

Studies pertaining to coagulant and proteolytic activities of plant proteases from *Cynara cardunculus*

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Abstract

Studies encompassing variation of milk clotting time with the concentration of proteases extracted from *Cynara cardunculus* were performed; milk clotting time dependence is not linear, and a model was postulated that fits well the experimental data obtained, and hence may be useful in predicting changes during the cheesemaking process. Parallel studies were also conducted pertaining to proteolysis of a mixture of ovine and caprine caseins, in attempts to investigate the stability of the aforementioned coagulating enzymes in crude or in pure form, with or without previous incubation for a certain time under typical ripening conditions. The enzymes exhibited an increase in activity whenever previous incubation had taken place. Moreover, the extent of enzyme-mediated proteolysis was always higher on caprine than on ovine milk caseins.

Introduction

Rennet substitutes of plant origin have been increasingly used to manufacture cheese, especially at the artisanal level. Application of plant coagulants allows target cheese production, and hence contributes to improve the nutritional input of those populations on whom restrictions are imposed by use of animal rennets (Gupta & Eskin, 1977). Several plant preparations have been shown (Aworth & Muller, 1987; Domingos et al., 1997; Gupta et al., 1977; Edwards & Kosikowski, 1983; Padmanabhan, Chitre, & Shastri, 1993; Pozsar-Hajnal, Vamos-Vigyazo, Nonn-Sas, & Hegedues-Voelgyesi, 1969; Singh, Chander, Bhalerao, & Dastur, 1973; Tamer, 1993) to clot milk; however, the majority proved unsuitable for cheese production owing to their excessively proteolytic character – which lowers cheese yield and produces bitter flavours in the final cheese. Moreover, their toxin content constrains their use as food. Among those that have been reported to produce sat-

isfactory final products, the proteases present in the flowers of *Cynara cardunculus* deserve special mention; aqueous extracts thereof have been used for centuries in the Iberian Peninsula for the manufacture, at the farm level, of ovine and/or caprine milk cheeses (Pires et al., 1994; Trujillo, Carretero, & Guamis, 1994; Vieira de Sá & Barbosa, 1972). The clotting activity of such extracts is accounted for by two aspartic proteases, cardosins A and B, which have been shown to act on κ -casein in a way similar to that of chymosin (Macedo, Faro, & Pires, 1993), but which can also hydrolyse caseins other than κ -casein (Silva & Malcata, 1998; Silva & Malcata, 1999).

Cheese production starts with milk coagulation, which involves the enzyme-mediated cleavage of κ -casein at the peptide bond Phe105–Met106; this chemical step renders the casein micelles unstable, and thus eventually causes cooperative aggregation to yield a clot – which is then ripened and gives rise to the final cheese. A few kinetic models were proposed (Hyslop, Richardson, & Ryans, 1979; Kopelman & Cogan, 1976; van Hooydonk & Walstra, 1987) for milk coagulation by rennet, but information is still scarce concerning the kinetics of milk coagulation by enzymes of plant origin; as of today, there is indeed no model whatsoever,

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proposed and tested for milk coagulation by cardosins A and B, either in a mixture or isolated. Such modelling would help, not only in attempts to illuminate the contribution of each cardosin to the general milk-clotting activity of that plant coagulant, but also for optimising clotting process so as to mechanistically predict its evolution as a function of a number of processing parameters.

Milk clotting enzymes affect final cheese quality via their hydrolytic action on α_s -, β - and κ -caseins, even though only a residual amount of enzyme is retained in the curd after syneresis. Manufacturing and ripening conditions influence the protein degradation pattern of cheese insofar as they constrain the activity/stability of enzymes in terms of prevailing environmental conditions (e.g. pH, a_w and temperature).

The major aim of this work was to generate experimental data on milk coagulation by cardosins A and B, and to examine the fit of existing mathematical models to these data – so as to ascertain whether milk clotting time by cardosins might be predictable. The integrity (in terms of kinetic stability) of these plant enzymes was also a secondary aim. A comparison was made with a mixture of both enzymes, in attempts to reveal interactions and to better understand what happens when such a coagulant is used in traditional cheesemaking.

Materials and methods

Enzyme source

Cardosins were extracted from the dried flowers of *C. cardunculus*, and purified by gel filtration followed by ion-exchange chromatography, according to the method described by Silva, Allmere, Malcata, and Andr n (2003). The lyophilized enzymes were reconstituted in citrate buffer (pH 6.2) just prior to use, or alternatively reconstituted in 10 mM citrate buffer (pH 5.0) containing 20 g/l of NaCl, at 10  C for 20 h, so as to mimic actual cheese ripening conditions.

Protein content determination

The protein concentration of each enzyme solution was determined by the phenol reagent method for biological fluids, using the micro protein determination method (Sigma, St. Louis MO, USA).

Coagulating activity determination

Milk clotting activity was determined following the procedure described by IDF (1992). Low-heat skim milk powder (NILACTM) (NIZO, Ede, The Netherlands) was reconstituted in 10 mM aqueous CaCl₂ (pH 6.5) to a final concentration of 0.12 kg/l. Enzyme extracts were

added at a rate of 0.1 ml per ml of milk. The coagulation point was determined by manual rotating of the test tube periodically, at short time intervals, and checking for visible clot formation. One rennet unit (R.U.) was defined as the amount of protein that coagulates 10 ml of reconstituted low-heat skim milk powder at 30  C in 100 s (Berridge, 1945). Standard solutions of each enzyme were prepared by diluting the enzymes in ultra-pure water.

Proteolytic activity determination

Ovine (or caprine) casein (SIGMA), at the 1% (w/v) level, was subjected to hydrolysis at 30  C in 10 mM citrate buffer (pH 6.2). Sodium azide was added at 0.03% (w/v) to prevent microbial growth. The hydrolysis was initiated by addition of 120  l of each enzyme solution (cardosin A, cardosin B, or a mixture of both enzymes) to 3 ml of substrate solution. At selected times (5 min, 1, 3, 6 and 10 h), aliquots were duly taken, and the reaction was quenched by heating at 100  C for 5 min. The proteolytic activity was quantified by evaluating the peptides soluble in aqueous 5% (w/v) trichloroacetic acid (TCA); aliquots of each sample were treated with 5% (w/v) TCA at a volumetric ratio of 1:2; the mixture was allowed to settle for 10 min, and then centrifuged at 12,000 g for 10 min. The absorbance of the supernatant was measured at 280 nm.

Mathematical and statistical analyses

Coagulating activity

The experimental data pertaining to variation of clotting time with enzyme concentration were examined by non-linear regression analysis, using the Statistica software (v. 5.1, Statsoft, Tulsa OK, USA), based on the model proposed by Payens, Wiersma, and Brinkhuis (1977).

$$tE_0^\gamma = C, \quad (1)$$

where t denotes clotting time and E_0 total enzyme concentration, and where γ and C are adjustable parameters.

Diagnostics of residuals of the data fitted to by the aforementioned model (not shown) were performed, in order to check the validity of the statistical analyses; since no biased trends were detected, no data transformation was required, nor was transformation of the functional form of the model warranted, so Eq. (1) was assumed to be appropriate for fitting.

Proteolytic activity

Assuming that each substrate protein molecule (denoted as A) is hydrolysed into two peptide molecules (denoted as B), then the mass balance prevailing during hydrolysis and based on said stoichiometry is:

$$2(C_{A0} - C_A) = C_B, \quad (2)$$

where C_x is the concentration of entity x , and subscript 0 denotes initial conditions; for simplicity, the initial concentration of product B was taken as nil.

The simplest mechanistic equation for this event can thus be written as



which, assuming it represents an elementary chemical step, leads to

$$r_A = -kC_A, \quad (4)$$

where r_x is the rate of production of entity x , and k denotes a first-order kinetic constant. Integration of Eq. (4) in a batch system, the mass balance of which is characterized by

$$\frac{dC_A}{dt} = -r_A, \quad (5)$$

leads to

$$C_A = C_{A0}e^{-kt}, \quad (6)$$

where t is the hydrolysis time.

Recall that the proteolytic activity is measured via quantification of the peptides soluble in 5% (w/v) TCA, which is based on a correlation with absorbance that is the sum of the absorbances (Abs) given by the intact caseins and the peptides formed so far, i.e.

$$\text{Abs}(t) = (\varepsilon_A LC_A) + (\varepsilon_B LC_B) \quad (7)$$

where ε_A and ε_B denote the molar absorption coefficients for the substrate and the product, respectively, and L denotes the absorption path length. Substituting Eqs. (2)

and (5) in Eq. (7), followed by rearrangement, one obtains

$$\text{Abs}(t) = 2L\varepsilon_B C_{A0} + LC_{A0}(\varepsilon_A - 2\varepsilon_B)e^{-kt}. \quad (8)$$

Analysis of variance was carried out on experimental data obtained from hydrolysis of ovine and caprine milk casein by all enzymes, using Excel 2000 (Microsoft Office 2000 Professional, C.P.C., Portugal). Student's t -test, with $P < 0.05$, was employed for comparison of means.

Results and discussion

Clotting activity

The variation of the clotting time with the enzyme concentration is shown in Fig. 1. All forms of enzyme were added to reconstituted milk at final concentrations in the range 0.00025–0.00500 g/ml, thus giving rise to milk clotting times in the ranges 303–4384, 378–3615 and 230–4187 s, respectively, for cardosin A, cardosin B and a 2:1 mixture of both enzymes. The dependence of the clotting time on the enzyme concentration was estimated by fitting the model depicted in Eq. (1) to the data. The goodness of the fit is apparent in Fig. 1, and was further confirmed by the data depicted in Table 1.

Our experimental data were also fitted to by the kinetic models proposed for milk coagulation by van Hooydonk and Walstra (1987), Kopelman and Cogan (1976) and Hyslop et al. (1979), which correlate the clotting time with the reciprocal enzyme concentration, the reciprocal clotting time with the enzyme concentration, and the

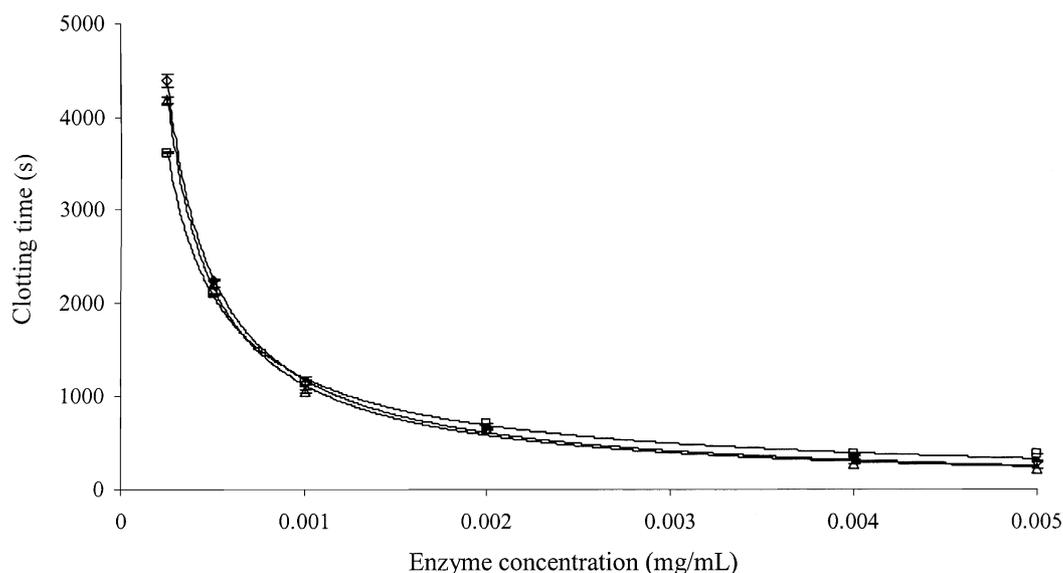


Fig. 1. Variation of clotting time with initial enzyme concentration, for cardosin A (◇), cardosin B (△), and a mixture of both enzymes at the ratio 2:1 (□); experimental data (◇, △, □), with error bar indicating Standard Deviation and mathematical fit (Eq. (1)).

Table 1

Best form of the model proposed by Payens et al. (1977) pertaining to clotting time (t) as a function of initial enzyme concentration (E_0), and associated correlation coefficient

Enzyme	Best model	r^2
Cardosin A	$t = 2.527E_0^{-0.8942}$	0.999
Cardosin B	$t = 5.837E_0^{-0.7724}$	0.997
Cardosin A + Cardosin B (ratio 2:1)	$t = 1.422E_0^{-0.9652}$	0.996

clotting time with the reciprocal square root of enzyme concentration, respectively (see Table 2).

The data encompassing milk coagulation by cardosin A were well fitted to by all proposed models, although yielding a not so good correlation coefficient in the case of the model by Hyslop et al. (1979) (see Table 2). Cardosin A is believed to be similar, in terms of specificity, to chymosin (Picon, Gaya, Medina, & Núñez, 1995); Verissimo, Esteves, Faro, and Pires (1995) reported that the best fit for chymosin was the one relating clotting time with reciprocal enzyme concentration.

Cardosin B is believed to resemble pepsin in terms of kinetic behaviour (Verissimo et al., 1995), and is known to possess a non-specific proteolytic activity; a slight deviation from the model proposed by van Hooydonk and Walstra was expected (Picon et al., 1995), and our results confirmed that to some degree. The other two models provided worse fits to the experimental data.

As far as the mixture of the two enzymes is concerned, Ketting and Pulay (1970) claimed that no linearly weighted relationships between clotting time and enzyme concentration are expected relative to the relationships pertaining to each enzyme in pure form; furthermore, no kinetic models pertaining to milk coagulation by mixtures of enzymes have, to our knowledge, been proposed to date. The milk coagulation by the mixture of the two aspartic proteases was fitted equally well by the models proposed by van Hooydonk and Walstra (1987) or by Kopelman and Cogan (1976); the poorer regression obtained with the model of Hyslop et al. (1979) was consistent with the not

so good regressions obtained when attempting to fit the pure enzyme performance in the first place. Cardosin A is always the most abundant within the cardosin inventory in crude aqueous extracts of the flowers of *C. cardunculus* (Frazão et al., 1999), so it is expected that the catalytic behaviour of the whole mixture is somehow dominated by that exhibited by cardosin A.

Proteolytic activity

One of the most expeditious methods for assessing proteolysis in cheese is quantification of the peptides soluble in TCA, at levels between 2% and 12% (w/v); the higher the TCA concentration, the shorter the sizes of the soluble peptides (Yvon, Chabanet, & Pélissier, 1989).

The hydrolysis of ovine and caprine caseins brought about by cardosin A, cardosin B and a mixture of both, as shown by the production of 5% (w/v) TCA-soluble peptides, is depicted in Fig. 2, over a period of 10 h of incubation under conditions prevailing during regular milk coagulation or cheese ripening conditions. The rate of hydrolysis (measured by the slope of the tangent at each point) decreases appreciably with time; the product concentration tends toward an asymptotic value by ca. 6 h; the limiting situation is always higher when the substrate is caprine casein, and when incubation takes place under cheese ripening conditions.

The result of the fit of Eq. (8) to our experimental data, using non-linear regression analysis, yields estimates for the adjustable parameters ε_A , ε_B and k , which are tabulated in Tables 3–5; the best fits are plotted in Fig. 2. The goodness of the fits is obvious from inspection of Fig. 2, and is further confirmed by the magnitude of the correlation coefficients (typically very close to unity), coupled with the significance of all parameters and the fact that the variance explained by the model is high. Note that parameters ε_A , ε_B and k should be regarded solely as adjustable parameters, so their individual physical meanings (as implied in derivation of Eq. (8)) should not be considered in a strict sense – the same

Table 2

Best form of the alternative models pertaining to clotting time (t) as a function of initial enzyme concentration (E_0), and associated correlation coefficient

Enzyme	Best model	r^2	Reference
Cardosin A	$t = 1.074E_0^{-1} + 89.51$	0.999	van Hooydonk and Walstra (1987)
	$t^{-1} = 0.6534E_0 + 0.0001556$	0.994	Kopelman and Cogan (1976)
	$t = 81.65E_0^{-1/2} - 1099$	0.96	Hyslop et al. (1979)
Cardosin B	$t = 1.161E_0^{-1} - 289.6$	0.996	van Hooydonk and Walstra (1987)
	$t^{-1} = 0.5191E_0 + 0.0002701$	0.976	Kopelman and Cogan (1976)
	$t = 65.88E_0^{-1/2} - 716.9$	0.984	Hyslop et al. (1979)
Cardosin A + Cardosin B (ratio 2:1)	$t = 0.9606E_0^{-1} - 54.73$	0.999	van Hooydonk and Walstra (1987)
	$t^{-1} = 0.8719E_0 - 0.000005700$	0.996	Kopelman and Cogan (1976)
	$t = 79.30E_0^{-1/2} - 1100$	0.973	Hyslop et al. (1979)

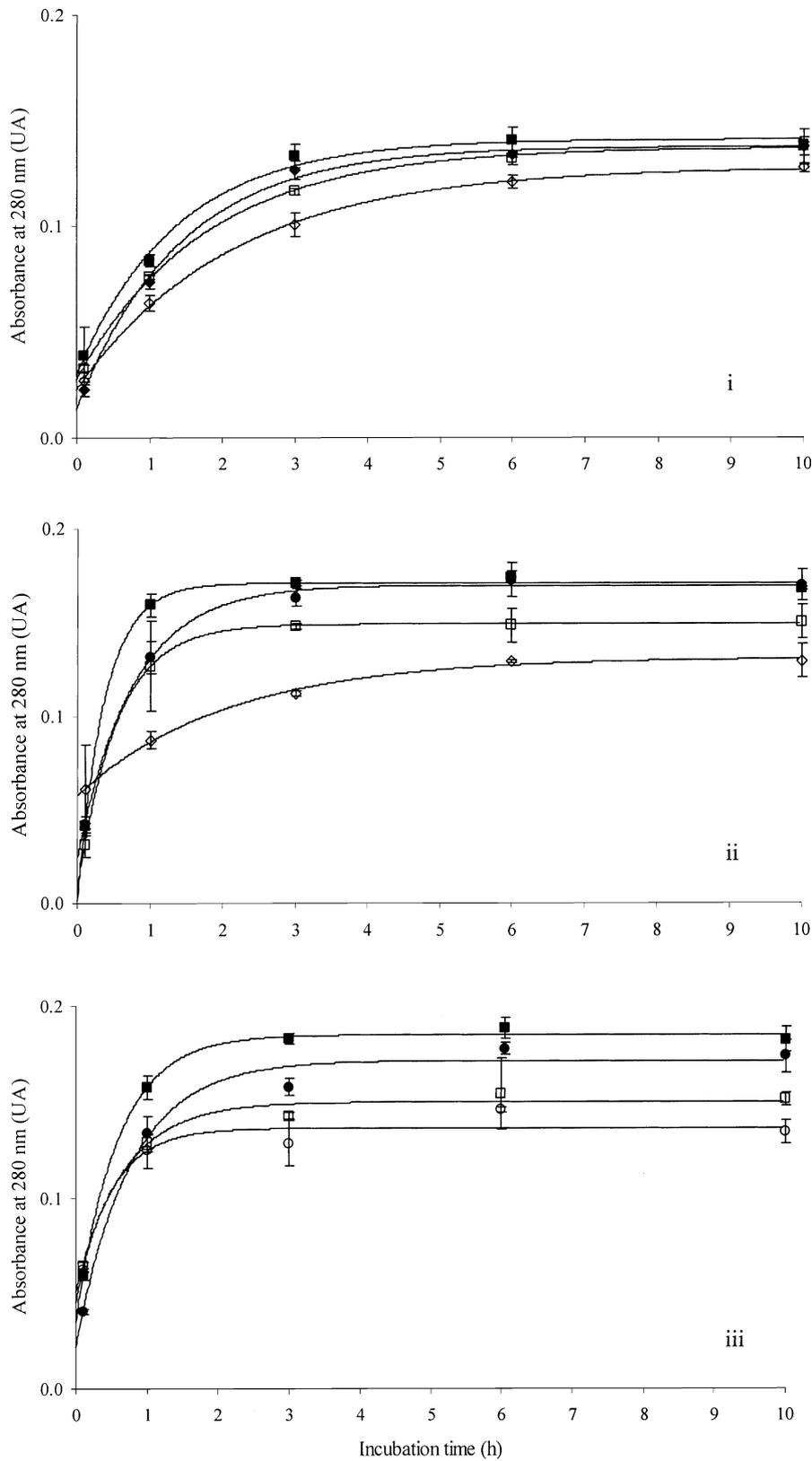


Fig. 2. Variation of absorbance (which correlates with proteolytic activity) with incubation time of ovine (○, ●) and caprine (□, ■) caseins, acted upon by cardosin A (i), cardosin B (ii), and a mixture of both enzymes at the ratio 2:1 (iii), under regular milk coagulation (○, □) and under cheese ripening (●, ■) conditions; experimental data (○, ●, □, ■), with error bar indicating standard deviation and mathematical fit (Eq. (8)).

Table 3

Parameter estimates of the mathematical model proposed (Eq. (5)) that describes proteolysis of ovine and caprine caseins by cardosin A, and statistical assessment of the fit

Parameters	Estimates \pm SD	Covariance matrix			Estimates \pm SD	Covariance matrix		
		Clotting conditions				Ripening conditions		
		ε_A	ε_B	k		ε_A	ε_B	k
<i>Ovine</i>								
ε_A (cm ⁻¹ mol ⁻¹)	1.365 \pm 0.324	0.1051	0.005148	-0.009143	2.981 \pm 0.635	0.4027	0.01569	-0.03753
ε_B (cm ⁻¹ mol ⁻¹)	6.890 \pm 0.104		0.01087	-0.003247	7.061 \pm 0.199		0.03959	-0.01282
k (s ⁻¹)	0.7006 \pm 0.0529			0.002796	0.7263 \pm 0.115			0.01315
		$R^2 = 0.994$				$R^2 = 0.973$		
<i>Caprine</i>								
ε_A (cm ⁻¹ mol ⁻¹)	2.301 \pm 0.239	0.05709	0.006511	-0.004874	2.750 \pm 0.159	0.02542	0.002152	-0.002369
ε_B (cm ⁻¹ mol ⁻¹)	6.394 \pm 0.108		0.01157	-0.002902	6.8566 \pm 0.0603		0.003638	-0.001085
k (s ⁻¹)	0.4627 \pm 0.0366			0.001336	0.5643 \pm 0.0265			0.0007010
		$R^2 = 0.995$				$R^2 = 0.998$		

Table 4

Parameter estimates of the mathematical model proposed (Eq. (5)) that describes the proteolysis of ovine and caprine caseins by cardosin B, and statistical assessment of the fit

Parameters	Estimates \pm SD	Covariance matrix			Estimates \pm SD	Covariance matrix		
		Clotting conditions				Ripening conditions		
		ε_A	ε_B	k		ε_A	ε_B	k
<i>Ovine</i>								
ε_A (cm ⁻¹ mol ⁻¹)	2.437 \pm 0.636	0.4049	0.01330	-0.05727	0.1321 \pm 0.551	0.3033	0.01171	-0.1165
ε_B (cm ⁻¹ mol ⁻¹)	8.498 \pm 0.148		0.002178	-0.01148	8.555 \pm 0.0708		0.005010	-0.008687
k (s ⁻¹)	1.304 \pm 0.159			0.02524	2.662 \pm 0.260			0.06781
		$R^2 = 0.987$				$R^2 = 0.997$		
<i>Caprine</i>								
ε_A (cm ⁻¹ mol ⁻¹)	5.744 \pm 0.765	0.5859	0.05694	-0.07003	0.7266 \pm 1.15	1.320	0.04284	-0.3002
ε_B (cm ⁻¹ mol ⁻¹)	6.561 \pm 0.328		0.1075	-0.03913	7.461 \pm 0.223		0.04979	-0.04333
k (s ⁻¹)	0.4845 \pm 0.167			0.02802	1.830 \pm 0.422			0.1778
		$R^2 = 0.908$				$R^2 = 0.964$		

Table 5

Parameter estimates of the mathematical model proposed (Eq. (5)) that describes proteolysis of ovine and caprine caseins by a mixture of cardosin A and cardosin B (ratio 2:1), and statistical assessment of the fit

Parameters	Estimates \pm SD	Covariance matrix			Estimates \pm SD	Covariance matrix		
		Clotting conditions				Ripening conditions		
		ε_A	ε_B	k		ε_A	ε_B	k
<i>Ovine</i>								
ε_A (cm ⁻¹ mol ⁻¹)	2.214 \pm 0.764	0.5836	0.02328	-0.08832	3.503 \pm 0.486	0.2363	0.007370	-0.04520
ε_B (cm ⁻¹ mol ⁻¹)	8.577 \pm 0.176		0.03087	-0.01761	9.250 \pm 0.0987		0.009748	-0.007086
k (s ⁻¹)	1.312 \pm 0.195			0.03814	1.704 \pm 0.155			0.02412
		$R^2 = 0.983$				$R^2 = 0.994$		
<i>Caprine</i>								
ε_A (cm ⁻¹ mol ⁻¹)	4.526 \pm 0.999	0.9982	0.3906	-0.4534	5.080 \pm 0.966	0.9329	0.03337	-0.2418
ε_B (cm ⁻¹ mol ⁻¹)	6.826 \pm 0.177		0.03143	-0.05678	7.506 \pm 0.208		0.04340	-0.04290
k (s ⁻¹)	2.032 \pm 0.667			0.4446	1.504 \pm 0.420			0.1766
		$R^2 = 0.944$				$R^2 = 0.944$		

applies to the functional forms of models in Tables 1 and 2 regarding clotting time.

Incubation of cardosin A in citrate buffer (pH 5.0), for 20 h at 10 °C, had a significant effect on its activity upon ovine and caprine caseins ($P < 0.01$). For cardosin B, as well as for the mixture of both enzymes, incubation at cheese ripening conditions also led to a significant increase of its activity ($P < 0.01$). These results are not surprising at all, because the cardosins are aspartic acid proteases, which possess an optimum pH in the acidic range and close to 5.2 (Sousa & Malcata, 1996). Moreover, as previously indicated, cardosin B resembles pepsin in its activity and specificity (Veríssimo et al., 1995), so its activity is expected to vary favourably with incubation under ripening conditions (Andrén & von Reedt, 1990).

In conclusion, a simple equation that relates variation of clotting time with initial enzyme concentration appropriately describes milk coagulation by enzymes of plant origin, namely cardosins A and B, either isolated or as a mixture. This model can be employed to predict milk clotting time by those enzymes, and eventually to optimise the coagulation process.

Previous incubation of the enzymes, under typical cheese ripening conditions, leads to a significant increase of their proteolytic activity, which is always higher upon caprine rather than ovine milk caseins. A mathematical model can appropriately describe the proteolytic activity; the overall rate of hydrolysis decreases appreciably with time, eventually reaching an asymptotic behaviour.

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