). After plasmid extraction, the *pE-pro.chym* plasmid was sequenced to identify the inserted fragment (Fig. 2). The correct plasmid was used further for sub-cloning into expression vector directly downstream *E. coli* strong *tet*-promoter. The resulting plasmid, *pASG-pro.chym*, had a 4263 bp (Fig.3.A) which was appeared as a strong band on gel

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electrophoresis after digestion with HindIII/ or SacI restriction enzyme (Fig.3.B). The transformed E. coli BL21 (DE3) with pASG-pro.chym plasmid was further analyzed with colonies PCR-screening using internal vector-based-primers (Seq.F and Seq.R) which produced a 1304 bp fragment (Fig.3.C). The correct colony carrying the pASG-pro.chym plasmid should produce a 1304 bp fragment (Fig.3.D).

>Camel chymosin Amino acid sequence

MASGITRIPLHKCKTLRKALKERGLLEDFLQRQQYAVSSKYSSLGKVAREPLTSYLDSQYFG KIYIGTPPQEFTVVFDTGSSDLWVPSIYCKSNVCKNHHRFDPRKSSTFRNLGKPLSIHYGTG SMEGFLGYDTVTVSNIVDPNQTVGLSTEQPGEVFTYSEFDGILGLAYPSLASEYSVPVFDNM MDRHLVARDLFSVYMDRNGOGSMLTLGAIDPSYYTGSLHWVPVTVOOYWOFTVDSVTINGVA VACVGGCQAILDTGTSVLFGPSSDILKIQMAIGATENRYGEFDVNCGSLRSMPTVVFEINGR DYPLSPSAYTSKDQGFCTSGFQGDNNSELWILGDVFIREYYSVFDRANNRVGLAKAI

Fig. 1. Complete Amino acids sequence of camel chymosin protein. The first 44 amino acids which present the prosequence are written in bold red italic line.

>Camel chyr	nosin DNA se	equence		
ATGGCCTCGG	GCATCACCCG	CATCCCGCTG	CACAAGGGCA	AGACCCTGCG
CAAGGCGCTG	AAGGAACGCG	GCCTGCTGGA	GGACTTCCTG	CAGCGCCAGC
AGTACGCCGT	GAGCAGCAAG	TACAGCTCGC	TGGGCAAGGT	GGCCCGCGAA
CCGCTGACCA	GCTACCTGGA	CAGCCAGTAC	TTCGGCAAGA	TCTACATCGG
CACCCCCCCG	CAGGAGTTCA	CGGTCGTGTT	CGACACCGGT	AGCTCGGACC
TGTGGGGTCCC	GAGCATCTAC	TGCAAGAGCA	ACGTGTGCAA	GAACCACCAT
CGCTTCGATC	CCCGCAAGAG	CTCGACGTTC	CGCAACCTGG	GCAAGCCGCT
GAGCATCCAT	TACGGCACCG	GCAGCATGGA	GGGCTTCCTG	GGCTACGACA
CGGTGACCGT	GAGCAACATC	GTCGACCCGA	ACCAGACCGT	GGGCCTGAGC
ACGGAGCAGC	CGGGCGAGGT	GTTCACCTAC	TCGGAGTTCG	ACGGCATCCT
GGGCCTGGCC	TACCCGTCCC	TGGCCAGCGA	GTACTCGGTG	CCGGTGTTCG
ACAACATGAT	GGACCGCCAC	CTGGTGGCGC	GCGACCTGTT	CAGCGTGTAC
ATGGACCGCA	ACGGCCAGGG	CAGCATGCTG	ACCCTGGGCG	CGATCGACCC
GTCGTACTAC	ACCGGTAGCC	TGCACTGGGT	GCCGGTGACC	GTCCAGCAGT
ACTGGCAGTT	CACCGTGGAC	AGCGTGACGA	TCAACGGTGT	GGCCGTCGCC
TGCGTCGGTG	GTTGCCAGGC	GATCCTGGAC	ACCGGCACGA	GCGTGCTGTT
CGGCCCGAGC	AGCGACATCC	TGAAGATCCA	GATGGCGATC	GGTGCCACCG
AGAACCGCTA	CGGCGAGTTC	GACGTGAACT	GCGGCAGCCT	GCGCTCCATG
CCCACCGTGG	TGTTCGAGAT	CAACGGCCGC	GACTATCCCC	TGAGCCCGAG
CGCGTACACC	AGCAAGGACC	AGGGCTTCTG	CACCAGCGGT	TTCCAGGGCG
ACAACAATAG	CGAGCTGTGG	ATCCTGGGCG	ACGTCTTCAT	CCGCGAGTAC
TATAGCGTGT	TCGACCGCGC	GAACAATCGC	GTGGGCCTGG	CGAAGGCGAT





Fig. 3. Construction of the camel pro-chymosin gene and expression plasmid. (A) Schematic representation of the pASGpro.chym plasmid. The sequence of encoding camel pro-chymosin was expressed under tet-promoter. (B) DNA electrophoresis for the pASG-pro.chym plasmid after digestion with HindIII (lane 1) and SacI (lane 2) restriction enzymes; (C) Schematic representation of DNAfragment (1,304 bp) containing pro.chym gene after amplification with the seq.F and seq.R primers. (D) DNA-electrophoresis of colonies PCR with seq.F and seq.R primers for 5 different colonies.

Heterologous expression of the pro-chymosin gene in *E. coli*:

Correct E. coli BL21 (DE3) colonies carrying pASG-pro.chvm plasmid were tested for pro-chymosin expression. Cells were grown in LB-media supplemented with ampicillin (100 µg/mL) and induced with 200 ng/mL anhydrotetracycline when the OD₅₇₈ was 0.4. Biomass was harvested before induction and 2 hr post-induction. Cellular extracts were subjected to electrophoresis on 12% SDS- polyacrylamide gel (Fig.4). Obtained result from the analysis of pro-chymosin by SDS-PAGE (Fig.4) revealed that pro-chymosin (367 amino acid residue) was visualized as a sick band. Also its protein had a molecular weight with about 41 kDa, which it is almost similar to the calculated molecular weight (40.6 kDa) as well as the previously reported molecular weight (Kappeler et al., 2007; Langholm Jensen et al., 2013 and Wang et al., 2015). Further analysis of the produced pro-chymosin band using LabWork3.0 indicating that the amount of recombinant protein was 250 mg/L and it was comparable with that reported before for recombinant bovine chymosin (Menzella, 2011). The reported expression level of E. coli BL21 (DE3) for recombinant bovine chymosin after 5 hr post-induction was ranged from 260 for the wild type sequence to 450 mg/L after codon optimization (Menzella, 2011). Since the sequence was optimized, it should produce the highest level but the presented data (250 mg/L) was only after 2 hr post-induction which make it comparable with the previously reported data (Menzella, 2011).



Fig. 4. Expression analysis of the pro-chymosin by SDS-PAGE. M, is molecular weight ladder. Lane 1, lysate of *E. coli* BL21 harbouring *pASG-pro.chym* expression plasmid without induction. Lane 2 and 3, lysate of two different colonies of *E. coli* BL21 harbouring *pASG-pro.chym* expression plasmid after 2hr induction.

Solubilization and renaturation of recombinant prochymosin:

Recombinant pro-chymosin is expressed in bacteria in insoluble form and it is presented in the inclusion bodies. This finding has been reported for the production of recombinant bovine chymosin in *E. coli* (Mule *et al.*, 2009; El-Sohaimy *et al.*, 2010 and Menzella, 2011). The recombinant camel chymosin behaves like the bovine one that is presented in inclusion bodies and is not presented in the cell free extract (Fig. 5). The protein is completely soluble in the solubilizing solution which contains 8.0 M urea. After renaturation and subsequently activation, the mature active form of chymosin shows a molecular weight of 35.6 kDa (Fig.5, lane 4 and 5). It should be noticed that, recombinant camel chymosin expressed in fungus or yeast is glycosylated and the apparent molecular weight is between 38-40 kDa, while after treatment with endoglycosidase, the protein show a band corresponding to a molecular weight of 35-37kDa (Kappeler, 2006; Langholm Jensen *et al.*, 2013 and Wang *et al.*, 2015). Since bacteria system does not glycosylate protein which is only happen in eukaryote systems, the apparent molecular weight of protein is almost similar to the calculated one.





Activity measurement of recombinant camel chymosin:

The clotting activity of recombinant camel chymosin obtained from E. coli was compared with commercial calf chymosin in Fig. (6). Recombinant chymosin without renaturation is not active and required renaturation process for the generation of active clotting enzyme. Furthermore, the active ammonium sulphate precipitated recombinant camel chymosin has doubled the clotting activity of commercial calf chymosin. This indicates that, the enzyme has a 300 IMCU/ ml. Therefore, it could be concluded that biologically active chymosin expressed in fungi (Kappeler, 2006) or yeast (Wang et al., 2015) can be produced in E. coli after solubilization, renaturation, and acidic activation. Although, calf chymosin has been expressed in E. coli and it was active after solubilization, renaturation, and acidic treatment (El-Sohaimy et al., 2010 and Menzella, 2011).



Fig. 6. Milk clotting activity of commercial calf chymosin and the recombinant camel chymosin obtained from *E. coli*. Serial dilutions were prepared in microtiter plate and the clotted milk in the wells is shown. A (column 1-3): Serial dilution of the standard calf chymosin (150 IMU/ ml) was prepared. B (column 5-7): Presented serial dilution of the recombinant camel chymosin derived from *E. coli* before renaturation. C (column 9-11): Presented serial dilution of the recombinant camel chymosin after renaturation and activation.

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Optimal pH for the produced recombinant camel chymosin:

The effect of the substrate pH on the clotting activity of recombinant chymosin was evaluated and the results are presented in Fig. (7). The recombinant enzyme has a narrow pH-range for its activity. The enzyme shows more than 80% of its relative activity with the pH value of 5.25 and more than 90% of its relative activity when the pH values were 5.5 or 6.0. The highest level of recombinant camel chymosin activity was observed at pH between 5.5 and 6.0 which is also the range of the animal's milk pH. Two different optimum pH values have been reported before for recombinant camel chymosin. The recombinantly produced camel chymosin in Asperigillus niger had an optimum pH of 6.0 (Kappeler, 2006), while the produced enzyme in Pichia pastoris had an optimum pH value of 5.04 (Wang et al., 2015). In contrast, recombinant bovine chymosin had a narrow pH-range for its activity between 5.4 to 5.6 (Cardoza et al., 2003; Starovoitova et al., 2006 and Noseda et al., 2013).



Fig. 7. Optimum substrate pH for the recombinant camel chymosin activity. The effect of pH values on the milk clotting activity of recombinant camel chymosin was determined and it was plotted against the relative activity.

Optimal temperature for the produced recombinant camel chymosin:

Figure (8) show the effect of temperature on the clotting activity of the recombinant camel chymosin. There was a significant effect of the temperature on the milk clotting activity. The enzyme show maximum activity (100%) when the temperature was in the range of 50-55 °C. When the temperature was changed within 5 °C down or above the optimum range, a significant decrease of the activity was observed and it was reached of 70 and 60% relative activity, respectively. Further increment of temperature had a significant inhibition effect on the activity due to denature the proteins (Wang et al., 2015). A further decrease in relative activity to reach about 35 and 5% was observed when the temperature was 65 and 70 °C, respectively. These data are in agreement with the previously published data, in which camel chymosin had an optimum temperature of 45-55 (Kappeler, 2006 and Wang et al., 2015). It is known that, animal source chymosin is temperature sensitive opposite to plant source rennin which is more resistance for high temperature (Gama Salgado et al., 2013 and Li Yuqiu and Qiyun, 2015).





Optimal concentration of CaCl₂:

The effect of CaCl₂ on the clotting activity of the recombinant camel chymosin was studied and the data were plotted in Figure (9). Calcium ions has high impact in milk clotting in which it bind with the insoluble peptide derived (Para-casein) from casein hydrolysis and form clot.

The data in Fig. (9) show that, the enzyme required narrow concentration of CaCl₂ for optimum activation. The maximum activity was observed when CaCl₂ concentration was 15- 30 mM and less than 90% relative activity was detected when CaCl₂ concentration was 10 or 45 mM. These data were closed with the reported results for recombinant camel chymosin (Wang *et al.*, 2015) and also with the microbial rennin that has optimum CaCl₂ concentration of 50 mM (Foda, 2012).



Fig. 9. Optimum concentration of CaCl₂. The effect of CaCl₂ on the milk clotting activity of recombinant camel chymosin was determined and it was plotted against the relative activity.

CONCLUSION

In this work, recombinant camel pro-chymosin was successfully constructed and expressed in *E. coli* BL21 under tet-promoter. The protein was produced in an insoluble form. It was *in vitro* solubilized, re-folded from the inclusion bodies, and activated under acidic conditions.

The optimum milk-clotting activity conditions for mature camel chymosin were pH of 5.75, temperature between 50-55 °C, and 15 mM of CaCl₂. The produced recombinant chymosin could serve as an economic alternative source to bovine chymosin in cheese manufacturing. This study provide that, it is possible to obtain an active mature camel chymosin from *E. coli* as a first step for large-scale production of recombinant enzyme to meet the needs of local market and maintain the foreign currency.

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الاستنساخ والتعبير الجيني لمولد- الكيموزين الجملي في خلايا بكتيريا الـ E. coli وتوصيف الانزيم النشط المتحصل عليه الحسينى احمد على ابوالنجا

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