# Tailoring the structure of casein micelles through a multifactorial approach to manipulate rennet coagulation properties

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21 Abstract:

The internal structure of the casein micelle remains a matter of debate. The two latest models published consider the presence of at least three levels of structure: the casein micelle; dense regions within the micellar structure, and calcium phosphate nanoclusters and/or proteins inhomogeneities. The properties of casein micelles are affected by modifications to the environment, such as variations in pH or the addition of salts: The scientific literature typically considers the effects of one factor at a time, while in industrial processes, several modifications are performed simultaneously.

The aim of this study was to assess the impact of multifactorial environmental modifications on the colloidal, structural and rennet coagulation properties of casein micelles. The experimental design involved variations in pH, together with the addition of NaCl and CaCl<sub>2</sub>.

A key finding was that dense regions (~ 20 nm in size) could be released from the casein 32 micelle. The addition of NaCl and CaCl<sub>2</sub> had opposing effects, *i.e.* enhancing or limiting this 33 34 micellar disruption, respectively. A decrease in pH had the strongest impact on the mineral 35 balance, causing the colloidal CaP to solubilize and the micelle to swell. The rennet clotting time was impacted by variations in pH and NaCl content. Interestingly, a consideration of all three 36 levels of casein micelle structure and their interactions was needed to explain variations in the 37 38 firmness of rennet gels. This study illustrates the complex interplay of factors affecting micellar 39 structure and improves our understanding of how micelles can be manipulated to control their 40 properties.

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<u>Keywords:</u> casein micelle, rennet properties, internal structure, multifactorial modifications, small
 angle neutron scattering, cryo transmission electron microscopy

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### 45 **1** Introduction

In milk, casein proteins ( $\alpha_{s1}$ ,  $\alpha_{s2}$ ,  $\beta$  and  $\kappa$ ) and minerals (mainly calcium phosphate (CaP) 46 47 nanoclusters) self-assemble to form a colloid referred to as the casein micelle. Electrostatic, 48 hydrophobic and Van der Waals forces hold the different components together leading to a polydisperse population of particles 100 – 200 nm in diameter (Dalgleish & Corredig, 2012; Holt, 49 50 Carver, Ecroyd, & Thorn, 2013; Holt, 2016; Holt & Horne, 1996; Horne, 2017; Thorn et al., 2005). There is a consensus that casein micelles are stabilized through electrostatic and steric 51 repulsions due to the presence of a polyelectrolyte brush of  $\kappa$ -casein at the micellar surface (de 52 53 Kruif, 1999; de Kruif & Zhulina, 1996; Tuinier & de Kruif, 2002). Despite extensive studies the internal structure of the casein micelle, *i.e.* the interactions between caseins and the minerals 54 55 located within the colloid structure, is still under debate. The recent use of small angle X-ray and 56 neutron scattering (SAXS and SANS, respectively) in parallel with cryo-transmission electron 57 microscopy (cryo-TEM) has enabled questions around the internal structure to be partially 58 answered, although without general agreement (Bouchoux et al., 2015; Bouchoux, Gésan-Guiziou, Pérez, & Cabane, 2010; Day, Raynes, Leis, Liu, & Williams, 2017; De Kruif, 2014; 59 Ingham et al., 2015, 2016; Marchin, Putaux, Pignon, & Léonil, 2007; Pignon et al., 2004; Shukla, 60 Narayanan, & Zanchi, 2009). To date, the previously most accepted submicelle model (Schmidt, 61 62 1982; Walstra, 1999) has been progressively abandoned in favor of a more open structure 63 composed of dense regions, water channels and CaP nanoclusters. Environmental factors such as variations in pH, temperature and the addition of salts or 64 65 chelating agents (de Kort, Minor, Snoeren, van Hooijdonk, & van der Linden, 2011; Gaucheron, 2004; Lazzaro et al., 2017; Silva et al., 2013) induce shifts in the mineral balance between the 66 67 diffusible and colloidal phases of the casein micelle. These mineral modifications also involve

68 colloidal modifications that lead to changes in the functional properties of the casein micelle,

- 69 such as the formation and stability of emulsions, thermal stability, and response to acid and
- rennet coagulation (Broyard & Gaucheron, 2015; Gaucheron, 2004). Although the mineral

fraction only represents a small proportion of the milk components (0.7%), it is used to control
the properties of numerous manufactured dairy products.

This study focuses on the gelation of the casein micelle following the addition of chymosin (rennet), which forms the first step of cheese manufacture. The rennet coagulation mechanism can be divided into three steps: firstly, chymosin hydrolyzes the  $\kappa$ -casein of the casein micelle by cleaving the Phe(105)-Met(106) bond. Next, once a sufficient amount of  $\kappa$ -casein is hydrolyzed the depleted casein micelles aggregate (Lucey, 2002) and finally, this aggregation is followed a reorganization and reticulation of the casein gel (Dalgleish & Corredig, 2012).

79 Rennet coagulation can be assessed by two main characteristics: (i) the rennet clotting time

80 (RCT), the time elapsed from the addition of rennet or chymosin to the detectable onset of

gelation and (ii) the firmness of the gel. The RCT mainly depends on the rate of the enzymatic

reaction and the aggregation of the paracaseinates (first and second steps), while the firmness
depends on the organization and the strength of the gel (third step).

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85 Variations in pH and the addition of NaCl and CaCl<sub>2</sub> are steps commonly applied in cheese manufacture to control the coagulation of milk. Many studies have been published describing the 86 separate influence of each of these parameters on the colloidal properties and on the rennet 87 88 coagulation properties of the casein micelle e.g.: (Bulca, Wolfschoon-Pombo, & Kulozik, 2016; 89 Choi, Horne, & Lucey, 2007; Daviau, Famelart, Pierre, Goudedranche, & Maubois, 2000; Deeth & Lewis, 2015; Famelart, Le Graet, & Raulot, 1999; Famelart, Lepesant, Gaucheron, Le Graet, & 90 91 Schuck, 1996; Grufferty & Fox, 1985; Karlsson, Ipsen, & Ardö, 2007; Karlsson et al., 2007; Sandra, Ho, Alexander, & Corredig, 2012; Sbodio, Tercero, Coutaz, & Revelli, 2006; Zhao & 92 93 Corredig, 2015; Zoon, van Vliet, & Walstra, 1988, 1989). Although necessary to understand the 94 dissociated effect of each parameter, this monofactorial approach does not correspond to the reality of the cheese industry where a multifactorial approach occurs with the simultaneous 95 variation of several parameters. 96

The aim of the current study was to investigate the effect of simultaneous modification of the 97 physico-chemical parameters on both the colloidal properties and the rennet induced 98 99 coagulation casein micelles. An experimental matrix of 27 different suspensions of casein 100 micelles in water, with variations in added NaCl and CaCl<sub>2</sub>, at three different pH was designed. A combination of multiple advanced biophysical techniques, such as cryo-TEM and SAXS, was 101 102 used to make a thorough characterization of the casein micelles in terms of physicochemical, structural and renneting properties. Appropriate statistical analyses were applied to establish the 103 104 relationships between the colloidal and the rennet coagulation properties of the modified casein micelles, leading to the first study that combines SAXS characterization with an assessment of 105 106 the functional properties of casein micelles.

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### 108 2 Materials and methods

### 109 **2.1 Chemicals**

All chemicals used for this study, hydrochloric acid (VWR chemicals, Fontenay-sous-bois,

111 France), NaCl (PanReac AppliChem, Barcelona, Spain), CaCl<sub>2</sub> (VWR International, Leuven,

Belgium) and sodium azide (Riedek-de Haën, Seelze, Germany) were of analytical grade.

113 2.2 Materials

114 Experiments were carried out using a native phosphocaseinate (NPC) powder resuspended in deionised water at 24.2± 0.8 g kg<sup>-1</sup> of protein. Concentrated NPC was supplied by Gillot SAS 115 (Saint Hilaire de Briouze, France) and obtained by microfiltration (0.1 µm pore size membrane) 116 117 of raw skimmed milk followed by diafiltration against milli-Q water. The concentrate was then spray dried according to the method described by Pierre, Fauquant, Le Graet, & Maubois (1992) 118 119 and Schuck et al., (1994) using Bionov facilities (Rennes, France). Casein and their associated minerals represented more than 90% of the total solid content of the powder. Residual whey 120 proteins (3%) (w/w) and traces of lactose were present in the powder. 121

For the coagulation experiments, Chr. Hansen (Hoersholm, Denmark) supplied commercial
 chymosin (CHY-MAX M 200, 200 IMCU ml<sup>-1</sup>).

## 124 **2.3 Preparation of casein micelle suspensions**

125 An experimental matrix was designed to assess the concomitant effects of variations in pH and the addition of NaCl or CaCl<sub>2</sub>. The range of pH and the final concentrations of added NaCl and 126 CaCl<sub>2</sub> were selected to produce suspensions which would form gels within one hour after the 127 addition of a set amount of chymosin. The pH values targeted were 5.7, 6.3 or 6.9 and the final 128 concentrations of NaCl and CaCl<sub>2</sub> in the suspensions were 0, 50 and 100 mmol kg<sup>-1</sup> or 0, 7.5 129 and 15 mmol kg<sup>-1</sup>, respectively. A full experimental design was carried out, where 27 different 130 131 suspensions of casein micelles were prepared in water, with different salt and pH environments. 132 The suspensions were named from A to Z, and CTRL to represent the control which consisted of NPC in water at pH 6.9, with no added salts (Fig.1). 133

134 Dispersion of the NPC powder in milli-Q water was performed as described in Lazzaro et al.,

135 (2017). Varying amounts of stock solutions of NaCl (2.5 mol kg<sup>-1</sup> in milli-Q water, pH 6.9) and

136 CaCl<sub>2</sub> (0.25 mol kg<sup>-1</sup> in milli-Q water, pH 6.9) were added to the concentrated suspensions of

137 NPC in milli-Q water, in such a way that after the dilution step (see below), the required

138 concentrations in salts were reached (Fig.1). The suspensions were stirred for 30 min. The pH

shift induced by the addition of salts was corrected using HCI 1M in milli-Q water and set to 5.7,

140 6.3 or 6.9 (Fig.1). The suspensions were then diluted to  $24.2 \pm 0.8$  g kg<sup>-1</sup> of protein and left

141 overnight at room temperature. If necessary, the pH was readjusted prior to experiments. For

142 convenience, Tables 1, 2, 3 and Figures 6, 7, 8 report the results of different analyses on a set

143 on 9 samples only. These selected samples correspond to the corners of the cubic experimental

design (suspensions A, B, D, E, J, L, M, CTRL ; extreme points) and the center point of the

145 experimental design (suspension T) (Fig. 1).

## 146 **2.4 Recovery of the diffusible phases of the suspensions**

147 Aliquots (15mL) of each NPC suspension were ultrafiltered by centrifugation for 30 min, at 20°C

and 1800 g. using Vivaspin concentrators (molecular weight cut-off 10 kD, Vivascience,

149 Palaiseau, France). The diffusible phases were analyzed to determine the concentrations of

diffusible ions and used for the dilution of the suspension for the turbidity (T) and nanoparticle

151 tracking analysis (NTA) measurements and for the background determination for the SAXS

- 152 measurements.
- 153
- 154 **2.5 Analysis**

# 155 2.5.1 Protein content

- 156 The total nitrogen content of each suspension was determined according to the Kjeldahl method
- 157 (IDF standard 20-1, 2014). A factor of 6.38 was used to convert nitrogen to protein
- 158 concentration. Measurements were performed in duplicate.

# 159 2.5.2 Mineral composition and distribution

160 Total and diffusible cations (calcium Ca, sodium Na) and anions (chloride Cl, inorganic

161 phosphate Pi) contents were determined as described in (Lazzaro et al., 2017). Colloidal

- 162 concentrations were deduced by subtracting the concentration of diffusible ions from the
- 163 concentration of total ions. The mineral concentrations were adjusted to account for the small
- 164 differences in protein content (see section 2.5.1.)

# 165 2.5.3 Turbidity measurements

166 Absorbance measurements were carried out at 600 nm and 20 °C using a UV-visible

167 spectrometer (UVmc<sup>2</sup>, Safas, Monaco). The casein micelle suspensions were diluted 10 times in

- their diffusible phases and analyzed immediately. The diffusible phase for each suspensions
- 169 was also analyzed. Absorbance measurements were converted into turbidity (T) according to the
- 170 following formula:

171  $\tau = 2.303 * \frac{OD(600nm)}{l}$ 

with OD(600nm) being the optical density of the suspension (difference between the absorbance of the diluted suspension and the absorbance of its diffusible phase); and *l* the light path length (l = 1 cm).

175 **2.5.4 Nanoparticle tracking analysis** 

176 Nanoparticle Tracking Analysis (NTA) was performed at 20°C using a Nanosight NS300 177 (Malvern Instruments, Malvern, United Kingdom) equipped with a Nanosight syringe pump. The principle of NTA is based on the tracking of individual particles in suspension. A large dilution 178 179 (40000 times) was necessary to meet the optimal settings of the apparatus, *i.e.* 20 - 100180 particles per frame during the measurement combined to a dark background image. A syringe was loaded with the diluted suspension, the focus was adjusted manually, the infusion rate was 181 182 set to 20, the camera level to 12 and 5 video images of 60 s recorded. The video images of the 183 movement of particles under Brownian motion were analyzed by the NTA image analysis 184 software (V 3.0 0064., Malvern Instruments). The minimum screen gain, and detection threshold of 3 were selected to maximize the detection of small particles (< 50 nm diameter). The particle 185 186 size distributions obtained (data not shown) were fitted with a log-normal population of particles using Schulz equation (Schulz, 1935): 187

$$W(R, r_{NTA}, \sigma) = \frac{R^Z}{\Gamma(Z+1)} \left(\frac{Z+1}{r_{NTA}}\right)^{Z+1} \times exp\left[-\frac{R}{r_{NTA}}(Z+1)\right]$$

188 Where  $r_{NTA}$  is the average radius of the particles and Z relates to the polydispersity ( $\sigma$ ) of the 189 particle radii (R) given by the expression:

$$\sigma = \left(\frac{\overline{R^2}}{r_{NTA}} - 1\right) = \frac{1}{Z+1}$$

190 The value of  $\sigma$  varied from 0.23 to 0.47 within the set of suspensions and  $r_{NTA}$  was further used 191 in the statistical analyses.

# 192 2.5.5 Cryogenic Transmission Electron microscopy

193 A thin vitrified film of casein micelle suspension was prepared as described in the method of 194 (Chen et al., 2011). A Formvar lacey carbon film mounted on a 300 mesh copper grids 195 (ProSciTech, Queensland, Australia) were glow discharged for 15 s and used as a hydrophilic 196 support on which the suspensions (4 µL) were adsorbed. After 30 s, the grids were plunged in liquid ethane using a Vitrobot (FEI Company, Eindhoven, Netherlands) to freeze the sample. 197 198 The grids were observed on a Technai G2 TF30 (FEI company, Eindhoven, Netherlands) 199 operating at 200 kV and equipped with a Gatan US1000 2kX2k CCD Camera (Gatan). Between 200 10 and 20 micrographs per suspension were recorded under low-dose conditions with defocus 201 values of 4 - 6 µm. Image analysis was performed using ImageJ software (National Institute of 202 Health, USA). Due to the large numbers of samples, this process was automated through the 203 use of macros set up within Image J. In total, between 1554 particles within suspension TEM 204 and 29 092 particles within suspension B were measured in three steps:

205 i) The region of interest (ROI) was defined:

Particle detection was strictly limited to the area free from the grid structure and ice particlespresent as a result of sample preparation.

208 ii) Particle detection:

209 To detect particles, the background was subtracted from images using a 'rolling ball' algorithm 210 and smoothed using Gaussian filtering before the threshold was applied and particles measured. 211 Touching particles were separated using the distance transform watershed plugin (Quasi-212 Euclidean). Most of the very large particles (diameter > 50 nm) were often partially hidden by the 213 grid. Therefore, the grid-obstructed particles were excluded from the detection only if more than 214 10% of the area defined by the best-fitted ellipse drawn around the particle was hidden under 215 the grid. In order to prevent any misrepresentative segmentation, which can be caused from 216 automatic detection, all segmentation results were visualized and the overlay was saved as a 217 separate image that was manually inspected.

218 iii) Shape measurement of the particles:

Feret's diameter was determined for each particle detected. Particles were defined as small (< 50 nm in Feret's diameter) or large (> 50 nm in Feret's diameter). The ratio of small to large particles was then calculated for each suspension and defined as  $\Gamma_{s/l}$ .

222 **2.5.6** Small angle X ray scattering measurements, data treatment and modelling

223 The SAXS measurements were carried out on the suspensions and their respective diffusible

224 phases at the SAXS/WAXS beamline of the Australian Synchrotron (Clayton, Melbourne,

Australia). The beamline is equipped with a Pilatus 1 M detector (170 mm x 170 mm, effective

pixel size of 172 x 172 μm). Each sample was measured at sample-to-detector lengths of 7.106

m and 0.721 m with respective photon energies of 8.2 keV (1.512 Å) or 18.1 keV (0.685 Å).

228 Merging the data from both distances provided a q range of 1.3 10<sup>-3</sup> to 1.93 Å <sup>-1</sup>. The samples

were drawn into a 1.5 mm glass capillary; allowing continuous flow through the X-ray beam

230 during measurements. The data were obtained from at least 10 exposures of 2 s intervals at

231 20°C. The capillary was rinsed with water, followed by 8 M guanidine and again with water

before being air-dried between each sample analysis.

The SAXS intensities were normalized to an absolute scale and at least 10 measurements per 233 sample were averaged to obtain the intensity profiles using ScatterBrain (V 2.71) (Australian 234 Synchrotron, Clayton, Australia). Measurements of the diffusible phases for each suspension 235 236 were subtracted from the corresponding scattering intensities of the suspensions to account for 237 background. Finally, the intensities were adjusted to account for the difference in total protein content of the suspensions in Primus (V 3.2) (ATSAS, Hamburg, Germany). Igor software (V 238 239 7.0.2.2) (Wave Metrics, Lake Oswego, USA) was used to merge the data from the 7.106 and 0.721 m detector distances to obtain the final curves I = f(q). 240

The SAXS scattering intensity curves were fitted according to the model of Bouchoux et al.,

242 (2010) with slight modifications. This model considers three populations of particles: Population

- A, observed at low q (up to  $6 \times 10^{-3} \text{ Å}^{-1}$ ) where the scattering intensity corresponds to the
- presence of casein micelles; population B, in the intermediate q regions (6 x  $10^{-3}$  to 2 x  $10^{-2}$  Å<sup>-1</sup>),

where the scattering corresponds to dense regions inside the casein micelles and population C; at high q (7-8 x  $10^{-2}$  Å<sup>-1</sup>) attributable to the CaP nanoclusters. The intensity depends on the form factor of each population P<sub>n</sub>(q) (approximated by the form factor of polydisperse spheres of mean radius r<sub>a</sub>, r<sub>b</sub> and r<sub>c</sub>) and on prefactors a, b and c:

$$I(q) = a P_a(q) + b P_b(q) + c P_c(q)$$

249 with:

$$a = \alpha \times n_a (Vma \times \Delta \rho_a)^2$$
$$b = \alpha \times n_b (Vmb \times \Delta \rho_b)^2$$
$$c = \alpha \times n_c (Vmc \times \Delta \rho_c)^2$$

where  $n_a$ ,  $n_b$ ,  $n_c$  were the number of scatterers for each population, Vma, Vmb, Vmc their

volume and  $\Delta \rho_a$ ,  $\Delta \rho_b$ ,  $\Delta \rho_c$  their contrast, respectively.

The absolute number of casein micelles was assumed to be the same in all the suspensions.

253 This hypothesis is based on the observations of Moitzi, Menzel, Schurtenberger, & Stradner,

254 (2011) that a decrease in pH left the number of casein micelles unmodified, even if some casein micelle materials are subdivided into individual monomers or smaller casein aggregates with a 255 256 resulting decrease in micelle size and mass. In the present study, it was also assumed that the 257 modifications of the pH and the addition of NaCl and CaCl<sub>2</sub> applied in the experimental design 258 were not sufficient to cause the complete disruption of the micellar structure. This implies that the n<sub>a</sub> value is not affected by the physical-chemical modifications of our samples. We chose to 259 set it to 1 so that n<sub>b</sub> and n<sub>c</sub> are defined relatively to one casein micelle. As a consequence, the 260 constant  $\alpha$  accounts for both the electron scattering length and the absolute number density of 261 262 casein micelles in the samples.

This model was tested on our data according to the procedure of Bouchoux et al., (2010). First, the value of the radius of each population,  $r_a$ ,  $r_b$  and  $r_c$ , and the prefactors, a, b, c, were determined by fitting the model to the experimental data with polydispersities set to  $\sigma_a = 1/3$ ,  $\sigma_b$ = 1/3 and  $\sigma_c = 0.2$ . In a second step, the value of the constant  $\alpha$  was calculated from the control sample (CTRL) using the prefactor a obtained from the fit, the size (and therefore volume) of the micelle in this case, and the contrast  $\Delta \rho_a$  of a native casein micelle in water, i.e. 0.018 e- Å<sup>-3</sup> ( $\rho_{water} = 0.334 \text{ e-} Å^{-3}$  and  $\rho_{caseinmicelle} = 0.352 \text{ e-} Å^{-3}$ ).

270 In a third step, the values of  $\Delta \rho_a$ ,  $n_b$  and  $n_c$  were calculated for the 27 suspensions using the sizes and prefactors obtained from the fits. The micelle,  $\Delta \rho_a$  was simply calculated from 271 272 prefactors a and sizes  $r_a$ , knowing that  $n_a = 1$ . The number of dense regions  $n_b$  was calculated from prefactors b, sizes  $r_b$ , and making the assumption that contrast  $\Delta \rho_b$  is relatively insensitive 273 to the physical-chemical modifications performed in this study.  $\Delta \rho_{b}$  is taken as 0.035 e- Å<sup>-3</sup>, i.e. 274 twice the contrast of the micelle assuming that dense regions occupy 50% of the total volume of 275 276 the casein micelle (Bouchoux et al. (2010)). The CaP nanoclusters / Protein inhomogeneities, n<sub>c</sub> was calculated from prefactors c, sizes  $r_c,$  and an estimated contrast  $\Delta \rho_c$  of 0.172 e-  ${\rm \AA}^{-3}$  that 277 was also assumed to be constant between samples. Note that in a recent work, Ingham et al., 278 (2016) suggest a new interpretation and assign the high g features of SAXS data to the 279 280 presence of inhomogeneous protein structures of 1-3 nm length scale instead of CaP 281 nanoclusters. As our purpose is not to take a position on this question, we decided not to restrict our analysis solely to the interpretation of Bouchoux et al. (2010). A number of possible 282 inhomogeneities n<sub>cPI</sub> was therefore calculated following Ingham's postulate, this time using an 283 estimated contrast of 0.126 e-  $Å^{-3}$  (Ingham et al., 2016). 284

285 2.5.7 Rennet coagulation properties

The coagulation properties of samples were assessed using a ChymoGRAPH® (Chr Hansen, Denmark) which uses a similar physical principal to the Formagraph (McMahon & Brown, 1982). The RCT corresponded to the time elapsed from chymosin addition to the detectable onset of gelation, where gelation was defined as at the time point when the firmness of the suspensions was > 0. The maximal firmness recorded during the 60 min duration of the experiment was defined as the firmness of the gel.

292 2.5.8 Statistical treatments

293 As mentioned in section 2.3., a complete experimental design was carried out to study the combined effects of variation in pH (5.7, 6.3, 6.9), NaCl addition (0, 50, 100 mmol kg<sup>-1</sup>) and 294 295 CaCl<sub>2</sub> addition (0, 7.5, 15 mmol kg<sup>-1</sup>) on the colloidal and renneting properties of the casein 296 micelles (Fig. 1). The data set was subjected to statistical analysis by principal component analysis (PCA) using the Facto-MineR package and the R software (Lê, Josse, & Husson, 2008) 297 298 to highlight the correlations between the different measurements. All the correlations mentioned in the results and discussion section were found to be significant (p < 0.05) using the paired 299 300 student t-test.

In addition, multiple linear regression was also applied to the SAXS variables  $r_a$ ,  $n_b$  and  $n_{cCaP}$ , using the software STATGRAPHICS Centurion XVII (V. 17.1.10, Statpoint Technologies, The Plains, USA) in order to evaluate the effect of these structural features on the firmness of the gel. A model of firmness was defined that included the quadratic effects of  $r_a$ ,  $n_b$  and  $n_{cCaP}$  and the second order interactions between these factors. The full equation of the model was: *Firmness* = *constant* +  $r_a$  +  $n_b$  +  $n_c$  +  $r_a^2$  +  $n_b^2$  +  $n_c^2$  + ( $r_a \times n_b$ ) + ( $r_a \times n_c$ ) + ( $n_b \times n_c$ ) The LS-means were calculated and differences regarded as significant for p < 0.05. Non-

307 significant effects were excluded from the model, except when first order effects were

308 participating in interaction effects.

309

# 310 3 Results and discussion

The results of the biophysical analyses are presented and discussed in two main sections. In the first section, the PCA analyses are used to (i) assess the impacts of variation in pH and the addition of NaCl and CaCl<sub>2</sub> on the mineral balance of the casein micelle, (ii) establish relationships between the structure of the casein micelle and its other colloidal properties. In the second section, the relationships between rennet coagulation properties and colloidal and structural features of casein micelle are considered. Here, the PCA analyses are able to explain

the variation observed in RCT, while linear regression is able to explain interactions between the
 structural properties and the firmness of the rennet gels.

### **319 3.1 Colloidal and structural properties of the modified casein micelles**

320 It was found that almost 80% of the variability observed within our experimental design could be 321 evaluated by considering the first four principle components of the PCA analysis (Figs. 2 and 4). 322

# 323 3.1.1 Impact of the environmental modifications on the mineral balance of the casein 324 micelle

Analysis of the partition of minerals Ca, Pi, Na and Cl between the colloidal and diffusible

326 phases of the suspensions (Table 1) revealed that the colloidal ions consisted mainly in Ca and

327 Pi, whereas Na and Cl were mainly present in the diffusible phases of the suspensions when

NaCl was added. Figure 2 shows the strong, positive correlations between the pH and colloidal

329 Ca and Pi (0.85 and 0.80, respectively) and a negative correlation with the concentration of

diffusible Pi (- 0.86). The diffusible Ca concentration was weakly, but still significantly, impacted

by variation in pH, with a correlation coefficient of - 0.5 (Fig. 3).

The strongest variation of Ca in the diffusible phase was attributed to the addition of CaCl<sub>2</sub>, with

a correlation coefficient of 0.84 (Fig. 4). Colloidal and diffusible concentrations of Pi were also

significantly impacted by the addition of CaCl<sub>2</sub> but to a much smaller extent, with correlation

coefficients of 0.42 and - 0.42 respectively (Fig. 3). These results confirm that modifications in

 $^{336}$  pH and CaCl<sub>2</sub> induced opposite effects on the mineral content of the casein micelle, with pH

inducing a stronger effect compared to CaCl<sub>2</sub> addition.

A decrease in pH led to the solubilization of the CaP nanoclusters, which has been reported in

the literature (Dalgleish & Law, 1989; Daviau, Famelart, Pierre, Goudrdranche, & Maubois,

2000; Famelart, Lepesant, Gaucheron, Le Graet, & Schuck, 1996; Le Graet & Brulé, 1993; Le

- Graet & Gaucheron, 1999; Le Ray et al., 1998; van Hooydonk, Boerrigter, & Hagedoorn, 1986;
- Zoon, van Vliet, & Walstra, 1989). Conversely, the addition of CaCl<sub>2</sub> limited CaP solubilization

presumably by shifting the Ca<sup>2+</sup> equilibrium through the saturation of the diffusible phase (Moitzi 343 et al., 2011). Added Ca would also directly associate with caseins and/or with the diffusible Pi 344 345 and precipitate as CaP (Le Ray et al., 1998; Philippe, Le Graet, & Gaucheron, 2005; Philippe, 346 Gaucheron, Le Graet, Michel, & Garem, 2003; Udabage, McKinnon, & Augustin, 2000). The addition of NaCl positively correlated with the diffusible Na and Cl concentrations (0.99 and 347 348 0.95, respectively) (Fig. 2), which was consistent with the presence of NaCl in the diffusible phase. No significant correlation was found between NaCl addition and the concentrations of 349 350 colloidal ions (Fig. 2) i.e. NaCl had no direct effect on the mineral content of the casein micelle within the range studied (0 to 100 mmol  $kg^{-1}$ ). This result is in agreement with the finding of 351 352 Karlsson, Ipsen, & Ardö (2007) who reported no change in the colloidal CaP content. However, 353 this observation differs from the results of (Aoki, Umeda, & Nakao, 1999; Famelart et al., 1996; Grufferty & Fox, 1985; Zhao & Corredig, 2015; Zoon et al., 1989) who reported that solubilization 354 of Ca, and occasionally Pi, occurred when NaCl was added to fresh or reconstituted skim milk or 355 356 casein micelle suspensions. These discrepancies could arise from differences in pH, as in most 357 cases, variations in pH induced by NaCl addition were not corrected, and/or the amount of added NaCl was 3 to 5 times higher than in the present study. 358

## 359 **3.1.2 Consequences on the structural properties of the casein micelle**

360 As mentioned in the section 2.5.6 of this paper, the SAXS data presented in Fig. 5 were treated 361 primarily using the sponge-like model defined by Bouchoux, Gésan-Guiziou, Pérez, & Cabane, (2010) which uses three populations to interpret the SAXS pattern of casein micelles. The 362 alternate interpretation of Ingham et al., (2016) also defines three population but attributes 363 population C; at high g (7-8 x  $10^{-2}$  Å<sup>-1</sup>), to protein inhomogeneties rather than the CaP 364 365 nanoclusters. Given that we have decided not to rely exclusively on the interpretation of either Ingham et al., (2016) or Bouchoux et al., (2010), the implications of the both of these studies are 366 discussed here. The SAXS patterns (Fig. 5) show variability in the intensity of signal across 367 these 3 different regions for the set of 9 selected samples. The structural features determined 368

from the SAXS data ( $r_a$ ,  $\Delta p_a$ ,  $r_b$ ,  $n_b$ ,  $r_c$ ,  $n_c$ ) for the 27 suspensions were compared to the other physico-chemical variables (concentrations of colloidal and diffusible minerals,  $\tau$ ,  $r_{nano}$ ,  $\Gamma_{s/l...}$ ) for each population of scatterers (A, B and C) and analysed by PCA (Figs. 2, 3, 4). These correlations and the relevance of the two SAXS models applied are discussed in the following sections.

### 374 **3.1.2.1 Population A: the casein micelle**

375 The modelling of the SAXS data enabled 2 variables,  $r_a$  and  $\Delta \rho_a$ , describing the casein micelle radius and its contrast to be defined, respectively. The radius, ra varied from 41.5 to 58.1 nm 376 (Table 2), which is consistent with values determined in earlier characterizations of milk or 377 378 casein micelle dispersions by SAXS (Bouchoux et al., 2010; Ingham et al., 2016; Pignon et al., 379 2004; Shukla et al., 2009). r<sub>a</sub> positively and strongly correlated with the mean micellar radius measured by NTA,  $r_{NTA}$  (53.8 <  $r_{NTA}$  < 74.8 nm) and the turbidity,  $\tau$  (12.6 <  $\tau$  < 43.8) of the 380 different suspensions (Fig. 2 and Table 2), with correlation coefficients of 0.72 and 0.70, 381 382 respectively. Major differences were found when comparing these radii to the values found by Tran Le, Saveyn, Hoa, & Van der Meeren (2008) for NPC dispersed in water (diameter of 212 383 nm). This discrepancy could arise from strong differences in the conditions of analysis (e.g. 384 higher dilution of x 40 000 for our suspensions compared to x 6 000 for Tran et al. (2008), or 385 386 different threshold settings).

387 The PCA also suggests a dependency between the micellar size and mineral balance. Indeed, r<sub>a</sub> showed negative correlations with colloidal concentrations of Ca and Pi (- 0.71 and - 0.38, 388 respectively) and a positive correlation with the concentration of diffusible Pi (0.38) (Fig. 2). The 389 solubilization of the micellar CaP caused by a decrease in pH resulted in an increase in micellar 390 391 size due to swelling. This size increase also affected the turbidity of the suspensions that 392 increased (correlation coefficient of - 0.46 between pH and T) (Fig. 2). These results were in 393 agreement with those of Daviau et al., (2000) and van Hooydonk et al., (1986) but differed to the 394 observations of Moitzi et al., (2011) and Ouanezar, Guyomarc'h, & Bouchoux, (2012). Indeed,

395 these last authors reported a decrease of micellar diameter, measured by multiple angle 3D light 396 scattering or by AFM microscopy, respectively. In these studies, skim milk and casein micelle 397 powders were resuspended in deionised water or synthetic milk ultrafiltrate (lactose free saline 398 solution), respectively, providing a different ionic environment for the casein micelles than in our study. In addition, the pH ranges covered in these studies were lower than in our study and this 399 400 could result in different behavior when the colloid is exposed to more severe conditions. 401  $CaCl_2$  addition had no impact on  $r_a$  (Fig. 3), which was in agreement with results reported by Philippe et al., (2005); Philippe et al., (2003) and Udabage et al., (2000). Conversely, the 402 403 concentrations of NaCl and diffusible Na significantly correlated with r<sub>a</sub> (- 0.46, - 0.43, 404 respectively) (Fig. 2), with NaCl addition slightly reducing the diameter of the casein micelle. This 405 differs from the findings of Zhao & Corredig, (2015) and Karlsson, Ipsen, Schrader, & Ardö, 406 (2005), where an increase in the size of casein micelles was observed. It is possible, however, that in these previous studies, the increase in size was due to the decrease in pH (not corrected) 407 408 induced by NaCl addition rather than the direct effect of the soluble NaCl salt. Here, T negatively 409 correlated with NaCl (- 0.46) (Fig. 2), reflecting a decrease in scattering following NaCl addition. This result was in accordance with the two studies cited above and could be caused by a 410 decrease in micellar size and/or internal rearrangements of the micelle structure. Diffusible Na 411 412 may screen the negative charge on  $\kappa$ -casein CMP, causing a partial collapse of the hairy layer 413 and a slight decrease in micellar size. The impact of NaCl on the size and turbidity of the casein micelles could also be related to the release of small casein aggregates (dense regions) from 414 415 case in micelles. This argument is developed further in section 3.1.2.2 of the present paper.  $\Delta \rho_a$ , defined as the contrast of the casein micelle, corresponds to the electron density of the micelle 416 417  $(\rho_{CM})$  relative to the electron density of the diffusible phase  $(\rho_{DF})$ , which consists of water containing ions (Ca, Na, Cl and Pi) from the NPC powder and/or the addition of NaCl and CaCl<sub>2</sub>. 418 The contribution of these diffusible ions to  $\rho_{DF}$  ranged from 0.02 to 0.73 %, and thus, was not 419 420 taken into account here and  $\rho_{DF}$  was considered constant and equal to the electron density of

water, (0.334 e<sup>-</sup>Å<sup>-3</sup>). Therefore, in the present study,  $\Delta \rho_a$  directly reflected the variation of the 421 electron density of the casein micelle. This value depended on the volume, the casein 422 423 concentration and the CaP content of the case micelles.  $\Delta \rho_a$  varied between 0.010 and 0.018 e<sup>-</sup>Å<sup>-3</sup> (Table 2), which is of same order of magnitude as the contrast of a native casein micelle 424 described by Bouchoux et al., (2010) and Ingham et al., (2016).  $\Delta \rho_a$  presented a negative 425 426 correlation (-0.86) with r<sub>a</sub> (Fig. 2). It also positively and strongly correlated with concentrations of colloidal Ca and Pi (0.90 and 0.67, respectively) (Fig. 2). These results were consistent with an 427 increase in the volume of the casein micelle due to a depletion in CaP, leading to a decrease in 428 the electron density of the micelle. 429

### 430 **3.1.2.2 Population B: the dense regions**

The scattering caused by population B is characterized by  $r_b$ , the radius, and  $n_b$ , the number of dense regions per casein micelle. Both features showed variability within the full sample set, with RSD's of 38.9 and 84.1 %, respectively (Table 2). PCA analysis (Fig 2.) indicates a strong and positive correlation (0.73) between  $n_b$  and  $\Gamma s/I$ , the ratio of small (< 50 nm in Feret's diameter) to large (> 50 nm in Feret's diameter) particles detected on cryo-TEM micrographs (section 2.5.5) (Fig. 6).

Large black and homogeneous strands crossing the cryo-TEM micrographs (Fig. 6) correspond 437 438 to the grids that support the suspensions and large circular spots (e.g. suspension E) or merged 439 spots (e.g. suspension D) were individual casein micelles and aggregates of casein micelles, respectively. Differences in the granularity of the background of the images were attributed to 440 441 the presence of small-dissociated parts of the casein micelle that were present in the diffusible phase. This feature was directly quantified by the ratio  $\Gamma_{sl}$ . Image analysis revealed that these 442 443 small particles have diameter of around 5 nm (around the resolution limit of the TEM microscope) to 50 nm (data not shown), which was is agreement with the size range of the 444 445 population B modelled by SAXS measurements (from 6.1 nm to 21.9 nm in radius – Table 2).

This observation suggests that the dense regions detected by SAXS were not only presentinside the micelle but also outside the casein micelle in the diffusible phase.

448 A positive correlation of 0.49 was also observed between n<sub>b</sub> and both NaCl and diffusible Na 449 (Fig. 2), indicating that NaCl addition increased n<sub>b</sub>. Conversely, enrichment of the suspensions with CaCl<sub>2</sub> weakly but significantly, reduced n<sub>b</sub> (correlation coefficient of - 0.39) (Fig. 7). Small 450 451 particles of around 20 nm in diameter were also observed by Müller-Buschbaum, Gebhardt, Roth, Metwalli, & Doster (2007) using atomic force microscopy under similar conditions. These 452 authors reported a decrease of the number of small particles in the presence of increasing Ca, 453 454 consistent with observations here. To date, only the present study and that of Müller-Buschbaum 455 et al., (2007) report the presence of dissociated aggregates outside of the casein micelle based 456 on observations by microscopy and SAXS. It is reasonable to assume, however, that such small 457 particles would not sediment by ultracentrifugation and a parallel can be established between our observations and the increasing presence of caseins in the supernatant obtained by 458 459 ultracentrifugation, defined as soluble caseins. Our observations are also consistent with 460 (Famelart et al., 1999; Zhao & Corredig, 2015), who reported an increase in the concentration of soluble casein after NaCl addition. Conversely, (Famelart et al., 1999; Philippe et al., 2005; 461 Udabage et al., 2000) observed a decrease in soluble caseins when CaCl<sub>2</sub> was added. This 462 463 may be explained by considering the actions of these two ions. NaCl would be responsible for 464 the disruption and loosening of the internal structure of the casein micelle by neutralizing negative charges on the casein chains, whereas CaCl<sub>2</sub> would either favor the creation of new 465 bonds between the phosphorylated caseins and/or prevent the dissociation of casein materials 466 by limiting the solubilization of CaP nanoclusters. Our finding of an increase in soluble casein 467 468 due to pH-induced dissociation of CaP nanoclusters is also consistent with reports by Dalgleish & Law, (1989); Le Graet & Gaucheron, (1999); and van Hooydonk et al., (1986). 469

470

In the present study, there was no significant correlation between pH and n<sub>b</sub> (Fig. 2). pH, There 471 was however, a direct and strong effect of pH on n<sub>cCaP</sub> (the number of population C scatterers 472 473 per micelle, correlation of 0.81), which correlates negatively with  $n_b$  (- 0.45) (Fig. 2). CaCl<sub>2</sub> 474 addition had no significant impact on the size of the dense regions (Fig. 7), while NaCl caused their decrease, as shown by the correlation between r<sub>b.</sub> and the concentration of NaCl and 475 476 therefore diffusible Na and Cl (- 0.50, - 0.53 and - 0.45, respectively) (Fig. 2). The decrease of 477 population C also led to a decrease in size of the dense regions (correlation coefficient of 0.47 between  $n_{cCaP}$  and  $r_b$ ) (Fig. 2). Finally, it is interesting to note that  $n_b$  and  $r_b$  were inversely 478 correlated (- 0.59) (Fig. 2), meaning that the more dense the regions within the micelle, the 479 480 smaller the size of these regions.

### 481 **3.1.2.3** Population C: CaP nanoclusters or protein inhomogeneities

482 The mean radius of population C ( $r_c$ ) ranged from 1.5 to 1.7 nm, for the set of suspensions studied with a RSD of 3.1 % (Table 2). This indicates that this population is of similar size in 483 484 each suspension, regardless of the physico-chemical modifications applied. However, the 485 number per casein micelle, n<sub>cCaP</sub>, varied from 75 to 244. Considering this population as CaP, the number is lower than the values of CaP nanoclusters in a native case in micelle of  $355 \pm 20$ 486 CaHPO<sub>4</sub>.2H<sub>2</sub>O unit reported by Holt, Timmins, Errington, & Leaver, (1998). In our experiments, 487 488 the suspension of NPC powder in water caused ~20 % of the colloidal CaP to dissolve 489 (calculated based on the colloidal and diffusible contents of the CTRL sample (Table 1) and this reduction in CaP could explain the discrepancy with the data of Holt and colleagues. The 490 491 number observed, however, is consistent with the findings of Bouchoux et al., (2010), who reported ~210 CaP nanoclusters per casein micelle. 492

If the population C scatterers were considered as protein inhomogeneities, their number per
casein micelle varied from 140 to 453, which is about 17 times lower than the value found by
Ingham et al., (2016) (Table 2). This difference may be due to the application of a simple sphere
form factor in this study compared to the combination of a Sorensen form factor and hard sphere

structure factor used by Ingham et al., (2016). Our approach was nevertheless sufficient to fit the 497 498 SAXS pattern in the high g-region (Fig. 5). The properties of the C-scatterers *i.e.* constant size 499 and varying number of C-scatterers per casein micelle indicated the disappearance of this 500 population adhered to an "all-or-nothing" rule, where population C either "dissolved" completely upon modification of the environment or remained intact within the casein micelle. 501 502 According to the PCA analysis, n<sub>cCaP</sub> correlated highly with: pH (0.80), concentrations of colloidal Ca and Pi (0.86 and 0.80, respectively) and with concentration of diffusible Pi (- 0.84) (Fig. 2), 503 504 indicating that the high g feature disappeared with the pH-induced dissolution of colloidal CaP. 505 Similar pH induced changes in SAXS patterns of casein micelles were reported by Ingham et al., 506 (2016) and Marchin et al., (2007). The disappearance of the high q feature was also observed 507 when colloidal CaP was removed from the casein micelle by the use of chelating agents (EDTA 508 or Na<sub>3</sub>Cit) (Day et al., 2017; Ingham et al., 2016; Marchin et al., 2007; Pitkowski, Nicolai, & 509 Durand, 2007). These correlations are therefore consistent with the assignment of this 510 population as CaP nanoclusters, as suggested by (Holt, de Kruif, Tuinier, & Timmins, 2003). 511 The correlations between  $n_{cCaP}$  noted above are not inconsistent, however, with the hypothesis of Ingham et al. (2016) i.e. that this population of particles corresponds to protein 512 513 inhomogeneities. In this case, protein inhomogeneities would be closely linked to micellar CaP. 514 Dissolution of the CaP from the casein micelle would also induce the disruption of the protein 515 inhomogeneities. Such an arrangement would be in agreement with the dual binding model of 516 (Horne, 1998), that considers CaP nanoclusters not only as crosslinking agents but also as 517 charge neutralizers between casein chains that allow proteins to form more hydrophobic 518 interactions. Further analysis in the form of cross comparisons between the evolution of the high g SAXS shoulder and the specific intensity variation at  $q = 0.035 \text{ Å}^{-1}$  observed either in SANS or 519 520 resonant X-ray scattering would bring interesting information about this CaP nanocluster / 521 protein inhomogeneity dependency. Finally,  $n_{cCaP}$  correlated negatively with  $n_b$  (- 0.45) and with  $\Gamma_{s/l}$  (-0.43), and correlated positively with  $r_b$  (0.47) (Fig. 2). These weak but significant 522

523 correlations suggest that casein micelles depleted in population C released more dense regions524 into the diffusible phase.

## 525 **3.2** Coagulation properties of the modified casein micelles

The RCT and the maximum firmness of the gel, defined here as firmness, were determined for the 27 suspensions using data obtained from the firmness curves (Fig. 8). These two parameters were linked to the other colloidal and structural variables through PCA and multiple linear regression analyses.

## 530 3.2.1 Rennet clotting time

The use of rennet made the 27 suspensions clot between 1.1 to 42.4 min (RSD of 113.5%) 531 (Table 3). This large variability was first ascribed to the variation in pH, as there was a significant 532 533 correlation between RCT and pH with a coefficient of 0.69 (Fig. 2). A consequence of the pH 534 decrease was the solubilization of the micellar Ca and Pi (section 3.1.1). Therefore RCT also positively correlated with the concentration of colloidal Ca and Pi, and with diffusible Pi with 535 536 coefficients of 0.49, 0.61 and - 0.61, respectively (Fig. 2). A reduction in RCT as a result of a decrease in pH has been well described in the literature (Choi et al., 2007; Daviau et al., 2000; 537 Karlsson et al., 2007; Zoon et al., 1989). This has been ascribed to the enhancement of enzyme 538 activity and a decrease in the electrostatic repulsion between paracaseinates at low pH that 539 540 favored aggregation.

A weaker but significant and positive correlation was also observed between diffusible Na and
RCT (0.39) (Fig. 2), meaning that an increased concentration of this ion in the diffusible phase
led to an increase in the RCT. Similar effects of added NaCl have been also reported by (Bulca
et al., 2016; Famelart et al., 1999; Grufferty & Fox, 1985; Karlsson et al., 2007; Sbodio et al.,
2006; Zhao & Corredig, 2015; Zoon et al., 1989) and were attributed to a decrease in the rate of
the rennet enzyme due to the screening of charges on κ-casein and the enzyme.

547

548 The negative correlations of RCT with  $\tau$  and  $r_a$  (- 0.47 and – 0.53, respectively) (Fig. 2) indicated 549 that large micelles clotted more quickly than small ones, a finding that was opposite to those made in the studies of Ekstrand, (1980) and Ford & Grandison, (1986). The increase of the 550 551 micellar size in the present study was a consequence of the pH decrease that caused the micelles to swell. The correlation between these two factors is probably a disguised effect of the 552 553 pH. In the two studies with opposing findings, the micelles were fractionated according to their 554 size by ultracentrifugation and did not undergo any physico-chemical treatment, potentially explaining these differences. 555

### 556 **3.2.2 Maximum firmness of the rennet gel**

Although the firmness was highly variable, with an RSD of 34 % (Table 3) within the set of 557 558 suspensions, this variable did not correlate directly with the other colloidal and structural characteristics of the casein micelle suspensions. Indeed, the firmness, concentrations of CaCl<sub>2</sub> 559 and concentrations of diffusible Ca were the only well-projected variables as defined by the 3rd 560 and 4<sup>th</sup> dimensions of the PCA analysis (Fig. 4). Vectors representing concentrations of CaCl<sub>2</sub> 561 562 and diffusible Ca were orthogonal to the vector for firmness, reflecting no correlations between these variables. However, PCA estimates the first order correlations between variables, and 563 does not take into account the interactions that might exist between the different features. 564 565 A second statistical approach, multiple linear regression, was therefore used to assess the effect 566 of possible interactions between variables on the firmness of the rennet gels. Structural SAXS features revealed to be excellent candidates for this complementary analysis for two reasons. 567 568 Firstly, these variables were unique and interesting descriptors reporting information at three different structural levels: 1) the casein micelle, 40 to 60 nm in radius and previously described 569 570 as population A; 2) the dense regions, 6 to 22 nm in radius and previously described as population B; 3) the CaP nanoclusters (or protein inhomogeneities), 1.5 to 1.7 nm in radius and 571 572 previously described as population C. Secondly, these variables correlated significantly with the 573 whole set of colloidal and mineral features determined by other techniques and constituted a

574 way to summarize the whole set of data. Therefore,  $r_a$ ,  $n_b$  and  $n_{cCaP}$  were subject to linear regression in order to define a predictive model of the firmness that considered the guadratic 575 576 effects and the second order interactions between these variables. For consistency, the values 577 of n<sub>cCaP</sub> used in this statistical analysis were determined considering population C as CaP nanoclusters. This would make no difference in the properties of the model, as there was a 578 579 proportional relationship linking  $n_{cCaP}$  for CaP nanoclusters and  $n_{cPl}$  for protein inhomogeneities. 580 Based on the experimental design, the following model equation was established to predict the 581 maximum firmness of the rennet gels made from the suspensions:

*Firmness* =  $187.3 - 3.5 \times ra - 1.2 \times n_b - 0.9 \times n_{cCaP} + 0.02 \times (r_a \times n_c)$ 

where  $r_a$  was the radius of the casein micelle,  $n_b$  and  $n_{cCaP}$  were the number of dense regions and CaP nanoclusters per casein micelle, respectively. This model explained 68.5% of the variability of the firmness. Statistical analysis also revealed that the interaction between  $r_a$  and  $n_c$ ( $r_a \ge n_c$ ) and the first order effect of  $n_b$  in this model were significant. These two contributions had a statistical weight of 35.3% and 27 % in the model, respectively.

Figure 9.A and B. displays each suspension in the first four dimensions of the PCA. The 587 588 suspensions are colored according to their firmness (orange for weak, red for medium, black for 589 strong). Figure 9.A represents the evolution of firmness within the set of samples, defined by PCs 3 and 4 with the arrow pointing in the direction of increasing firmness (the same direction as 590 seen in Figure 4). The negative coefficient assigned to n<sub>b</sub> in the firmness equation model 591 592 indicates that the release of dense regions from the casein micelles led to the formation of 593 weaker gels. This direct effect is well illustrated when reading Figure 9.A from the bottom right 594 corner (suspension G, U - poor in dense regions and stronger gels) to the upper left corner 595 (suspension B - rich in dense regions and a weaker gel).

As mentioned in section 3.1.2.2, the release of dense regions was favored by the addition of

597 NaCl and limited by the addition of  $CaCl_2$ . The influence of NaCl on the firmness has been

598 examined by several groups but conflicting results have been reported. Consistent with the

599 results reported here, Famelart et al., (1999) and Grufferty & Fox (1985) did not observe any 600 modification of the moduli or the curd tension of the rennet gels upon addition of NaCI. However, 601 Bulca et al., (2016) and Zhao & Corredig, (2015) reported a decrease in the firmness or stiffness 602 of the rennet gels. While Zoon et al., (1989) observed higher moduli for 8h aged gels supplemented in NaCl but lower moduli only 1h after the addition of rennet to the milk, which 603 604 corresponded to the experimental conditions of the present study. The negative effect of NaCl on the firmness of rennet gels is poorly explained in the literature. A competition between Na<sup>+</sup> 605 and Ca<sup>2+</sup> has been proposed, as well as the screening of casein charges and in some cases, the 606 607 solubilization of the micellar CaP (Grufferty & Fox, 1985; Zhao & Corredig, 2015). Based on the 608 significant correlations that link  $n_b$  and NaCl and the significant negative effect of  $n_b$  on the 609 firmness of the rennet gel, we would argue that the decrease in firmness observed with the 610 addition of NaCl is due to the release of dense regions from the casein micelles. This explanation is similar to that of Gaygadzhiev, Massel, Alexander, & Corredig, (2012), who found 611 612 that the addition of sodium caseinate to milk inhibited the aggregation of casein micelles. In this 613 case the soluble dense regions may adsorb on the surface of the paracaseinate formed after rennet addition, causing an increase in the steric repulsion between the rennet-altered particles 614 615 which would limit the aggregation phenomenon and the formation of a firm network. 616 In contrast, there is a general consensus that CaCl<sub>2</sub> addition increases gel firmness (Deeth & 617 Lewis, 2015; Sandra et al., 2012; Zoon et al., 1988). In this case, this improvement was 618 attributed to the ability of Ca to preserve the number of CaP bonds between the caseins within 619 the micelle but also within the casein gel network. It was demonstrated in section 3.1.2.2, that CaCl<sub>2</sub> addition limited the release of dense regions, which consequently had a positive impact 620 on the firmness of the gels through the decrease of  $n_b$ . 621

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The effect of the interaction between  $r_a$  and  $n_{cCaP}$  appears to be more subtle. The direct effect of r<sub>a</sub> could be seen in figure 9.B from the upper right corner (suspension M - small casein micelle)

to the bottom left corner (suspension A, D, O - large and swollen casein micelle). There were no 625 626 direct and simple consequences of r<sub>a</sub> on the gel firmness. This can be illustrated by the medium 627 size micelles (center of the graph, suspensions B, S, V, L for instance) that can either form 628 weak, medium or strong gels. This result was quite consistent with the observation of Dalgleish, Brinkhuis, & Payens, (1981) who found no dependence between the size of micelles and their 629 630 coagulation. Yet, several authors have reported that small casein micelles form stronger gels (Ford & Grandison, 1986; Logan et al., 2015; Niki, Kohyama, Sano, & Nishinari, 1994). However, 631 632 in these studies, the micelles were in their native state because they were either isolated from milk by ultracentrifugation or were in milk samples that were selected from cows who produced 633 634 small casein micelles. In contrast, the present study involved modifications to the environment 635 that affected the size of the casein micelles.

The direct effect of  $n_{cCaP}$  can be observed in figure 9.B from the upper left corner (suspension B – poor in population C) to the bottom right corner (suspensions L, Y – rich in population C) but no direct dependency of gel firmness on  $n_{cCaP}$  was observed. As mentioned in section 3.1.2.3, this population may be either CaP nanoclusters or protein inhomogeneities. In both cases, the presence of such interactions, whether they are mineral-protein or protein-protein interactions, would create more crosslinking points resulting in a stronger gel network.

642 The literature reports a quadratic relationships between pH and rennet gel firmness, *i.e.* there is 643 a parabolic increase in the gel firmness with pH up to a maximum value, followed by a parabolic decrease in firmness (Choi et al., 2007; Karlsson et al., 2007; Lucey, Johnson, & Horne, 2003; 644 645 Zoon et al., 1989). A change in pH modifies the ionization of individual amino acids, either increasing or decreasing the electrostatic interactions between casein chains. A simultaneous 646 647 consequence is the solubilization of the micellar CaP at lower pH, which decreases the attractive interactions between casein molecules. The addition of a chelating agent to milk or casein 648 649 suspension similarly leads to the solubilization of micellar CaP (de Kort et al., 2011; McCarthy et

al., 2017; Mizuno & Lucey, 2005; Pitkowski et al., 2007) and causes a decrease in the firmness
of the rennet gels (Choi et al., 2007).

652 At this point, it is important to remember that the variations of r<sub>a</sub> and n<sub>cCaP</sub> were not impacted by 653 only one factor (size fractionation, or pH, or chelating agent addition) but by the simultaneous effect of three factors (pH, NaCl and CaCl<sub>2</sub>). Therefore, the firmness modeling reveals that the 654 655 interaction between those variables should be considered. The interaction between  $r_a$  and  $n_{cCaP}$ on the rennet gel firmness means that the firmness at a given  $r_a$  depended on  $n_{cCaP}$  and vice 656 versa. As an example to illustrate this interaction, suspensions containing medium sized casein 657 micelles (from 45 to 47 nm in radius) can lead to the formation of weaker gels if the amount of C-658 659 particles is too low (suspension B). Yet these micelles formed medium strength gels 660 (suspensions P, E, C) if their C-particle content increased, or even stronger gels when those 661 casein micelles are rich in C-particles (suspensions X, I, T). Similarly, small casein micelles that were rich in C-particles formed weaker gels (suspension W, K, M) but an increase in micelle size 662 663 led to an increase in gel firmness (suspension H, V, T). Large casein micelles, depleted in C particles (suspension A, O, D) also formed weaker gels. 664

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### 666 **4** Conclusion

667 The multifactorial experimental design applied here allowed the effect of three variables: pH, 668 NaCl and CaCl<sub>2</sub> on the colloidal and rennet coagulation properties of casein micelles to be assessed and ranked. Variations in pH had the strongest influence on the mineral balance of the 669 670 casein micelles. A decrease in pH caused the colloidal CaP to solubilize. In contrast, NaCl addition had no impact on the mineral content of the casein micelle. The solubilization of 671 672 colloidal CaP caused the micelle to shrink while the addition of NaCl reduced the size of the casein micelle due to the release of small particles into the diffusible phase. The presence of 673 674 such particles, around 25 nm in diameter, was strongly supported by experimental SAXS data 675 combined with observations by cryo-TEM. These particles are believed to be part of the dense

regions described by Bouchoux et al. (2010); here they were observed both inside and outside of the casein micelle. CaCl<sub>2</sub> had no effect on the casein micelle size but prevented disruption of the dense regions within the micelle. The SAXS data also revealed the presence of a high-q structural feature (the C-population), that were of a constant size (~1.6 nm in radius) but varied in number with different environmental conditions. Their presence was strongly dependent to the CaP content of the casein micelle. This feature could be assigned to the presence of either CaP nanoclusters or protein inhomogeneities.

The renneting properties were most impacted by a decrease in pH, causing a reduction in RCT, while NaCl supplementation led to longer RCT. Variations in gel firmness were more complex but could be explained by considering the interactions between the size of the casein micelle,

the C-population and the dissociation of dense regions within the casein micelle.

Together the data presented here illustrate the complex interactions of three variables on the properties of casein micelles, this study provides a framework that links existing literature on the effect of single variables and improves our understanding of how the properties of casein micelles can be manipulated to control micelle size, structure and functional properties.

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**Table 1: Distribution of the mineral salts in the suspensions.** Colloidal concentrations were determined by subtracting the concentration of diffusible ions from the concentration of total ions. Average, standand deviation (SD), relative standard deviation (RSD), minimum and maximum values were determined on the complete set of 27 samples.

Table 2: Size-related parameters determined by different analytical methods including Turbidimetry, NTA, SAXS and Cryo-TEM. Average, standard deviation (SD), relative standard deviation (RSD), minimum and maximum values were determined on the complete set of 27 samples.  $n_{cCaP}$  and  $n_{cPl}$  correspond to the number of C scatterers per casein micelle, in the case where these scatterers are considered as CaP nanoclusters or protein Inhomogeneities, respectively.

**Table 3: Rennet coagulation properties of Firmness and Rennet Coagulation Time (RCT).** Average, standand deviation (SD), relative standard deviation (RSD), minimum and maximum values were determined on the complete set of 27 samples.

**Figure 1: Cubic representation of the experimental design.** Each of the 27 suspension is represented by a letter or designated as CTRL (control suspension, pH 6.9, no salts added). The pH of the suspensions was set to 5.7, 6.3 or 6.9 and the suspensions contained 0, 50 or 100 mmol kg<sup>-1</sup> of added NaCl and 0, 7.5 or 15 mmol kg<sup>-1</sup> of added CaCl<sub>2</sub>.

**Figure 2: PCA showing the correlation circle of 21 variables in the plan delimited by the two first principal components (PCs).** 57.8 % of the variability of the set of data is represented in this PCA plan.

**Figure 3: PCA showing the correlation circle of 21 variables in the plan delimited by the first and the fourth principal components (PCs).** 44.5 % of the variability of the set of data is represented in this PCA plan.

**Figure 4: PCA** showing the correlation circle of **21** variables in the plan delimited by the third and the fourth principal components (PCs). 21.3 % of the variability of the set of data is represented in this PCA plan.

**Figure 5: Fit to the SAXS data shown in a log – log plot.** The scattering curves have been translated along the y axis for clarity. The scattering profiles have been fitted using the model of Bouchoux et al. (2010) that contains 3 populations of scatterers: the casein micelle (population A - up to 6 x 10-3 Å<sup>-1</sup>), the dense regions (population B - 6 x 10-3 to 2 x 10-2 Å<sup>-1</sup>) and either the CaP nanocluster or the proteins inhomogeneities (population C - 7-8 x 10-2 Å<sup>-1</sup>). Open diamonds show experimental data; the solid lines represent fits for which the parameter values are given in Table 2.

**Figure 6: Cryo-TEM micrographs of the selected suspensions.** Large black strands crossing the images are carbon grids, circular or merged spots are casein micelles or aggregated casein micelles. Dark round spots in suspension L are ice particles that formed during storage of the grids in liquid nitrogen. Granular backgrounds point out the presence of small casein aggregates dissociated from the casein micelles. This feature is quantified for each suspensions by a  $\Gamma_{s/l}$  (ratio of small < 50 nm to large > 50 nm particles). The values of  $\Gamma_{s/l}$  are reported in Table 2.

**Figure 7: PCA showing the correlation circle of 21 variables in the plan delimited by the second and the fourth principal components (PCs).** 33.6 % of the variability of the set of data is represented in this PCA plan.

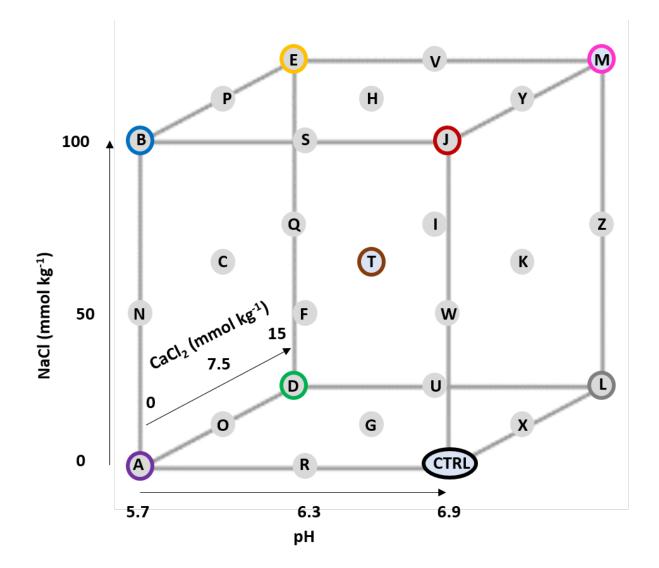
**Figure 8: Evolution of firmness as a function of time for selected samples.** Arrows point to the Rennet Clotting Time (RCT) for each of the suspensions, while firmess was defined as the maximum firmness reached within 60 min of the addition of chymosin to the suspensions. Values of RCT and Firmness are reported in Table 3.

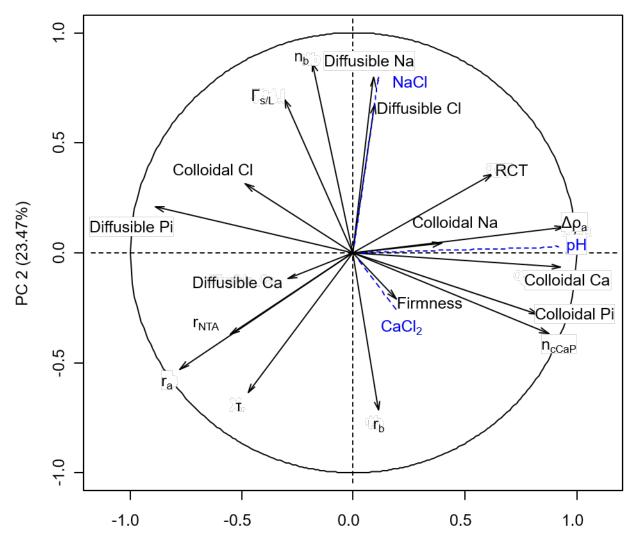
**Figure 9: PCA showing the similarity of suspensions determined by A.** The third and fourth Principal Components (PCs) and B. the two first PCs. Suspensions leading to weak gels are colored in orange, gels with medium firmness are red and gels with strong firmness are black. Arrows indicate the direction of correlation for firmness and SAXS variables in the same PC plans (A. see Figure 4 and B. see Figure 2).

	Diffusible Ca	Colloidal Ca	Diffusible Na	Colloidal Na	Diffusible Cl	Colloidal Cl	Diffusible Pi	Colloidal Pi
	(mmol kg <sup>-1</sup> )	(mmol kg <sup>-1</sup> )	(mmol kg <sup>-1</sup> )	(mmol kg <sup>-1</sup> )	(mmol kg <sup>-1</sup> )	(mmol kg-1)	(mmol kg <sup>-1</sup> )	(mmol kg <sup>-1</sup> )
	Full experimental plan - 27 suspensions							
Average	12.4	14.8	64.8	0.2	67.4	0.5	2.4	4.6
SD	6.0	4.3	45.5	0.5	45.7	1.4	1.3	1.3
RSD (%)	48.2	29.2	70.3	300.9	67.8	287.6	55.1	28.1
minimum	2.1	3.6	6.7	0.0	0.0	0.0	0.0	1.9
maximum	22.9	20.2	135.6	2.2	138.0	6.2	5.0	7.2
Selected individual suspensions								
А	9.6	8.2	8.9	0.0	2.2	1.6	3.8	3.3
В	11.0	8.8	114.4	0.0	102.7	6.2	4.8	1.9
D	20.1	12.0	24.0	0.0	26.9	3.3	2.9	3.5
Е	22.8	12.3	121.7	0.0	130.7	0.0	3.4	3.5
J	3.5	15.8	120.2	0.0	94.4	0.0	1.7	4.9
L	14.7	20.2	10.5	0.0	17.9	1.8	0.8	5.4
М	16.3	18.8	135.6	0.0	129.4	0.0	0.0	6.9
Т	11.5	15.8	65.4	0.0	69.2	0.0	2.3	5.0
CTRL	2.1	18.0	6.7	0.8	0.0	0.0	1.6	4.9

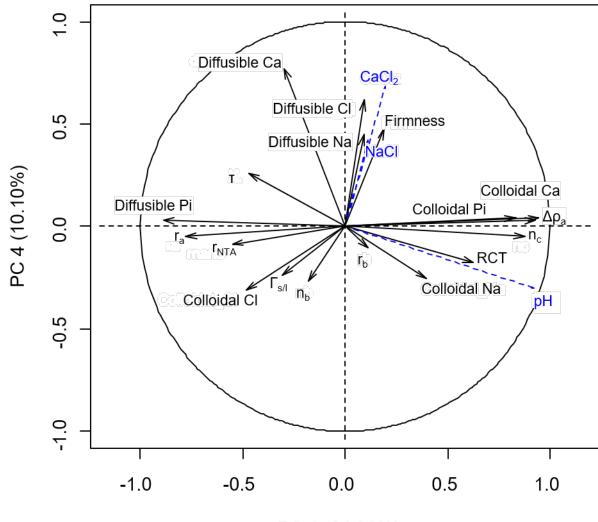
	Turbidimetry	NTA	SAXS					Cryo TEM		
	τ (cm <sup>-1</sup> )	r <sub>nta</sub> (nm)	Δρ <sub>a</sub> (e <sup>.</sup> .A <sup>.3</sup> )	r <sub>a</sub> (nm)	r <sub>b</sub> (nm)	r <sub>c</sub> (nm)	n <sub>b</sub>	n <sub>cCaP</sub>	n <sub>cPI</sub>	Γ <sub>s/I</sub>
			Full ex	perimental	design - 27	suspensior	ıs			
Average	21.2	61.0	0.015	45.6	10.7	1.6	3.3	171.5	318.9	8.5
SD	6.8	5.5	0.002	4.2	4.2	0.0	2.8	54.0	100.4	7.8
RSD (%)	32.1	9.0	13.5	9.3	38.9	3.1	84.1	31.5	31.5	91.0
minimum	12.6	53.8	0.010	41.5	6.1	1.5	0.2	75.3	140.0	0.9
maximum	43.8	74.8	0.018	58.1	21.9	1.7	13.3	243.7	453.3	35.6
			Se	elected indi	vidual susp	ensions				
А	21.4	70.8	0.011	54.7	7.6	1.7	2.1	87.0	161.9	6.7
В	14.6	53.8	0.012	45.5	6.9	1.6	13.3	75.3	140.0	35.6
D	43.8	73.4	0.012	54.5	6.1	1.6	1.6	127.4	236.9	3.0
E	18.0	59.6	0.014	46.4	7.6	1.6	4.0	121.6	226.2	6.2
J	14.0	59.3	0.016	41.8	8.7	1.6	6.6	174.2	324.0	16.8
L	13.8	59.8	0.018	43.5	12.8	1.6	1.2	231.7	431.0	6.7
М	12.6	61.8	0.017	42.3	8.3	1.6	5.1	202.6	376.9	1.7
Т	24.5	62.1	0.017	44.3	12.2	1.6	1.3	203.1	377.8	6.0
CTRL	15.7	57.0	0.018	41.5	10.1	1.5	4.0	228.0	424.1	4.6

	Firmness	RCT					
	(A.U.)	(min)					
full experimental design - 27 suspensions							
Average	14.9	8.6					
SD	5.1	9.8					
RSD (%)	34.0	113.5					
minimum	3.0	1.1					
maximum	20.9	42.4					
Selected individual suspensions							
А	3.3	1.4					
В	11.5	2.9					
D	14.0	1.4					
E	16.2	2.9					
J	8.0	24.6					
L	17.5	11.2					
М	3.0	42.4					
Т	20.9	3.4					
CTRL	13.9	11.0					

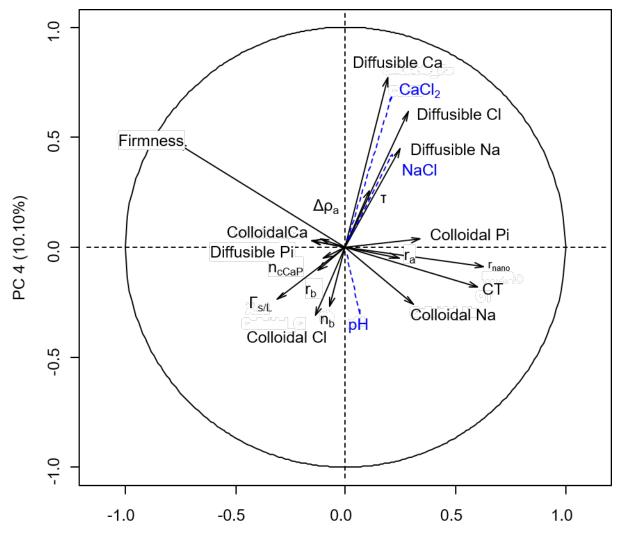




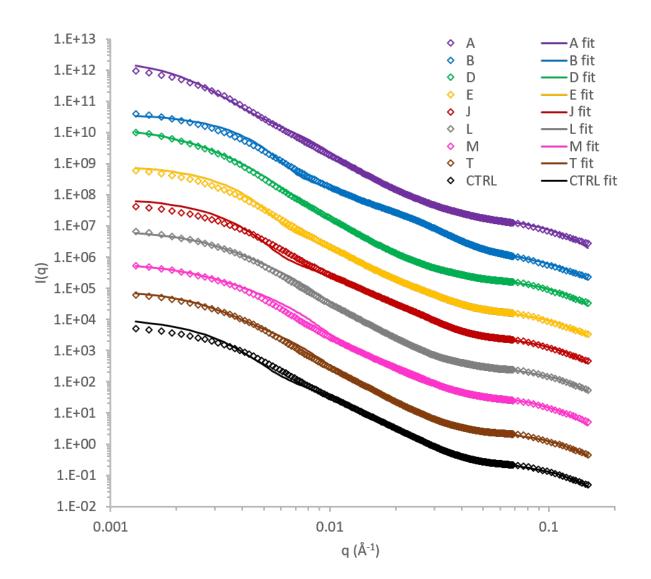
PC 1 (34.36%)

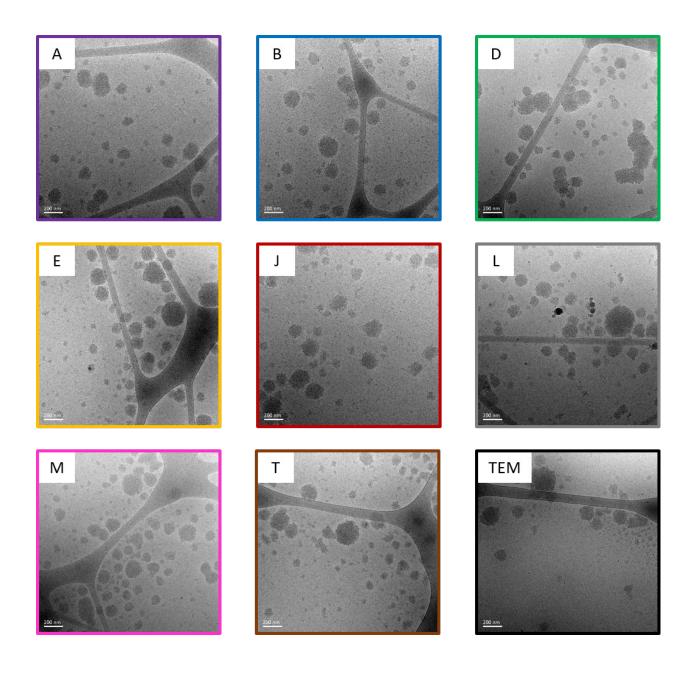


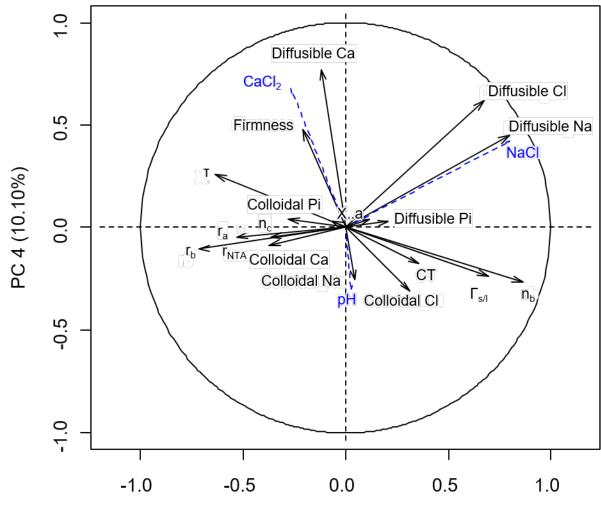
PC 1 (34.36%)



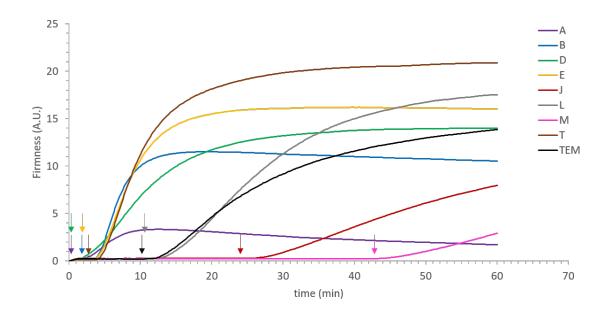
PC 3 (11.17%)

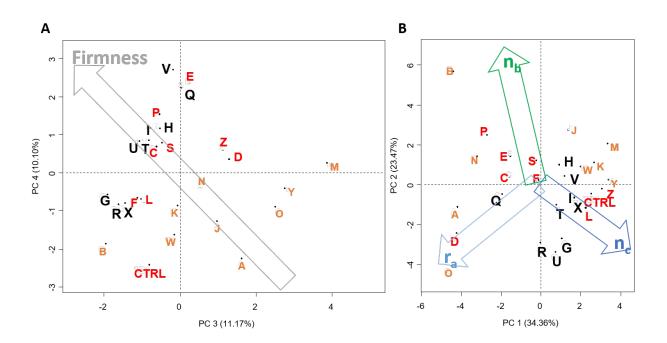






PC 2 (23.47%)





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