

# Identification of Polymorphisms of Gene CSN2 of B Casein in Greek Cow Breeds (Holstein) by Restriction Fragment Length Polymorphism

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## Abstract

The research focused to detect and identify genetic polymorphisms in exon 7 of the  $\beta$ -casein CSN2 gene in blood samples from Greek Holstein cows. For this purpose, DNA was extracted from 120 blood samples of cows. The desired region of exon 7 was amplified by PCR, resulting in a 121 bp product. The PCR product was digested by restriction fragment length polymorphism (RFLP) method. The results suggest that the A1A2 genotype prevails over the others. Specifically, of the 120 cattle, 72 showed triple bands of 121 bp, 86 bp, and 35 bp indicating the A1A2 genotype. The 42 cattle showed a single band at 121 bp, indicating that they carried the A1A1 genotype. The remaining 6 showed only two bands of 86 and 35 bps, indicating that they carried the A2A2 genotype. In the total population of heterozygotes A1A2-0.60 were the most frequent, while homozygotes A2A2-0.06 were the least frequent ones. This suggests a slight superiority of allele A-0.65.

**Keywords:**  $\beta$ -casein, b-casomorphin-7, polymorphism, PCR-RFLP

## 1. Introduction

Milk has always played an important role in the human diet, as it is considered a considerable source of all essential amino-acids, vitamins, metals, minerals. More specifically, bovine milk contains a total of 3.5% protein (Davoodi et al., 2016). Proteins are one of the most important milk components, that have received the greatest attention due to their recognized health-related properties. Casein (as1, as2, beta, kappa) accounts for 80% of the total protein content (Haug, Høstmark, & Harstad, 2007). The remaining 20% of the protein content is the serum protein (Davoodi et al., 2016). Beta casein's gene (CSN2) belongs to the cluster of four casein genes located on chromosome 6 (Ferretti, Leone, & Sgaramella, 1990; Jann, Ceriotti, Caroli, & Erhardt, 2002; Kumar et al., 2019). It is known that beta casein has 12 genetic variants, (A1, A2, A3, B, C, D, E, F, G, H1, H2 and I) (Kumar et al., 2019; Massella et al., 2017; Sebastiani et al., 2020). Among them, the most frequent genetic variants of CSN2 are A1 and A2 (Balteanu, Vlaic, Suteu, & Carsai, 2010; Cieślińska et al., 2012; Dinc, Ozkan, Koban, & Togan, 2013). The variants A1 and A2 are composed of 209 amino-acids. The A1 variant differs to the A2 at one amino acid of the amino acid chain. More specifically, proline (codon CCT) on the 67th position (A2 variant) has been replaced by histidine (codon CAT) (A1 variant) (Kumar et al., 2019; Miluchová et al., 2014; Massella et al., 2017). This mutation has an effect on the proteolytic digestion of the primary protein structure, leading to the production of different peptides. B-casomorphin is a bioactive peptide produced from  $\beta$ -casein variants A1 and has a significant opioid activity (Brooke-Taylor, Dwyer, Woodford, & Kost, 2017; Massella et al., 2017). BCM-7 is a small molecule that can infiltrate blood circulation more easily, and cause various health problems, such as gastrointestinal disorders, insulin-dependent diabetes, atherosclerosis, ischemic heart diseases and sudden infant death syndrome (Brooke-Taylor et al., 2017; Boztepe, Aytakin, & Şahin, 2018; Elliott, Harris, Hill, Bibby, & Wasmuth, 1999; Kumar et al., 2019; McLachlan, 2001; Sun et al., 2003). To our knowledge, there is a lack of genetic polymorphism of b-casein in cows that are bred in Greece. The aim of this study was to identify the frequency and type of genetic polymorphisms in exon 7 of the  $\beta$ -casein CSN2 gene in blood samples from Greek cows Breeds (Holstein).

## 2. Materials and Methods

### 2.1 Experimental Animals and Farms

A total of 120 Greek Holstein-Friesian cows studied in this research were taken from two different regions of Greece. The first farm was located in Korinthia, an area of the northeastern Peloponnese and the second in Larissa, a city in Thessaly. A representative genotype and correlation study was performed for the A1/A2 sites of the  $\beta$ -casein gene in Greek cows in the country.

### 2.2 Institutional Animal Care and Use Committee (IACUC)

Animal blood sampling was based on the 2010/63 EU guidelines of European community and council on the protection of animals used for scientific purposes. According to directive article 1, paragraph 5, element f, 'practices not likely to cause pain, suffering, distress or lasting harm equivalent to, or higher than, that caused by the introduction of a needle in accordance with good veterinary practice'.

### 2.3 Selection and Preparation of Control Samples

Blood samples from Cow identified as A2A2 (characterised as control samples) were acquired directly from the animals and were kept at -18 °C until the analysis.

### 2.4 Selection and Preparation of Blood Samples

During the study period (September 2019 to March 2020) a total of 120 blood samples were collected from Holstein-Friesian cows, in tubes containing Ethylene diamine tetra-acetic acid (EDTA) and were kept at -18 °C until the analysis.

### 2.5 DNA Isolation

350  $\mu$ l of blood from each cow was used. The control samples were centrifugated (10 minutes at 12000  $\times$  g) in order to obtain a pellet. The DNA was extracted by using NucleoSpin Blood kit, by Machery-Nagel. According to the manufacturer's instructions with a modification of an overnight incubation with the Lysis Buffer and the Proteinase K at 65 °C instead an incubation of the 30 min incubation. The extracted DNA was quantified spectrophotometrically at 260 nm.

### 2.6 PCR Amplification and Restriction Fragment Length Polymorphism Detection

The region of exon 7 of CSN2 was amplified by PCR. The DNA primers used for the PCR amplification, were forward primer 5'-CCTTCTTTCCAGGATGAACTC CAGG-3' and reverse primer 5'-GAGTAAGAGGAGGG ATGTTTTGTGGGAGGCTCT-3'. DNA primers described by Saran, Gurao, Joshi, and Kashyap (2019). The PCR mix composed: 0.3  $\mu$ l of primers Csn4F and Csn4R (HS ReadyMix and dye, KAPABIOSTSTEMS). The amplification was conducted by a thermal cycler (96 Well thermal cycler applied Biosystems, Singapore), as follows: The chosen temperatures were set at 94-96 °C for 4 min, 35 cycles of 95 °C for 60 sec, 58 °C for 60 sec, 72 °C for 60 sec, and final extension of 7min at 72 °C, for Csn4F and Csn4R. PCR products were separated in 2% agarose gel. The yield and specificity of the PCR products were evaluated after electrophoresis in 2% agarose gel stained with ethidium bromide (0.5  $\mu$ g/ml) and documented under UV illumination using MiniBIS Pro device (DNR Bio-Imaging Systems Ltd., Israel) producing a 121 bp fragment of the CSN2 gene. The PCR products were then digested with 5 U of DdeI, and the restriction digestion fragments were analyzed by electrophoresis on a 3% agarose gel, in order to identify the genotype. The RFLP pattern is used to identify the genotypes as A1A1 (121 bp), A2A2 (86 and 35 bp) or A1A2 (121, 86 and 35 bp).

Each A2A2 homozygote was confirmed a second time via a genotyping procedure. Genetic equilibrium of the examined population was estimated due to the Hardy-Weinberg principle and tested with the chi-square test (<http://www.dr-petrek.eu/documents/HWE.xls>). The study was conducted with compliance to local bioethics committee guidelines (18/2013).

## 3. Results

Genomic DNA was isolated from 120 blood samples from Holstein-Friesian breed cows. The extracted DNA collected was checked for its concentration and purity by photometry. All samples had a concentration of 80-120 ng/ $\mu$ l and a purity of 1.5-1.9 ng/ $\mu$ l. Subsequent amplification of exon 7 of  $\beta$ -casein was followed by the appropriate primers. Electrophoresis on 1.8% agarose gel was performed to verify the method. A PCR of a known molecular weight of 50 bp (ladder) was added during the electrophoresis of the PCR products to calculate the molecular size of the fragment. The amplified segments were 121 bp in size. The figure below shows a typical result of a PCR amplification of the portion of the gene studied. All the 120 blood samples had the amplified size.

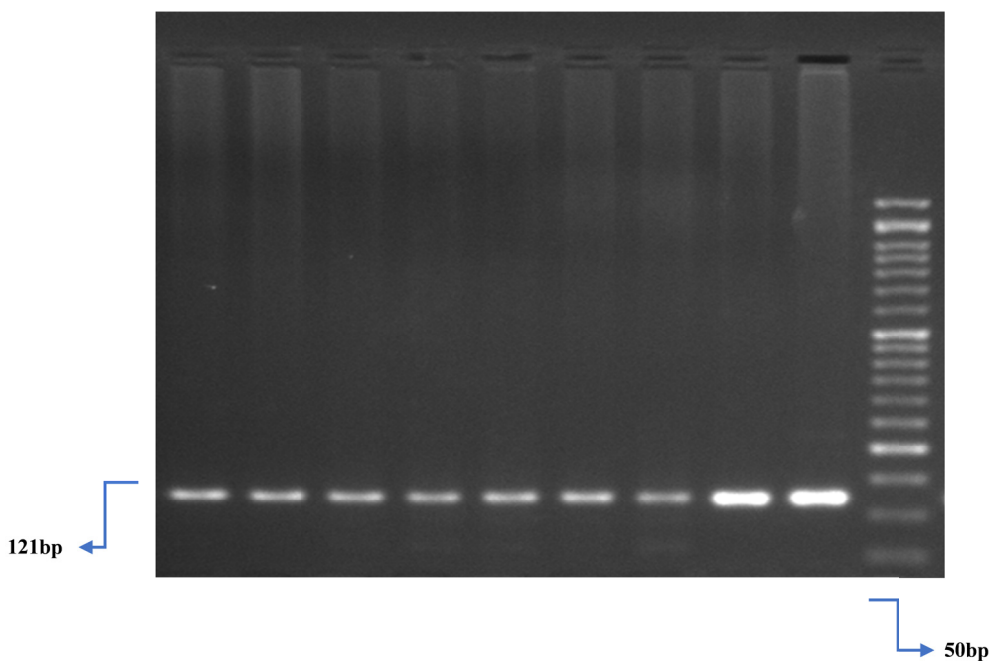


Figure 1. Electrophoresis showing PCR products (121 bp) on 1.8% w/v agarose gel with ethidium bromide staining; DNA ladder: 50 bp

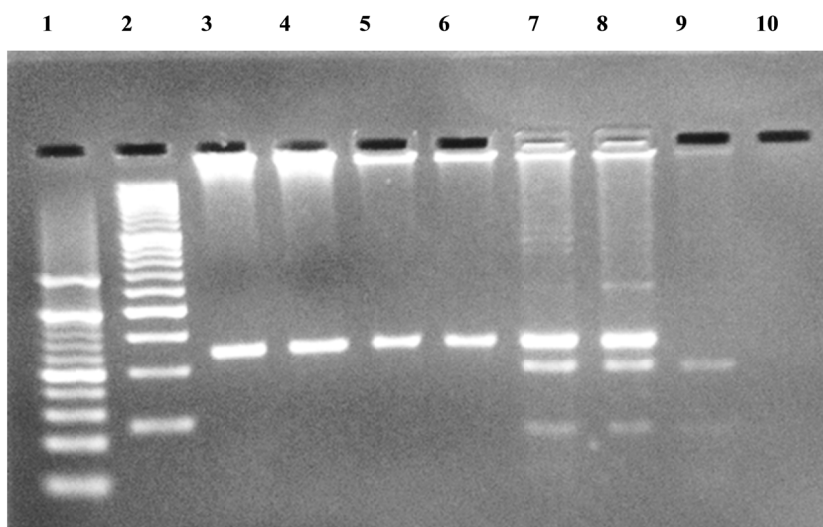


Figure 2. Results of PCR-RFLP analysis for CSN2 gene by DdeI on 3% agarose gel. 1: marker 20 bp DNA ladder; 2: marker 50 bp DNA ladder; 3,4: PCR product (121 bp); 5,6: genotype A1A1 (121 bp); 7,8: genotype A1A2 (121 bp, 86 bp, 35 bp); 9: genotype A2A2 (86 bp, 35 bp); 10: Negative control

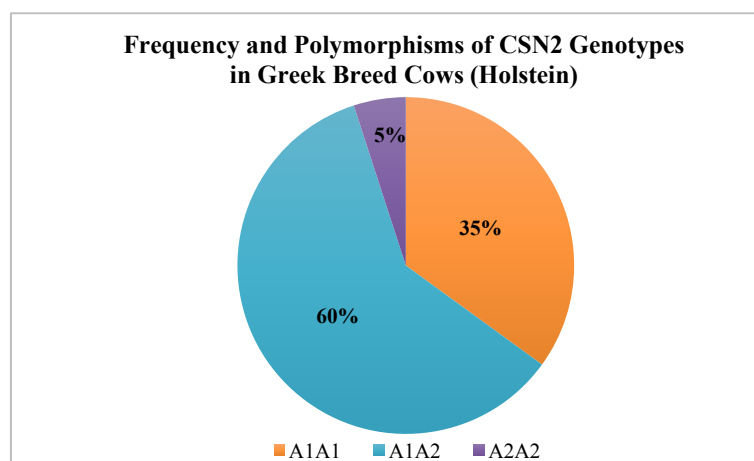


Figure 3. The above figure shows the results obtained in this study. Of the 120 cattle, 72 (60%) showed triple bands of 121 bp, 86 bp, and 35 bp, indicating the A1A2 genotype. 42 (35%) cattle showed a single band at 121 bp, indicating that they carried the A1A1 genotype. The remaining 6 (5%) showed only two bands of 86 and 35 bps, indicating that they carried the A2A2 genotype. In conclusion, the A1A2 genotype predominates over the others

In the amplified product, the restriction site for the enzyme DdeI has been established. The resulting RFLP template from DdeI distinguishes three different band patterns/genotypes: A1A1 (121 bp), A2A2 (86 and 35 bp) and A1A2 (121, 86 and 35 bp). All these bands were clearly visible on 3% agarose gels except the 35 bp band which could not be detected since very small. Allele A1 produced 121 bp fragments, and allele A2 produced a 86 bp and 35 bp fragments as the PCR-RFLP. The genotypic frequencies of these three variants varied across the breeds examined. It was detected 3 genotypes: 2 homozygote genotypes (A1A1, A2A2) and 1 heterozygote genotypes (A1A2). The heterozygote A1A2 genotype was the most frequent. Of the 120 cattle, 72 (60%) showed triple bands of 121 bp, 86 bp, and 35 bp indicating the A1A2 genotype. The 42 (35%) cattle showed a single band at 121 bp, indicating that they carried the A1A1 genotype. The remaining 6 (5%) showed only two bands of 86 and 35 bps each, indicating that they carried the A2A2 genotype.

Frequencies of genotypes and alleles determined in the total population (120 animals) are presented in the Table 1. In the total population heterozygotes A1A2 (0.6) were more predominant compared to homozygotes A1A1 (0.35) and A2A2 (0.05) which were the least frequent ones. As it can be inferred it can be observed two alleles A1 and A2. In the population included in the study the most common allelic variant of beta casein was A1, followed by A2, with frequencies of 0.65 and 0.35, respectively. As shown in Table 2 it is observed high significance at 5% ( $\alpha = 0.05$ ) and one degree of freedom since  $p < 0.01$ .

Table 1. Frequencies of genotypes and alleles of the CSN2 gene in the population of cows

Frequency	Genotypes (n = 120)		
	A1A1	A1A2	A2A2
<b>Absolute</b>	42	72	6
<b>Relative</b>	0.35	0.60	0.05

Table 2. Allele Frequencies and probability (P) chi-square test of genotypes and alleles of the CSN2 gene in the population of cows

Alleles		p-value
A1	A2	
156 (0.65)	84 (0.35)	$3.36 \times 10^{-6}$

Table 3. Occurrence of  $\beta$ -casein gene variants in Holstein-Friesian (HF) in Poland and Red in other countries (data sorted by increasing A1 allele frequency) (Kamiński et al., 2007)

Breed	Country	Allele Frequency of $\beta$ -Casein		N	References
		A1	A2		
HF	Denmark	0.266	0.614	415	(Gustavsson et al., 2014)
	The Netherlands	0.28	0.50	1929	(Visker et al., 2011)
	The Netherlands	0.029	0.69	1629	(Heck et al., 2009)
	Poland	0.32	0.68	177	(Cieślińska et al., 2012)
	Poland	0.35	0.65	650	(Cieślińska et al., 2012)
	Thailand	0.363	0.602	231	(Molee et al., 2011)
	Italy	0.371	0.546	1226	(Massella et al., 2017)
	Italy	0.395	0.57	100	(Chessa et al., 2013)
	Poland	0.40	0.60	143	(Kamiński et al., 2006)
	China	0.432	0.459	133	(Dai et al., 2016)
	Turkey	0.485	0.456	49	(Dinc et al., 2013)
	Iran	0.50	0.50	119	(Gholami et al., 2016)
Red	Sweden	0.48	0.51	392	(Gustavsson et al., 2014)
	Denmark	0.71	0.23	169	(Bech and Kristiansen, 1990)
	Poland	0.53	0.47	201	Present data

Note. Other variants of the  $\beta$ -casein gene are not included.

#### 4. Discussion

Milk contains all the nutrients a young body needs to grow and is particularly rich in proteins and salts. Milk is the only food in nature that contains the high-protein protein known as casein. Casein-derived peptides enhance the body's natural defenses, regulate blood pressure and help fight stress. In the present study, we investigated the polymorphism of exon 7 of the CSN2 gene encoding bovine milk  $\beta$ -casein. Several studies have been conducted linking A1 polymorphism with the production of the opioid peptide BCM-7 associated with human health problems such as type 1 diabetes, autism, schizophrenia and heart disease. In contrast, milk with A2 polymorphism has been found not to cause similar problems in human health (Caroli, Chessa, & Erhardt, 2009; Cieślińska et al., 2015; Elliott et al., 1999; Kumar et al., 2019; McLachlan, 2001; Woodford, 2007, 2011). To draw safe conclusions, the results of the present study were compared with similar studies on  $\beta$ -casein polymorphisms. The results of the present study showed that the A1A2 genotype predominates in relation to the A1A1 genotype. On the contrary, the A2A2 genotype is in a significant minority. Specifically, heterozygous cattle (A1A2) are 72 (60%) with the A2 allele while homozygous cattle (A1A1) with the A1 allele are 42 (35%). Only 6 (5%) showed the A2A2 genotype. The results were that the A1A2 genotype outperformed the others. A similar study was conducted by Kumar et al. (2019) where blood was collected from 429 Frieswal cows. The results were that A1A2 genotype and A2 polymorphism predominated. Maximum A1A2 genotypic frequency was observed in 221 (51.5%) samples, followed by 133 (31%) A2A2 and 75 (17.5%) A1A1 samples. Another survey with similar results was by Sodhi et al. (2018) participated 85 cattle of Ladakh region, India. Results showed that 82 (96.5%) animals carried the A2A2 genotype and the rest (3.5%) A1A2. None of the animals carried the A1A1 genotype. In conclusion, this region is dominated by the A2 allele. The results of the study were similar to those of Miluchová, Trakovická, & Gábor, (2009). A total of 89 cattle were included. The results of the study showed that there were 27 (30.34%) animals with A1A1 genotype, 46 (51.69%) animals with A1A2 and 16 (17.97%) animals with A2A2. Of the entire bovine population, the A2 allele was predominant. Another related study is by Miluchová et al. (2014) where 287 cattle from three different breeds participated. 111 belonged to the Simmental tribe, 89 Pinzgau and 87 Holstein. Results showed that in the Holstein breed, A1A1 genotype was detected in 12 (13.79%) cows, A1A2 genotype in 40 (45.98%) cows and A2A2 genotype in 35 (40.23%) cows. In the Pinzgau breed, the A1A1 genotype was detected in 27 (30.34%) cows, the A1A2 genotype in 46 (51.68%) cows, and the A2A2 genotype in 16 (17.98%) cows. Finally, in the Simmental breed, the A1A1 genotype was detected in 14 (12.61%) cows, the A1A2 genotype in 37 (33.33%) cows and the A2A2 genotype in 60 (54.06%) cows. In conclusion, this study also dominates the A2 allele in all three cattle breeds. Taking into account the results of this study and the numerous studies that have been carried out, it can be concluded that A2 polymorphism predominated over A1 and that the beta-casein polymorphisms in the different cattle breeds are still highly variable, which have not been adequately researched. Of course, to draw safe conclusions a complete mapping of the beta-casein variants would have to be done in order to correlate the various qualitative and quantitative characteristics of the milk and

also to the effects on human health.

## 5. Conclusions

Taking into account the results of the present study as well as studies conducted abroad, we conclude that the beta-casein polymorphisms in bovine breeds vary widely.

To our knowledge, as there is a lack of studies in Greece about the genetic polymorphism of b-casein in cows, this work serves as a preliminary study on the existing genetic polymorphisms of b casein of our local cow breeds. Moreover, this study will play an important purpose in aiding selection of cows with respects to traits of milk quality.

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### Abbreviations

RFLP: restriction fragment length polymorphism;  $\beta$ -LG: beta-lactoglobulin;  $\alpha$ -LA:  $\alpha$ lactalbumin; BSA: bovine serum albumin; LF: lactoferrin; LP: lactoperoxidase; Igs: immunoglobulins; BCM-9: beta-casomorphin-9; BCM-7: beta-casomorphin-7; EDTA: ethylene diamine tetra-acetic acid.

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