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A Stochastic Frontier Approach to Study the Relationship between the Hygienic Quality of Bulk Tank Sheep Milk and Technical Efficiency of the Coagulation Process

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Abstract: Sheep milk from local breeds is important for the production of high-quality cheeses throughout the Mediterranean region, such as Manchego cheese in Spain. To maintain sustainable and efficient production, it is necessary to reach a better understanding of how the composition and hygiene of the milk affect the coagulation process, with the aim of optimizing production yield. This study implemented a stochastic production frontier function to estimate the potential production of curd and efficiency using data from the four seasons of a study of 77 Manchega sheep farms. The Cobb–Douglas production frontier model was estimated using the maximum likelihood estimation method. The results showed that the content of protein, lactose, and fat exhibited increasing returns to scale, with protein content being the most significant factor for curd production. Approximately half of the inefficiency was due to factors related to the technological properties and the hygiene of the milk. The pH, curd firmness, and concentration of lactic acid bacteria improved the efficiency of coagulation, while the concentration of spores of lactate-fermenting *Clostridium* spp., *Pseudomonas* spp., staphylococci, and catalase-negative gram-positive cocci favored the inefficiency of the coagulation process. To date, this is the first study to evaluate the effect of different factors, such as microbial groups, milk composition, and technological properties, on the efficiency of the coagulation process in dairy sheep.

Keywords: coagulation efficiency; stochastic frontier analysis (SFA); Manchego cheese; dairy industry

1. Introduction

In Mediterranean regions, the production of traditional cheeses using milk from local sheep and goat populations plays a significant role in the economy and supporting communities in disadvantaged rural areas [1,2]. The region of Castilla-La Mancha (Spain) is noteworthy, and is renowned for Manchego Cheese, a product covered under a Protected Designation of Origin (PDO), made exclusively with milk from Manchega, probably the most important Spanish native dairy sheep breed [3]. This specific production supports over half a million sheep on 538 farms and aids 72 cheese factories, annually yielding around 17 million kilograms of Manchego cheese [4]. Considering these figures and values of production, it is essential to understand and control milk transformation into

cheese, as impairments in this process can severely compromise the economic sustainability of dairy farms [5].

Milk coagulation is a complex transformation process where curd is obtained primarily from fat and protein as inputs. The interaction of biochemical and physical factors during the coagulation process is decisive in the curd consolidation, establishing the beginning of a series of events that define the curd quality and yield [6]. Optimizing this process also minimizes raw material waste in the whey, thereby contributing to the sustainability of cheese production [7]. To achieve an efficient yield, it is critical to understand and manage critical basic points such as the milk composition, hygienic conditions of the process, and technological factors like coagulation time and curd firmness.

The hygienic and sanitary quality of milk can significantly impact the coagulation process and, consequently, the organoleptic properties of cheese and its final yield [8–10]. Milk with high somatic cell counts—an indicator of mammary infection—tends to coagulate slower and often leads to lower curd yields [11,12]. Furthermore, the microbiological profile of milk is affected by both its native microbiota and other external sources of post-milking contamination, such as environmental conditions at the farm or cheesemaking facilities [13]. These sources comprise various microorganisms, including yeasts, molds, and bacteria. The influence of these microorganisms on the coagulation process of cow milk has been underexplored, and even less is known about their impact on milk from small ruminants [10]. Overall, these studies show a partial vision of milk coagulation, since only a small number of factors are studied together.

The inherently complex process of milk coagulation could be effectively analyzed and optimized through modeling and analysis within stochastic frontier production functions [14,15]. This approach is of interest to the dairy industry, as it facilitates the assessment of the impacts of various factors, such as microbial groups, milk composition, and its technological properties, on the efficiency of the coagulation process. The inclusion of these elements into the analytical framework of stochastic frontier production functions allows for the identification of determinants of efficiency and the optimal conditions to maximize yield, enabling precise adjustments for more efficient and sustainable production of cheese and other derived dairy products [16,17].

Previous studies have used deterministic parametric methods to assess process efficiency, including coagulation [18,19]. However, the introduction of stochastic frontier analysis (SFA) has led to a more nuanced approach [20]. SFA separates deterministic efficiency from stochastic noise, providing deeper insights into process inefficiencies by considering the inherent variability in production. This approach, therefore, offers a more comprehensive understanding of efficiency.

The primary goal of this study was to utilize SFA for modeling and assessing the technical efficiency of the coagulation process of bulk tank milk from Manchega sheep. Considering that bulk tank milk reflects the quality and composition of milk in a cheese-making industrial context, the findings of this study are expected to provide a comprehensive and representative analysis of the actual production conditions. In addition, the present study also aims to quantify the relative contributions of the principal inputs to the coagulation process and to examine the impacts of key technological properties and milk microbial loads on the inefficiency of the coagulation process, which is an important advanced innovation that could be applied in the Manchego cheese industry.

2. Materials and Methods

2.1. Dataset and Sample Collection

This study involved 77 Manchega sheep flocks located in the region of Castilla-La Mancha, Spain. These flocks currently comprise approximately 15% of the farms registered under the PDO "Manchego Cheese" [4]. Each of the studied flocks consisted of a single breed and operated under a semi-extensive production system associated with grazing on natural pastures, residues, and cereal crop remains. The flock sizes varied,

ranging from 150 to 5500 ewes with an average production of 180 kg in 150 days, all of which were mechanically milked. A full description of the farming system can be found in Rivas et al. [21].

Bulk tank samples were collected from each flock at 4 time points (once each season), making a total of 308 samples. These samples were obtained from the tank, composed of a blend of milk from the morning milking of the sampling day and the evening milking of the previous day. The samples were collected into clearly labeled sterile containers, and transported at 4 °C to the Laboratory of Lactology at the Regional Center for Animal Selection and Reproduction (CERSYRA-IRIAF, Valdepeñas, Ciudad Real, Spain) within a maximum timeframe of 2 h. At arrival, the samples were aliquoted and prepared for compositional analysis, microbiological studies, and coagulation tests, all conducted no longer than 48 h after sample collection.

2.2. Laboratory Analysis

A first 50 mL aliquot of each milk sample was sent to the Small Ruminant Dairy Laboratory at the University of Córdoba (Córdoba, Spain) for an analysis of the milk coagulation properties (MCP), which was performed within 24 h after sampling. Rennet clothing time (RCT, min), curd firmness at 60 min (A_{60}, mm), and curd yield (CY, g/100 mL) were determined at 32 °C using a Formagraph lactodinamograph (Foss Electric A/S, Hillerød, Denmark) [18].

A second 50 mL aliquot of milk had azidiol added and was sent to the Interprofessional Dairy Laboratory of Castilla-La Mancha (LILCAM, Talavera de la Reina, Spain) for a milk composition analysis and somatic cell count (SCC), which were carried out within 48 h after sampling. SCC was subsequently expressed as somatic cell scores (SCS) to normalize its distribution by applying a logarithmic transformation [22]. Fat, protein, and lactose contents were determined using a Milkoscan 6000 FT device (Foss Electric, Hillerød, Denmark). SCC was obtained with a Fossomatic FC (Foss Electric, Hillerød, Denmark), and pH was measured using a Crison Basic20 pHmeter (Crison Instruments, Barcelona, Spain).

A third 50 mL aliquot was used for microbiological analyses at CERSYRA-IRIAF, which were performed within 24 h after sample collection. From each sample, serial dilutions were made to inoculate 0.1 mL onto different culture media. Bacterial counts for the following groups of microorganisms were determined on PCA media (Panreac, Spain): standard plate count (SPC) was incubated in aerobic conditions at 30 °C for 72 h; thermoduric bacteria (THERMO) were incubated in the same conditions as SPC, after pasteurizing milk at 62.8 °C for 30 min; and psychrotrophic bacteria (PSYCHRO) were incubated at 6.5 °C for 10 days. *Pseudomonas* spp. (PSEUDO) were cultured on Cetrimide agar (Panreac, Spain) and incubated at 35 °C for 48 h. The determination of coliforms (COLI) was conducted using CromoIDTM Coli medium (bioMérieux, Spain), incubated at 37 °C for 24 h. Gram-positive catalase-negative cocci count (GPCNC) was determined in modified Edwards medium with colistin and oxolinic acid supplement (Oxoid, UK), incubated at 35 °C for 48 h. Lactic acid bacteria (LAB) were seeded on MRS medium (Panreac, Spain) acidified to pH 5.7, and incubated at 30 °C for 72 h. Lactate-fermenting *Clostridium* spores (LFCS) count was performed using the most probable number (MPN) technique, in Bryant and Burkey Broth (BBB, Merck, Germany). For the enumeration of coagulase-positive staphylococci (CPS) and coagulase-negative staphylococci (CNS), Baird Parker RPF Agar medium (bioMérieux, Spain) was used, and incubated at 37 °C for 24 h. A full description of the microbiological analysis can be found in a previous study from our research group [10]. The microbial counts were subsequently subjected to a decimal logarithmic transformation to normalize their distribution [10].

2.3. Stochastic Frontier Analysis of the Milk Coagulation Process

2.3.1. Theoretical Concept

In 1977, several authors [20,23] formulated a stochastic frontier production function that can be specified for panel data as [24]:

$$Y_{it} = e^{X_{it}\beta + V_{it} - U_{it}} \tag{1}$$

where Y_{it} denotes the output of the *i-th* unit (*i = 1, …, N*) in the *t-th* time period (*t = 1, …, T*), X_{it} is a *k × 1* vector of input quantities used by the *i-th* unit in the *t-th* time period, β is a k × 1 vector of coefficients to be estimated, and V_{it} and U_{it} are components of the error term assumed to be independent. The first component, V_{it}, is a normally distributed random variable with a zero mean and variance σ_v^2, accounting for measurement errors and other random factors. The second component, U_{it}, is a non-negative random variable that measures the deviation from the efficient frontier for the *i-th* observation, derived from the normal distribution truncated at zero, with a mean $Z_{it}\delta$ and variance σ^2. Z_{it} is a *1 × m* vector of explanatory variables associated with technical inefficiency over time, and δ is an *m × 1* vector of unknown coefficients [25].

The formulation for inefficiency effects in the panel data model [24] allows for the joint estimation of the stochastic production function, and the model for inefficiency effects linked to this function is presented as follows:

$$U_{it} = Z_{it}\delta + W_{it} \tag{2}$$

where U_{it} represents the estimated one-sided inefficiency of unit *i* in time period *t*, Z_{it} is the set of variables explaining the inefficiency of unit *i* in period *t*, δ is a set of coefficients estimated in the inefficiency model, and W_{it} is defined by the truncation of the normal distribution with a mean zero and variance σ^2.

The stochastic production function (1) and the inefficiency effects model (2) are estimated simultaneously using the maximum likelihood method. The technical efficiency (TE) estimates for unit *i* in time period *t* are presented as follows:

$$TE_{it} = \frac{E(Y_{it}^*/u_{it}, X_{it})}{E(Y_{it}^*/u_{it} = 0, X_{it})} = e^{-u_{it}} = e^{-z_{it}\delta - w_{it}} \tag{3}$$

where Y_{it}^* is the production, which is equal to $e^{Y_{it}}$ when expressed in logarithms. Therefore, TE is calculated as the ratio of the level of production obtained with respect to the maximum achievable production given the quantities of the inputs (i.e., when $u_{it} = 0$). The value of TE ranges from 0 to 1, with the latter being the most favorable.

2.3.2. Empirical Model

The empirical analysis was based on the estimation of a Cobb–Douglas production function, in which both production and inputs are expressed in logarithmic form. Therefore, the estimated coefficients reflect the production elasticities [26]. A translog function was also estimated, whose preliminary results led to the rejection of the functional form [27–29].

Each of the 77 Manchega sheep farms was considered as a production unit, and each of the seasons (spring, summer, autumn, and winter) as a time period. The model assumes that the production of curd (CY, g/100 mL) is a function of three inputs, fat (FAT, g/100 mL), protein (CP, g/100 mL), and lactose (LAC, g/100 mL), expressed as:

$$LnY_{it} = \beta_0 + \sum_{k=i}^{3} \beta_k LnX_{kit} + V_{it} - U_{it} \tag{4}$$

where Y_{it} is the total curd production from a milk sample of farm *i* in season *t*, X_{1-3} are the three previously defined variable inputs of farm *i* in season *t*, V_{it} is the random noise, and U_{it} is the inefficiency term.

It is assumed that inefficiency follows a half-normal distribution, with the inefficiency model being specified as:

$$U_{it} = \delta_0 + \sum_{k=1}^{14} \delta_k Z_{kit} \tag{5}$$

where Z is the explanatory variables (Table 1) and δ is a set of parameters to be estimated. The estimation of Equations (5) and (6) is carried out simultaneously by maximum likelihood [24] using the program FRONTIER 4.1. [30].

Table 1. Descriptive statistics for variables in the stochastic frontier production function and variables in the inefficiency equation.

Variable	Symbol	Mean	SD	Minimum	Maximum
CY [1]	Y	31.52	4.88	20.10	44.70
FAT [1]	X_1	7.87	0.82	5.74	10.38
CP [1]	X_2	6.05	0.60	5.03	8.05
LAC [1]	X_3	4.60	0.24	3.71	5.16
pH	Z_1	6.61	0.15	6.00	7.00
RCT [2]	Z_2	31.25	12.12	4.45	50.10
A_{60} [3]	Z_3	33.24	10.49	1.66	54.90
SCS [4]	Z_4	6.40	0.77	4.44	8.90
LFCS [5]	Z_5	3.41	0.65	0.00	5.04
LAB [5]	Z_6	4.89	0.72	2.91	6.48
THERMO [5]	Z_7	3.23	0.87	0.00	5.48
PSYCHRO [5]	Z_8	4.51	2.08	0.00	8.00
PSEUDO [5]	Z_9	3.12	0.85	0.00	4.53
SPC [5]	Z_{10}	5.65	0.83	0.00	8.01
GPCNC [5]	Z_{11}	4.10	0.93	0.00	5.48
COLI [5]	Z_{12}	3.13	0.98	0.00	5.48
CPS [5]	Z_{13}	2.46	1.41	0.00	5.48
CNS [5]	Z_{14}	4.25	0.71	0.00	5.48

[1] g/100 mL; [2] min.; [3] mm; [4] SCS = $\log_2$(SCC/100,000) + 3/mL; [5] $\log_{10}$CFU/mL; CY = curd yield; FAT = fat; CP = protein; LAC = lactose; A_{60} = curd firmness at 60 min; RCT = rennet clotting time; SCS = bull tank milk somatic cell score; LFCS = lactate-fermenting *Clostridium* spores; LAB = lactic acid bacteria; THERMO = thermodurics, PSYCHRO = psychrotrophs, PSEUDO = *Pseudomonas* spp.; SPC = total mesophilic bacteria (standard plate count); GPCNC = gram-positive catalase-negative cocci; COLI = coliforms other than *Escherichia coli*, CPS = coagulase-positive staphylococci, and CNS = coagulase-negative staphylococci.

The existence of inefficiency (H0: $\gamma = \delta_0 = \delta_1 = ... = \delta_{14} = 0$), the relevance of exogenous variables in explaining the inefficiency component (H0: $\delta_0 = \delta_1 = ... = \delta_{14} = 0$), the existence of stochastic efficiency (H0: $\gamma = 0$), and the assumption of the truncated normal distribution of the inefficiency component (H0: $\mu = 0$) were assessed using the generalized likelihood ratio statistic (λ), defined as [27–29]:

$$\lambda = -2\{ln[L(H_0)]/ln[L(H_1)]\} \tag{7}$$

where ln[(H0)] corresponds to the value of the log-likelihood function for the restricted model (specified in the null hypothesis) and ln[(H1)] is the value of the log-likelihood function for the general model stipulated in the alternative hypothesis. This test is asymptotically distributed as a chi-squared distribution with degrees of freedom equal to the difference in the number of parameters estimated under both hypotheses [30].

Finally, the milk samples were classified according to the TE percentile into three groups: low (<P15), medium (P15 to P85), and high (>P85). The three efficiency groups were compared using simple ANOVA and the SNK test. Additionally, the association between TE and the season of the year was analyzed using the same methods (ANOVA and SNK test). Statistical analyses were performed using the software XLSTAT v.19.4 [31].

3. Results and Discussion

Table 2 presents the results from the estimation of the stochastic frontier production function for the coagulation of Manchega sheep milk, using a Cobb–Douglas production model. All obtained β coefficients were different from zero ($p < 0.05$). The highest output elasticity was for protein, at 0.94, implying that a 1% increase in milk protein content would increase curd production by 0.94%. The lowest output elasticity corresponded to lactose content, while fat content had an elasticity of 0.46.

Table 2. Maximum-likelihood estimates parameters for the stochastic frontier production function and inefficiency function equation.

Variable	Parameter	Coefficient	SE	t-Ratio	p-Value
Constant	β_0	0.103	0.176	0.59	0.336
Ln(FAT) [1]	β_1	0.463	0.064	7.20	<0.001
Ln(CP) [1]	β_2	0.941	0.112	8.39	<0.001
Ln(LAC) [1]	β_3	0.429	0.148	2.89	0.006
Inefficiency model					
Constant	δ_0	0.734	0.254	2.88	0.007
pH	δ_1	−0.128	0.039	−3.27	0.002
RCT [2]	δ_2	0.000	0.000	0.33	0.378
A_{60} [3]	δ_3	−0.001	0.001	−2.29	0.029
SCS [4]	δ_4	−0.007	0.007	−1.05	0.229
LFCS [5]	δ_5	0.009	0.005	2.39	0.023
LAB [5]	δ_6	−0.022	0.008	−2.69	0.011
THERMO [5]	δ_7	−0.003	0.007	−0.48	0.356
PSYCHRO [5]	δ_8	0.001	0.003	0.43	0.363
PSEUDO [5]	δ_9	0.025	0.010	2.47	0.020
SPC [5]	δ_{10}	0.011	0.010	1.17	0.200
GPCNC [5]	δ_{11}	0.014	0.008	2.16	0.039
COLI [5]	δ_{12}	−0.003	0.006	−0.48	0.355
CPS [5]	δ_{13}	0.005	0.003	2.11	0.043
CNS [5]	δ_{14}	0.025	0.013	2.22	0.034
Error variance	σ_e	0.002	0.000	5.44	<0.001
Variance	γ	0.508	0.167	3.04	0.004
Log-likelihood value	-	492.77			

[1] g/100 mL; [2] min.; [3] mm; [4] SCS = $\log_2$(SCC/100,000) + 3/mL; [5] $\log_{10}$CFU/mL; FAT = fat; CP = protein; LAC = lactose; A_{60} = curd firmness at 60 min.; RCT = rennet clotting time; SCS = bulk tank milk somatic cell score; LFCS = lactate-fermenting *Clostridium* spores; LAB = lactic acid bacteria; THERMO = thermodurics, PSYCHRO = psychrotrophs, PSEUDO = *Pseudomonas* spp.; SPC = total mesophilic bacteria (standard plate count); GPCNC = gram-positive catalase-negative cocci; COLI = coliforms other than *Escherichia coli*, CPS = coagulase-positive staphylococci, and CNS = coagulase-negative staphylococci.

The elasticity of milk components is influenced by a range of factors, both environmental (e.g., diet) and intrinsic (e.g., genetics of the breed). Fat content in milk can be adjusted through dietary modifications, but milk protein shows a more pronounced genetic dependence [32,33]. Considering the significant role of protein in enhancing the efficiency of coagulation, prioritizing its improvement through genetic selection programs is deemed essential. The necessity for such targeted improvements is underscored by the lack of research in this area, with only a few studies addressing this topic [19]. Furthermore, expanding this line of research to include other breeds, species, and production systems would be valuable, as it would provide a more comprehensive understanding of the efficiency of the coagulation process, encompassing both technical and economic aspects.

The sum of all output elasticities was 1.83, indicating that, on average, the Manchega dairy sheep system has increasing returns to scale. This means that if the fat, protein, and lactose contents in Manchega sheep milk were to increase by 1%, there would be a 1.83% rise in curd yield, leading to financial benefits for the industry. It is important to consider the negative correlation that usually exists between lactose content and fat and protein content, due to the role of lactose in the regulation of milk volume, which can lead to a dilution effect of the other major milk components. This fact could pose a challenge to achieving an effective increase in scale performance [34]. From a practical perspective, despite this negative correlation, an effective strategy could be a combination of milks with different fat, protein, and lactose contents to try to achieve an optimal balance by taking advantage of variations in population. An assessment of the technical and economic feasibility of this approach would be of interest.

On the other hand, milk quality payment systems often equally value fat and protein content, considering the sum of both ("cheese extract") or total solids. However, the findings of this study suggest that protein content has a greater importance than fat content regarding curd yield in Manchega sheep. Therefore, it is important to conduct a more detailed economic assessment to adapt these discoveries to the pricing system. Additionally, it should be noted that this study did not take into account the composition of the protein, meaning that the impact of casein was not distinguished from that of whey proteins. Such an adjustment would allow for a fairer and more accurate reward for producers, based on the true cheesemaking yield of the supplied milk. This reassessment is essential not only for reasons of social equity, but also to enhance the economic efficiency of the dairy sector. By better aligning financial rewards with factors that genuinely improve the quality and yield of cheese production, producers would be incentivized to optimize milk quality, thus promoting continuous improvement in the sector [35]. This more exhaustive and quality-based approach could encourage innovations in farm management and feeding practices, which, in turn, could lead to more sustainable and profitable long-term dairy production.

The average TE for coagulation was 0.95, ranging from 0.86 to 0.99 (Figure 1). Therefore, the average inefficiency in curd production was 0.05 (1-TE), which translates to a mean marginal loss of 52.6 g of curd per kilogram of curd produced, according to the average composition shown in Table 1. Previous studies have reported lower TE values, although their results are not entirely comparable, as they modeled a Cobb–Douglas function with two predictors (fat and protein content) using ordinary least squares (OLS) and assessed efficiency through a deterministic frontier [19].
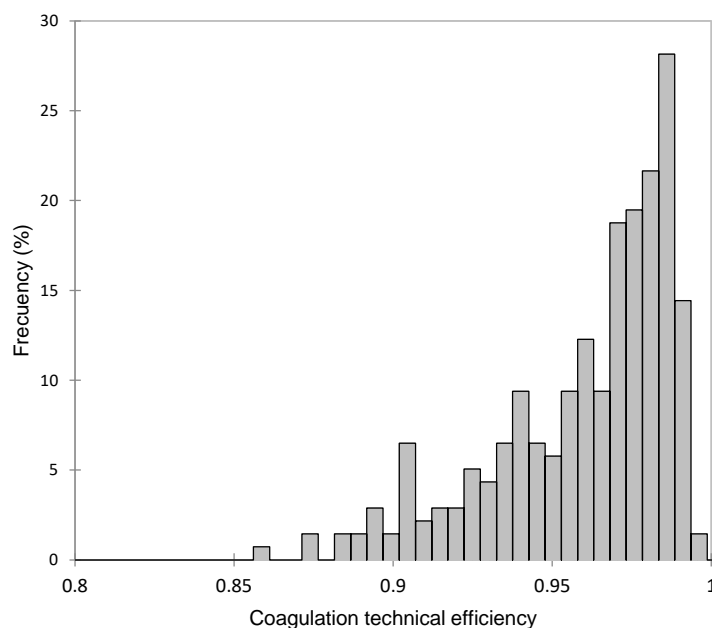
Figure 1. Frequency distribution of coagulation technical efficiency (TE) in Manchega sheep.

The tests on the specifications of the model for technical inefficiency led to the rejection of all null hypotheses under consideration ($p < 0.05$). Therefore, it is confirmed that an average production function constitutes an inadequate representation of the data, the necessity to incorporate technical inefficiency in the production function, the significance of the variables that explain technical inefficiency, and the fit to the truncated normal of the inefficiency component.

The variance parameter γ was 0.51, indicating that half of the variation in the error of the function was due to the inefficiency error u_{it}, while the other half was due to the stochastic random error v_{it}.

Table 2 presents the technical inefficiencies identified in the model. Positive parameter estimates indicate relative technical inefficiency, while negative ones signify relative technical efficiency. pH and A_{60} were statistically significant in the inefficiency model, suggesting that coagulation efficiency improves with an increased curd firmness and higher pH levels.

Ten groups of microorganisms were considered in the inefficiency model. Of these, six were statistically significant in their effect on the (in)efficiency of the coagulation process: LFCS, LAB, PSEUDO, GPCNC, CPS, and CNS. The concentration of these groups of microorganisms, with the exception of LAB, was associated with an increase in the inefficiency of the coagulation process. In contrast, a higher concentration of lactic acid bacteria (LAB) was associated with a greater efficiency in the process, suggesting a positive effect of this group of microorganisms on coagulation.

The more pronounced the magnitude of efficiency or inefficiency, the greater the deviation of the estimated value from zero. A detailed examination of each parameter estimate reveals noteworthy insights. Notably, pH emerged as the most influential factor in the inefficiency model, showing a positive impact on the efficiency of the process. This relationship is also reflected in Table 3, where three levels of efficiency are differentiated based on the 15th and 85th percentiles. An increase in average pH is observed, going from 6.49 in the least efficient group (15th percentile) to 6.74 in the most efficient group (85th percentile). pH is related to the coagulation process [36]: acidic pHs are associated with quicker coagulations, whereas pHs leaning towards alkaline tend to slow down the process, resulting in firmer curds [37], which, according to our findings, lead to a greater coagulation efficiency due to improved solid retention [38].

Table 3. Comparison between the three coagulation efficiency groups using ANOVA.

Variable	Low (<P_{15})	Medium (P_{15} to P_{85})	High (>P_{85})	SEM	*p*-Value
TE (%)	90.58 [a]	96.52 [b]	98.90 [c]	0.01	<0.001
CY (g/100 mL)	25.62 [a]	31.84 [b]	36.25 [c]	0.29	<0.001
FAT (g/100 mL)	7.72	7.86	7.96	0.05	0.410
CP (g/100 mL)	5.92	6.07	6.16	0.03	0.086
LAC (g/100 mL)	4.65	4.59	4.62	0.01	0.414
pH (−log [H+])	6.49 [a]	6.60 [b]	6.74 [c]	0.01	<0.001
RCT (min)	24.90 [a]	32.21 [b]	33.20 [b]	0.73	0.001
A_{60} (mm)	30.25 [a]	33.13 [ab]	35.86 [b]	0.64	0.035
SCS [1]	6.37	6.46	6.47	0.05	0.657
LFCS [2]	3.30	3.38	3.49	0.04	0.422
LAB [2]	4.41 [a]	4.87 [b]	5.47 [c]	0.04	<0.001
THERMO [2]	3.51	3.23	3.11	0.05	0.077
PSYCHRO [2]	5.56 [b]	4.30 [a]	4.59 [a]	0.13	0.002
PSEUDO [2]	4.02 [c]	3.13 [b]	2.22 [a]	0.05	<0.001
SPC [2]	5.60 [a]	5.63 [a]	5.91 [b]	0.05	0.138

GPCNC [2]	4.87 c	4.12 b	3.48 a	0.05	<0.001
COLI [2]	3.76 c	3.08 b	2.76 a	0.06	<0.001
CPS [2]	2.81	2.42	2.36	0.08	0.236
CNS [2]	4.68 c	4.27 b	3.79 a	0.04	<0.001

[1] SCS = $\log_2$(cells/100,000) + 3; [2] $\log_{10}$UFC; TE = technical efficiency; CY = curd yield; FAT = fat; CP = protein; LAC = lactose; A_{60} = curd firmness at 60 min.; RCT = rennet clotting time; SCS = bulk tank milk somatic cell score; LFCS = lactate-fermenting *Clostridium* spores; LAB = lactic acid bacteria; THERMO = thermodurics, PSYCHRO = psychrotrophs, PSEUDO = *Pseudomonas* spp.; SPC = total mesophilic bacteria (standard plate count); GPCNC = gram-positive catalase-negative cocci; COLI = coliforms other than *Escherichia coli*, CPS = coagulase-positive staphylococci, and CNS = coagulase-negative staphylococci. a–c: Means without a common superscript are statistically different.

Although RCT was not found to be significant in the inefficiency model, Table 3 shows an average increase in both RCT and A_{60} values from the lowest efficiency group to the highest. This indicates that an optimal pH is crucial for improving the TE of milk coagulation by influencing both the physical properties of the curd and its interaction with the microbiota. This approach confirms the idea that, in order to optimize the quality and efficiency of cheese production, it is necessary to consider milk physicochemical and microbiological attributes.

PSEUDO and CNS were the bacterial groups with the most negative impact on the milk coagulation process (Table 2). CNS is a group that includes a range of microorganisms that typically cause subclinical intramammary infections in sheep, characterized by moderately elevated but persistent cell counts, mainly affecting animals with high productivity [39]. Such infections can lead to minor changes in milk composition, like alterations in protein levels and somatic cell count, which can negatively influence curd quality and yield [10,40].

PSEUDO are ubiquitous microorganisms capable of surviving and proliferating at low temperatures, often associated with poor hygiene conditions, and spread during extended periods of milk storage [41,42]. Specifically, the presence of enzymes such as proteases and lipases, produced by some members of this group, is particularly problematic for cheesemaking. These enzymes break down the fat and protein in milk, potentially altering the structure and integrity of the curd. This not only affects the texture and quality of cheese, but can also reduce the curd yield due to a lower retention of solids and essential nutrients [10,43]. Although the PSYCHRO group was not significantly relevant in the inefficiency model, Table 3 indicates a decrease in the average concentration of these microorganisms when moving from low to high efficiency groups.

The GPCNC group exhibited an adverse effect on the efficiency of the coagulation process, though its impact was moderate compared to other microbial groups. This group includes various bacterial species that are indicators of poor hygiene conditions in the milk production environment, as well as mammary health issues in the breed [13,44,45]. From a coagulation perspective, these microorganisms can negatively impact the process by altering the concentration of minerals, the balance of proteins and fats, and enzymatic activity, which, in turn, can influence the formation and texture of the curd [10,46].

LFCS also showed a moderate negative effect on the efficiency of Manchega sheep milk coagulation. Furthermore, these microorganisms cause late blowing in pressed cheese, leading to the formation of cracks and cavities due to acid-butyric fermentation by vegetative cells once the sporulated forms germinate inside the cheese, causing significant economic losses for the cheese industry. These microorganisms are primarily introduced into the milk through silage and other by-products used in livestock feed, as well as by poor hygiene in the milking parlor [47]. Additionally, a correlation has been described between high spore counts and the technological characteristics of the milk, with increases in coagulation time and curd firming time [10]. Therefore, the presence of LFCS is a critical factor that must be controlled to ensure quality and efficiency in the production of fermented dairy products.

The CPS group also showed a minor negative impact on the efficiency of the milk coagulation process. *Staphylococcus aureus* is one of the main pathogens causing clinical mastitis in dairy ruminants [48,49]. Its significance extends beyond animal health, also impacting public health, as they are known for producing thermostable toxins [50,51]. From the perspective of milk coagulation, the presence of CPS can interfere with the normal process due to several factors. Firstly, mammary infections caused by these organisms can alter the chemical composition of the milk, affecting its ability to form an adequate curd due to the presence of enzymes from the plasmin–plasminogen complex associated with high somatic cell counts [52,53]. Although their impact on coagulation efficiency is not as marked as other microorganisms, the presence of coagulase-positive staphylococci is an important factor to consider in managing the quality and safety of dairy products. Their control is essential not only for maintaining production efficiency, but also for ensuring the safety of the final product.

On the contrary, LAB were revealed in the inefficiency model as a factor with a significant positive influence on the TE of the coagulation process. This finding is consistent with previous expectations, given the known beneficial role of LAB in lactic fermentation and in the production of dairy products, providing differential organoleptic and sensory characteristics to cheeses [54,55]. In addition, LAB contribute to the inhibition of other undesirable microorganisms present in raw milk [56,57]. Therefore, their presence helps to maximize curd efficiency and cheese production. It is important to consider that expecting both alkaline pH and LAB to improve coagulation performance may seem contradictory. It is crucial to note that the initial pH of milk, ranging between 6.00 and 7.00, according to Table 1, does not show more acidic values indicative of the massive proliferation of LAB. The acidifying capacity of LAB depends on the strain, as well as its homo- or heterofermentative aptitude [58,59]. Moreover, the microbiome of raw sheep milk is highly complex [60], and the behavior of LAB may differ from that of commercial LAB starters, which are designed, among other aspects, to have a high acidifying capacity. It could be of interest to explore whether there are specific conditions under which alkaline pH and LAB could synergistically interact to improve coagulation performance, even within a range of initial milk pH close to neutrality.

The TE showed significant variations depending on the season, increasing in autumn and reaching its highest values in winter, then decreasing in spring and recording the lowest values in summer, as observed in Figure 2. This variation is even more important, because the production of Manchega sheep milk does not have large seasonal fluctuations as occurs in other dairy breeds such as Sarda [61] or Latxa [62]. Therefore, this seasonal variability could well be due to climatic conditions, which selectively affect contamination by different groups of microorganisms [63–65] or in the composition of the milk due to aspects related to grazing, types of forage and preserved foods, periods of stabling, ventilation, and other related factors [66].
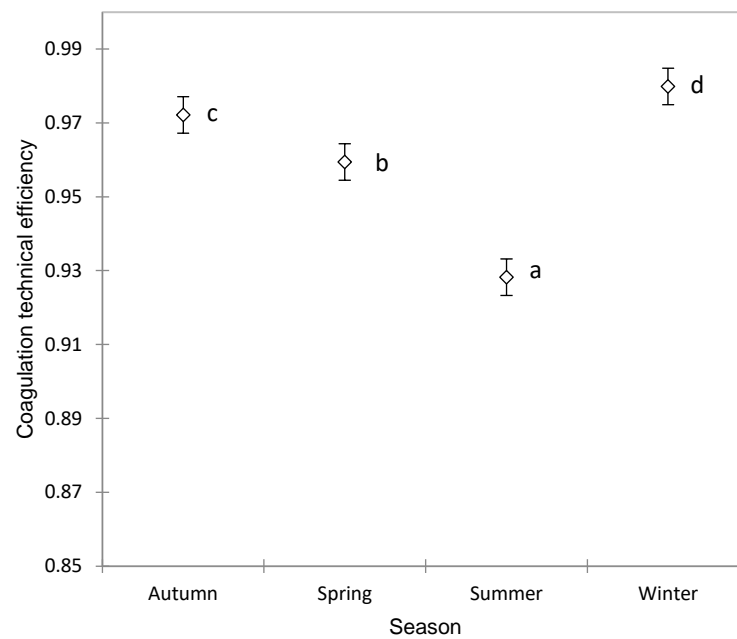
Figure 2. Association between the season and coagulation technical efficiency in Manchega sheep using ANOVA (mean ± standard error). Means without a common superscript (a–d) are statistically different (Student–Newman–Keuls, SNK *p* < 0.05).

4. Conclusions

This study implements a Cobb–Douglas stochastic production frontier function to estimate the potential production and technical efficiency of curd production from Manchega sheep milk. Using data from 77 farms, this study details, for the first time, the determinants of potential yields for the dairy sheep industry under different production constraints using a stochastic approach. The main findings of this study are: (1) empirical results showing that the Cobb–Douglas stochastic production frontier function model fits the data better than the translog specification; (2) curd production shows an increasing returns to scale, meaning a 1% increase in all input factors would result in almost 2% increase in production; (3) this study estimates substitution elasticities to identify that milk protein content is the most relevant input for curd production; and (4) approximately half of the inefficiency was due to factors related to the technological properties and hygiene of the milk. The pH, curd firmness, and concentration of lactic acid bacteria improve the efficiency of coagulation, while the concentration of spores of lactate-fermenting *Clostridium* spp., pseudomonas, staphylococci, and catalase-negative gram-positive cocci favor the inefficiency of the coagulation process.

Author Contributions: Conceptualization, J.M.P. and R.A.; methodology, J.M.P.; software, J.M.P.; formal analysis, J.M.P. and E.A.; investigation, L.J., J.C.-V., B.O. and R.A.; resources, L.J., J.C.-V. and R.A., writing—original draft preparation, L.J and J.M.P.; writing—review and editing, J.C.-V., A.C., N.A. and B.O.; visualization, L.J. and J.M.P.; supervision, R.A.; project administration, J.M.P. and R.A.; funding acquisition, J.M.P. and R.A. All authors have read and agreed to the published version of the manuscript.

Institutional Review Board Statement: Ethical approval was not required for this project, since all samples were bulk tank milk.

Informed Consent Statement: Not applicable.

Data Availability Statement: The data presented in this study are available on request from the corresponding author (accurately indicate status).

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