Effects of composite casein and β-lactoglobulin genotypes on renneting properties and composition of bovine milk by assuming an animal model

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The effects of κ-β-casein genotypes and β-lactoglobulin genotypes on the renneting properties and composition of milk were estimated for 174 and 155 milk samples of 59 Finnish Ayrshire and 55 Finnish Friesian cows, respectively. As well as the random additive genetic and permanent environmental effects of a cow, the model included the fixed effects for parity, lactation stage, season, κ-β-casein genotypes and β-lactoglobulin genotypes. Favourable renneting properties were associated with κ-β-casein genotypes ABA, A, and AAA, A, in the Finnish Ayrshire, and with ABA, B, AAA, A, AAA, A, ABA, A, and ABA, A, in the Finnish Friesian. The favourable effect of these genotypes on curd firming time and on firmness of the curd was partly due to their association with a high κ-casein concentration in the milk. The effect of the κ-casein E allele on renneting properties was unfavourable compared with that of the κ-casein B allele, and possibly with that of the A allele. The β-lactoglobulin genotypes had no effect on renneting properties but they had a clear effect on the protein composition of milk. The β-lactoglobulin AA genotype was associated with a high whey protein % and β-lactoglobulin concentration and the BB genotype with a high casein % and casein number.

Key words: coagulation properties, milk protein polymorphism

Introduction

Several studies have discussed the environmental and genetic factors that influence milk renneting properties. A number of workers have reported the favourable effect of the κ-casein B allele on renneting properties (e.g., Schaar 1984, Aaltonen and Antila 1987, Pagnacco and Caroli 1987). There is therefore interest in using κ-casein genotypes as a selection criterion when breeding for more favourable renneting properties and, thereby, better use of milk in cheese production. Conflicting results have been obtained for the effects of genotypes of αs1- and β-caseins and β-lactoglobulin on renneting properties. The origin, size and structure of the data, milk protein genotype frequencies, statistical
methods and models used or linkage disequilibrium between the casein loci may explain some of the discrepancies.

The effects of milk protein genotypes on renneting properties have been estimated by a least squares method (e.g., Tervalan et al. 1985, Pagnacco and Caroli 1987, Davoli et al. 1990, Machboeuf et al. 1993). When used for estimating single-gene effects on quantitative traits, this method ignores some or all of the polygenic effects (Kennedy et al. 1992). Because of probable confounding between single-gene effects and polygenic effects, it is thus possible to find an excess of spurious significant effects of the single genes. Kennedy et al. (1992) showed that the use of mixed model procedures under an animal model treating single-gene effects as fixed effects can provide unbiased estimates of single-gene effects and exact tests of associated hypotheses for pedigreed populations.

Various models have been applied for estimating milk protein genotype effects. A few studies have estimated the effect of genotypes of one protein at a time (Schaar 1984, Schaar et al. 1985, Davoli et al. 1990, Machboeuf et al. 1993). Others have included the genotypes of some or all major milk proteins in a model simultaneously (Feagan et al. 1972, Pagnacco and Caroli 1987, Tervalan et al. 1985, Olofss et al. 1992). Only a few authors have estimated the effects of composite genotypes of some or all major milk proteins (El-Negoumy et al. 1973, Pagnacco and Caroli 1987). Because the casein loci are tightly linked (e.g., Grosclaude et al. 1973, Threadgill and Womack 1990), the genotype effect of a casein locus may not be independent of the genotype effect of another locus. It seems therefore reasonable to estimate casein genotype effects simultaneously by using combined genotypes (Ojala et al. 1997).

In this study we estimated the effects of composite κ-β-casein genotypes and β-lactoglobulin genotypes on the renneting properties and composition of bovine milk by assuming an animal model. We also studied the associations of renneting properties with the composition of milk.

Material and methods

Milk samples

A total of 59 Finnish Ayrshire (FAy) cows from Helsinki University's experimental herd Viikki and 55 Finnish Friesian (FFr) cows from the experimental herd Suiita (FFr) were genotyped for α1s-, β- and κ-caseins and β-lactoglobulin by isoelectric focusing in polyacrylamide gels (Erhardt 1989). The FAy cows were born between 1980 and 1989, and the FFr cows between 1982 and 1989.

The effects of milk protein genotypes on the renneting properties and composition of milk were estimated by sampling the cows three times during lactation: 1, 3 and 5 months after calving. The cows calved from July 1990 to June 1991, and the sampling period lasted from the end of September 1990 to the end of October 1991. When the cows were housed indoors, the average proportion of concentrates in the feed, as determined on energy bases, was 37% for the FAy and 42% for the FFr. In 1991, the FAy cows were on pasture from the end of May to the end of September, and the FFr cows from mid-June to the end of September.

Because the cows were at different stages of lactation during sampling, the number of samples per cow varied from two to three, but for most cows it was three. The total number of milk samples was 174 for the FAy and 155 for the FFr. The milk samples (evening + morning milkings) were analysed for the following characteristics: daily milk yield, renneting properties, gross composition and protein composition.

Milk renneting properties

The renneting properties of individual milk samples (10 ml) determined by a Formagraph (Foss Electric, Hillerød, DK-3400, Denmark) at 32 °C for 30 min with 0.20 ml rennet (Renco) liquid diluted in 0.07 M sodium acetate buffer (1:100) were: renneting time (R), curd firming time (K25)
and firmness of the curd (E_{30}). R was the time from the addition of rennet to milk to the beginning of coagulation. K_{20} was the time from the beginning of coagulation to the moment the width of the curve was 20 mm. E_{30} was the width of the curve 30 min after the addition of rennet. Because milk samples were allowed to coagulate for only 30 min, renneting or curd firming times, or both, were not achieved for some samples owing to poor coagulation. Because the samples that did not coagulate in 30 min (nine samples from six FAy cows) were divided more or less equally among the milk protein genotypes, these samples were omitted from the statistical analyses of renneting properties.

**Composition of milk**

The fat and protein percentages were determined with a Milko-Scan 605 (Foss Electric) and the somatic cell count was made with a Fossomatic cell counter. Because the frequency distribution for the somatic cell count was far from normal in both breeds, the somatic cell counts were logarithmically transformed. pH was also measured.

The protein composition values determined were: casein and whey protein percentages, non-protein nitrogen (mg/g), casein number and the concentrations of \( \alpha_s \), \( \alpha_{\text{s'}}, \beta \)- and \( \kappa \)-caseins, \( \alpha \)-lactalbumin and \( \beta \)-lactoglobulin (g/l) in milk. The casein and whey protein percentages and non-protein nitrogen were determined according to International Dairy Federation (IDF) standards (1979 and 1986). Casein number was the proportion of casein in total protein. Concentrations of individual caseins in milk were obtained by multiplying proportions of individual caseins in total casein by casein content. The proportions of individual caseins were determined by fast protein liquid chromatography (FPLC) (Pharmacia Biotech, Uppsala, Sweden) as described by Syväoja (1992). Individual whey proteins were fractionated by FPLC gel filtration on a Superdex 75 HR 10/30 column (Pharmacia Biotech) as described by Syväoja and Korhonen (1994).

**Statistical analyses**

The effects of parity, lactation stage, season, \( \kappa \)-\( \beta \)-casein (\( \kappa \)-\( \beta \)-CN) genotypes and \( \beta \)-lactoglobulin (\( \beta \)-LG) genotypes on the renneting properties and composition of milk were estimated using an animal model. Owing to the difference in \( \kappa \)-\( \beta \)-CN genotypes formed in the FAy and FFr, the records from the two breeds, and thus from the two herds, were analysed separately. The following linear model was assumed:

**Model 1:**

\[
y_{ijklmn} = \mu + \text{parity}_i + \text{lstage}_j + \text{season}_k + \text{casge}_l + \text{lactge}_m + a_n + p_n + e_{ijklmn}
\]

where

- \( y_{ijklmn} \) = the \( o_{ih} \) observation of a milk renneting trait or a milk composition variable of the \( n_{ih} \) cow
- \( \mu \) = general mean
- \( \text{parity}_i \) = fixed effect of the \( i_{th} \) parity class
- \( \text{lstage}_j \) = fixed effect of the \( j_{th} \) lactation stage class
- \( \text{season}_k \) = fixed effect of the \( k_{th} \) season class
- \( \text{casge}_l \) = fixed effect of the \( l_{th} \) \( \kappa \)-\( \beta \)-CN genotype class
- \( \text{lactge}_m \) = fixed effect of the \( m_{th} \) \( \beta \)-LG genotype class
- \( a_n \) = random additive genetic effect of the \( n_{ih} \) cow (0, \( \mathbf{A} \sigma_a^2 \))
- \( p_n \) = random permanent environmental effect of the \( n_{ih} \) cow (0, \( \mathbf{I} \sigma_p^2 \))
- \( e_{ijklmn} \) = random residual effect, \( N(0, \mathbf{I} \sigma_e^2) \)

Parity was grouped into three classes: first, second and third to ninth lactation; lactation stage into three classes: 1, 3 and 5 months after calving; and season into four classes: Sep to Nov, Dec to Feb, Mar to May and Jun to Aug. The classification of \( \kappa \)-CN and \( \beta \)-LG genotypes is presented in Table 1. Because the FAy was monomorphic for \( \alpha_s \)-casein and there were a few cows with the \( \alpha_s \)-casein C allele in the FFr, \( \alpha_s \)-casein genotypes were not considered in the formation of composite genotypes.

The 59 FAy cows with records were daughters of 25 sires and the 55 FFr cows daughters of 32 sires. The number of daughters per sire ranged

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from one to six in the FAy and from one to four in the FFr. Ten FAy sires and 16 FFr sires had only one daughter each. The pedigrees of the cows with records were known for at least two generations, and the total number of animals in the statistical analyses was 352 for the FAy and 568 for the FFr.

In subsequent analyses, the associations of renneting properties with the composition variables of milk were estimated using Model 2, in which one milk composition variable at a time was included as a covariate in Model 1. Otherwise Model 2 worked like Model 1.

Variance components for the random effects (\( \sigma^2_g \), \( \sigma^2_e \) and \( \sigma^2_\varepsilon \)) in Models 1 and 2 were estimated from the data sets with the REML VCE package (Groeneveld 1993). The effects of parity, lactation stage, season, \( K\)-p-CN genotypes and \( \beta\)-LG genotypes on various characteristics were tested with the PEST program of Groeneveld (1990). The hypothesis tested was \( K'\varepsilon=0 \), in which \( K'\varepsilon \) contained the maximum number of independent estimable contrasts between classes of a fixed factor in the model. The statistical significance of regression coefficients (Model 2) was obtained by calculating \( F \) values using the difference between \( \sigma^2_\varepsilon \) from Model 1 without a covariate and \( \sigma^2_\varepsilon \) from Model 2 with a covariate. However, no consideration was made about the effect due to the number of independent tests generated by several traits within two different populations.

**Results**

**Frequencies of \( \kappa\)-\( \beta\)-CN and \( \beta\)-LG genotypes**

The expected number of all possible combinations of \( \kappa\)- and \( \beta\)-casein genotypes was 15 in the FAy and 18 in the FFr. Owing to the small size

| Finnish Ayrshire | Finnish Friesian |
|------------------|------------------|
| No. of cows      | Percentage of cows | No. of obs. | No. of cows | Percentage of cows | No. of obs. |
| \( \kappa\)-\( \beta\)-casein |
| AAA\(_1\)          | 2                | 3.0            | 6             | 10               | 18.0        | 29         |
| AAA\(_2\)          | 7                | 12.0           | 21            | 21               | 38.0        | 60         |
| AAA\(_3\)          | 14               | 24.0           | 40            | 2                | 4.0         | 4          |
| AAA\(_{123}\)      | 2                | 3.0            | 6             | 1                | 4.0         | 3          |
| ABA\(_1\)          | 8                | 14.0           | 24            | 8                | 15.0        | 23         |
| ABA\(_2\)          | 2                | 3.0            | 6             | 4                | 7.0         | 11         |
| ABA\(_B\)          | 2                | 4.0            | 5             | 2                | 7.0         | 11         |
| AEA\(_1\)          | 3                | 5.0            | 9             | 4                | 7.0         | 11         |
| AEA\(_2\)          | 13               | 22.0           | 38            | 3                | 5.0         | 9          |
| AEA\(_3\)          | 2                | 3.0            | 6             | 2                | 7.0         | 11         |
| AEA\(_A\)          | 2                | 3.0            | 6             | 4                | 7.0         | 11         |
| EEA\(_A\)          | 4                | 7.0            | 12            | 2                | 7.0         | 11         |
| \( \beta\)-LG |
| AA                 | 5                | 9.0            | 15            | 7                | 13.0        | 20         |
| AB                 | 28               | 47.0           | 82            | 34               | 62.0        | 97         |
| BB                 | 26               | 44.0           | 77            | 14               | 25.0        | 38         |

\( ^1\text{AAA}_{123} = \text{AAA}_{123} \) + \( \text{AAA}_{13} \) + \( \text{AAA}_{23} \)

\( \beta\)-lg = \( \beta\)-lactoglobulin
of the data sets and linkage disequilibrium in the casein loci, the observed number of combinations was 11 in the FAy and 9 in the FFr (Table 1). The most common $\kappa$-$\beta$-CN genotypes were $AAA_A^1A^2$ and $AEE_A^1A^2$ in the FAy, and $AAA_A^1A^2$ and $AEE_A^1A^2$ in the FFr. Consequently, 46% of the FAy cows and 56% of the FFr cows had one of the two most common $\kappa$-$\beta$-CN genotypes. The rarest $\kappa$-$\beta$-CN genotypes were carried by only one or two cows. The $\beta$-LG AB and BB genotypes were almost equally frequent in the FAy whereas AB was most frequent in the FFr. The $\beta$-LG AA genotype was rather rare in both breeds.

Means and Variation

Renneting properties

The average renneting and curd firming times were longer and the firmness of the curd was poorer for milk of the FAy than for milk of the FFr (Table 2). There was considerable variation in renneting properties in both breeds. The coefficients of variation for renneting and curd firming times would have been even larger had the poorly coagulating milk samples reached their extremely long renneting or curd firming times, or both.

Table 2. Milk renneting traits, daily milk yield, and gross and protein composition of milk.

|                  | Finnish Ayrshire | Finnish Friesian |
|------------------|------------------|------------------|
|                  | $\bar{x}$ | s.d. | cv | $\bar{x}$ | s.d. | cv |
| Milk renneting   |            |       |    |            |       |    |
| $R$, min         | 12.4      | 4.9   | 40 | 11.3      | 3.6   | 33 |
| $K_{20}^0$, min  | 8.1       | 3.3   | 41 | 7.4       | 3.5   | 47 |
| $E_{30}^0$, mm   | 25.8      | 12.5  | 48 | 31.2      | 10.1  | 33 |
| $E_{30}^0$, mm   | 27.2      | 11.2  | 41 |           |       |    |
| Milk yield and composition |       |       |    |            |       |    |
| Daily milk yield, kg | 26.2     | 5.7   | 21 | 26.0      | 6.0   | 23 |
| Fat %            | 4.51      | 0.66  | 15 | 4.27      | 0.69  | 16 |
| Protein %        | 3.20      | 0.29  | 9  | 3.14      | 0.25  | 8  |
| pH               | 6.76      | 0.07  | 1  | 6.77      | 0.08  | 1  |
| Somatic cell count (ln) | 4.59    | 1.63  | 36 | 5.67      | 1.35  | 24 |
| Protein composition |       |       |    |            |       |    |
| Casein %         | 2.49      | 0.25  | 10 | 2.44      | 0.22  | 9  |
| Whey protein %   | 0.53      | 0.08  | 15 | 0.54      | 0.07  | 13 |
| Non-protein nitrogen, mg/g | 0.29 | 0.06  | 21 | 0.26      | 0.03  | 12 |
| Casein number    | 78        | 2.49  | 3  | 78        | 2.41  | 3  |
| $\alpha_s$-casein, g/l | 9.47  | 0.98  | 10 | 9.32      | 0.89  | 10 |
| $\alpha_s$-casein, g/l | 3.19  | 0.67  | 21 | 3.97      | 0.45  | 15 |
| $\beta$-casein, g/l | 9.40  | 0.99  | 11 | 9.12      | 0.99  | 11 |
| $\kappa$-casein, g/l | 2.87  | 0.41  | 14 | 2.98      | 0.45  | 15 |
| $\alpha$-lactalbumin, g/l | 0.96  | 0.13  | 14 | 0.96      | 0.14  | 15 |
| $\beta$-lactoglobulin, g/l | 3.28  | 0.61  | 19 | 3.31      | 0.54  | 16 |

Finnish Ayrshire; total 59 cows and 174 observations, for $R$ and $E_{30}^1$: 58 cows and 165 observations and for $K_{20}^0$: 53 cows and 134 observations.

Finnish Friesian; total 55 cows and 155 observations, for $K_{20}^0$: 55 cows and 142 observations.

$\bar{x}$ = mean, s.d. = standard deviation, cv = coefficient of variation $E_{30}^1$ = samples with renneting time, used in statistical analyses.
**Gross and protein composition of milk**

Even though the renneting properties were somewhat weaker in milk of the FAy, the fat, protein and casein contents and the concentrations of $\alpha_1$-, $\alpha_2$- and $\beta$-caseins were higher than in milk of the FFr (Table 2). The somatic cell count and concentration of $\kappa$-casein were higher in milk of the FFr than in that of the FAy. There were no major differences in concentrations of $\alpha$-lactalbumin and $\beta$-lactoglobulin between the FAy and FFr.

**Estimates of genetic variation**

The moderately high heritability estimates for milk renneting properties in both breeds suggested that additive genetic effects made an important contribution to variation in these characteristics (Table 3). When $\kappa$-$\beta$-CN genotypes and $\beta$-LG genotypes were excluded from Model 1, the heritability estimates increased by 3–16 percentage units. A moderate proportion of the additive genetic variation in renneting properties was therefore due to milk protein genotypes. The magnitude of heritability estimates for renneting properties was about the same as that for protein and casein contents and concentrations of $\beta$- and $\kappa$-caseins in both breeds, and for fat content, and concentrations of $\alpha_1$- and $\alpha_2$-caseins and $\beta$-LG in the FAy (Table 3). Because the data sets were small, the standard errors of the heritability estimates were high for some traits, but reasonable for renneting properties.

**Effects of $\kappa$-$\beta$-CN genotypes**

Of the several traits studied, $k$-b-CN genotypes had a statistically significant effect on firmness

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### Table 3. Heritability ($h^2$) and repeatability ($r$) estimates for renneting properties, daily milk yield, and gross and protein composition characteristics of milk from the Finnish Ayrshire and Finnish Friesian.

|                      | Finnish Ayrshire | Finnish Friesian |
|----------------------|------------------|------------------|
|                      | $h^2 \pm \text{s.e.}^1$ | $r$       | $h^2 \pm \text{s.e.}^1$ | $r$       |
| Milk renneting       |                  |                  |                  |                  |
| $R$, min             | 0.62 $\pm$ 0.14 | 0.66             | 0.35 $\pm$ 0.21 | 0.58             |
| $K_{20'}$, min        | 0.54 $\pm$ 0.13 | 0.63             | 0.66 $\pm$ 0.10 | 0.71             |
| $E_{20'}$, mm         | 0.41 $\pm$ 0.19 | 0.64             | 0.57 $\pm$ 0.06 | 0.57             |
| Milk yield and composition |              |                  |                  |                  |
| Daily milk yield, kg | 0.12 $\pm$ 0.11 | 0.47             | 0.06 $\pm$ 0.20 | 0.60             |
| Fat %                | 0.37 $\pm$ 0.07 | 0.57             | 0.14 $\pm$ 0.15 | 0.30             |
| Protein %            | 0.34 $\pm$ 0.23 | 0.59             | 0.19 $\pm$ 0.18 | 0.57             |
| pH                   | 0.08 $\pm$ 0.19 | 0.46             | 0.05 $\pm$ 0.23 | 0.43             |
| Somatic cell count (In) | 0.18 $\pm$ 0.09 | 0.57             | 0.38 $\pm$ 0.10 | 0.41             |
| Protein composition  |                  |                  |                  |                  |
| Casein %             | 0.50 $\pm$ 0.07 | 0.52             | 0.20 $\pm$ 0.25 | 0.56             |
| Whey protein %       | 0.01 $\pm$ 0.04 | 0.37             | 0.15 $\pm$ 0.18 | 0.47             |
| Non-protein nitrogen, mg/g | 0.00 $\pm$ 0.00 | 0.42             | 0.02 $\pm$ 0.06 | 0.17             |
| Casein number        | 0.20 $\pm$ 0.10 | 0.31             | 0.30 $\pm$ 0.35 | 0.33             |
| $\alpha_1$-casein, g/l | 0.52 $\pm$ 0.07 | 0.55             | 0.06 $\pm$ 0.13 | 0.46             |
| $\alpha_2$-casein, g/l | 0.31 $\pm$ 0.22 | 0.46             | 0.00 $\pm$ 0.00 | 0.40             |
| $\beta$-casein, g/l  | 0.40 $\pm$ 0.07 | 0.41             | 0.33 $\pm$ 0.24 | 0.51             |
| $\kappa$-casein, g/l | 0.27 $\pm$ 0.18 | 0.43             | 0.42 $\pm$ 0.19 | 0.61             |
| $\alpha$-lactalbumin, g/l | 0.00 $\pm$ 0.00 | 0.26             | 0.27 $\pm$ 0.08 | 0.27             |
| $\beta$-lactoglobulin, g/l | 0.35 $\pm$ 0.29 | 0.48             | 0.21 $\pm$ 0.14 | 0.54             |

`s.e. = standard error of heritability estimate`
of the curd and concentrations of \( \alpha_{s2} \) and k-caseins in the FAy, and on curd firming time, firmness of the curd and k-casein concentration in the FFr. In the FAy, the ABA1A2, ABA1A3, and AAA1A3 genotypes had a favourable effect on firmness of the curd and k-casein concentration, and the ABA1A2 and AAA1A3 genotypes on \( \alpha_{s2} \)-casein concentration (Table 4). In the FFr, the k-b-CN genotypes associated with the most favourable renneting properties and the highest k-casein concentration were ABA2B, ABA1A3, AAA2A3, ABA1A2 and ABA2A2 (Table 5).

**Effects of \( \beta \)-LG genotypes**

The \( \beta \)-LG genotypes had no statistically significant effect on renneting properties in either breed but they had a strong effect on the protein composition of milk in both breeds (Table 6). Casein content and casein number were highest for the \( \beta \)-LG BB genotype, and whey protein and \( \beta \)-lactoglobulin concentrations for the AA genotype.

### Associations between renneting properties and composition of milk

An increase in the pH of milk had an unfavourable effect on each renneting characteristic in both breeds (Table 7). Some of the milk samples from the FFr had a very high somatic cell count. The somatic cell count did not, however, have a statistically significant effect on renneting properties in either breed. High protein and casein concentrations of milk were a characteristic of the Finnish Ayrshire.

| Table 4. Estimates of \( \kappa \)-\( \beta \)-casein genotype effects (with standard errors below the estimates) on firmness of the curd, and \( \alpha_{s2} \) and k-casein (cn) concentrations of milk from the Finnish Ayrshire. |
|-----------------|----------------|----------------|-----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|-----------------|
| \( \kappa \)-casein | \( \alpha_{s2} \)-casein | k-casein | \( \kappa \) | F-test |
| AA | AA | AA | AA | AB | AB | AB | AE | AE | AE | BE | EE |
| \( \kappa \)-casein | 0.93 | 0.12 | 0.35 | 0.90 | 0.00 | 0.29 | 0.00 | 0.21 | 0.23 | 0.16 | 0.00 | 0.22 | 0.30 | 0.10 | 0.00 | 0.05 | 0.00 | 0.00 | 0.00 | 0.00 |
| AA | 6, 21, 36, 6, 24, 3, 7, 38, 6, 6, 12 |

| Table 5. Estimates of \( \kappa \)-\( \beta \)-casein genotype effects (with standard errors below the estimates) on curd firming time, firmness of the curd and k-casein concentration of milk from the Finnish Friesian. |
|-----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|
| \( \kappa \)-casein | \( \kappa \)-casein | k-casein | \( \kappa \) | F-test |
| AA | AA | AA | AA | AB | AB | AB | AB | AB | AE | AE |
| \( \kappa \)-casein | 0.29 | 0.40 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 |
| \( \kappa \)-casein | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |

| \( \kappa \)-casein | \( \kappa \)-casein | k-casein | \( \kappa \) | F-test |
| AA | AA | AA | AA | AB | AB | AB | AB | AE | AE |
| \( \kappa \)-casein | 0.29 | 0.40 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 | 0.29 | 0.29 | 0.00 |
| \( \kappa \)-casein | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |

1. \( \kappa \)-casein number of observations in genotype classes = 23, 4, 53, 3, 23, 11, 5, 11, 9
Table 6. Estimates of β-lactoglobulin genotype effects (with standard errors below the estimates) on protein composition of milk from the Finnish Ayrshire and the Finnish Friesian.

|                      | Finnish Ayrshire |                  | Finnish Friesian |                  |
|----------------------|-----------------|-----------------|-----------------|-----------------|
|                      | AA n=15         | AB n=82         | BB n=77         | AA n=20         | AB n=97         | BB n=38         | F-test          |                  |
| Casein %             | -0.23 (0.10)    | 0 (0.06)        | 0.06 P=0.018    | -0.11 (0.07)    | 0 (0.05)        | 0.10 P=0.020    |                  |
| Whey protein %       | 0.03 (0.03)     | 0 (0.02)        | -0.08 P<0.001   | 0.08 (0.02)     | 0 (0.02)        | -0.03 P<0.001   |                  |
| Casein number        | -2.38 (0.80)    | 0 (0.45)        | 2.52 P<0.001    | -2.90 (0.73)    | 0 (0.51)        | 1.68 P<0.001    |                  |
| β-Ig, g/l            | 0.34 (0.19)     | 0 (0.11)        | -0.70 P<0.001   | 0.50 (0.15)     | 0 (0.11)        | -0.55 P<0.001   |                  |

β-Ig=β-lactoglobulin

Contents and concentrations of α\textsubscript{32}, β- and κ-caseins and β-lactoglobulin had favourable effects on curd firming time and firmness of the curd in both breeds. In addition, a high concentration of α\textsubscript{s1}-casein and a high casein number in the FF\textsubscript{r} and a high concentration of α-lactalbumin in the FA\textsubscript{y} were favourably associated with milk renneting properties.

The κ-β-CN genotypes with favourable renneting properties were associated with a high κ-casein concentration in both breeds and with a high α\textsubscript{32}-casein concentration in the FA\textsubscript{y}. When

Table 7. Statistically significant regression coefficients of milk renneting properties on daily milk yield, and gross and protein composition characteristics of milk.

|                      | Finnish Ayrshire |                  | Finnish Friesian |                  |
|----------------------|-----------------|-----------------|-----------------|-----------------|
|                      | R               | K\textsubscript{20} | E\textsubscript{30} | R               | K\textsubscript{20} | E\textsubscript{30} |
| Dmy, kg              |                 |                  |                  |                 | -0.1**           |
| Fat %                |                 |                  |                  |                 | -0.6*            |
| Protein %            | 28.4***         | 23.3***          | -41.7***         | 10.5***         | 16.3***          | 10.3**           | -34.5***         |
| pH                   |                 |                  |                  |                 | -5.1***          | 17.4***          |
| Casein %             | -6.4*           | 11.3***          |                  |                  | -5.6***          | 20.6***          |
| Whey protein %       | 12.7***         |                  |                  |                  |                  |                  |
| Npn, mg/g            | -13.9**         |                  |                  |                  |                  |                  |
| Casein number        |                 |                  |                  |                  |                  |                  |
| α\textsubscript{s1}-casein, g/l |                |                  |                  |                  |                  |                  |
| α\textsubscript{32}-casein, g/l | -1.6*           | 3.6*             |                  |                  |                  |                  |
| β-casein, g/l        |                 |                  |                  |                  |                  |                  |
| κ-casein, g/l        | -4.1***         | 8.2***           |                  |                  |                  |                  |
| α-la, g/l            | -11.3***        | 14.4***          |                  |                  |                  |                  |
| β-Ig, g/l            | -3.6***         | 5.3**            |                  |                  |                  |                  |

Dmy=daily milk yield, α-la=α-lactalbumin, β-Ig=β-lactoglobulin
* = P<0.05, ** =P<0.01, *** = P<0.001
Npn=Non-protein nitrogen
Discussion

Genetic variation of characteristics

The small data sets in this study were not suitable for estimating variance components and heritability values for the traits studied. Errors in the heritability values assumed can change the significance levels, and possibly lead to bias in estimates of the effects under study (Kennedy et al. 1992). We, however, used variance components estimated from the data, there being no estimates in the literature of the variance components deduced using a repeatability model. The heritability estimates for renneting properties we obtained were about twice as high as those reported by Lindström et al. (1984) for milk renneting time and by Tervalta et al. (1985) for each milk renneting trait. However, in Tervalta et al. (1985), the standard errors of heritability estimates were very high. In both previous studies, the cows were sampled only once.

Effects of milk protein genotypes

κ-β-CN genotypes with the κ-casein B allele had a favourable effect on firmness of the curd in both breeds. There was, however, some variation between the effects of κ-casein AB, AA and AE genotypes depending on the effect of β-casein genotypes or alleles in genotype combinations. In the FAy the β-casein A,A2 genotype, and in the FFr the β-casein A, and B alleles also had a favourable effect on firmness of the curd.

There was a difference between the effects of the κ-casein A and E alleles on renneting properties. The κ-casein AA genotype had a favourable effect on renneting properties when in combination with the β-casein A1A2 genotype in the FAy and with A,A1 and A2A, genotypes in the FFr. The effect of the κ-casein E allele was, in contrast, rather unfavourable in each κ-β-CN genotype. In the FAy, the κ-casein E allele was rather common (30%), whereas in the FFr it was rare (6%). It is possible that the differences in κ-casein E allele frequency and the κ-casein concentration in milk (Table 2) between the FAy and FFr were partly responsible for the differences in renneting properties between the breeds. A higher frequency of the κ-casein E allele in the FAy than in the FFr was also observed by Ahlfors (unpublished) in data on about 800 FAy and 100 FFr cows.

The favourable effect of the κ-casein B allele on milk renneting properties has been reported in several other studies (e.g., El-Negoumy 1972, Schaar 1984, Schaar et al. 1985, Pagnacco and Caroli 1987, Davoli et al. 1990, Olofss et al. 1992, Van den Berg et al. 1992, Macheboef et al.1993, Walsh et al. 1995). Nothing has previously been known, however, of the effect of the κ-casein E allele on renneting properties. A favourable effect of the β-casein B allele on renneting properties similar to that we observed in the FFr was reported by Feagan et al. (1972). According to Marziali and Ng-Kwai-Hang (1986), β-casein genotypes had no statistically significant effect on renneting properties.

A statistically non-significant effect of β-LG genotypes on renneting properties such as observed in this study was also reported by Feagan et al. (1972) and Pagnacco and Caroli (1987). According to van den Berg et al. (1992), the β-LG AA genotype was associated with the shortest renneting and curd firming times. The favour-
able effect of the β-LG BB genotype on casein concentration and casein number, and that of the AA genotype on whey protein and β-lactoglobulin concentrations were also reported by McLean et al. (1984) and Schaar et al. (1985).

As well as renneting properties, the milk samples were analysed for several gross and protein composition characteristics to establish whether the variation in renneting properties due to milk protein genotypes could be explained by differences in gross or protein composition characteristics between the genotypes. Of the several characteristics, the high κ-casein concentration in milk explained part of the favourable effect of certain κ-β-CN genotypes on the renneting properties in both breeds.

We estimated the direct effects of milk protein genotypes on renneting properties by assuming an animal model. We did this because the results for the κ-casein genotype effects on renneting properties are consistent suggesting that the κ-casein locus itself affects the renneting properties. There are no previous reports of an animal model being used for estimating the effects of milk protein genotypes on renneting properties. It is, however, possible that there are other quantitative trait loci near the κ-casein locus that have a considerable effect on renneting properties. Thus, it would be interesting to estimate associations between milk protein genotypes and renneting properties within sires. We could not do so here due to the restricted size of the data sets.

Conclusions

The κ-β-CN genotypes ABA₁A₂, ABA₁A₃ and AAA₁A₂ in the FAy and genotypes ABA₂B, AAA₁A₃, AAA₂A₃, ABA₁A₂, ABA₂A₂ in the FFr were associated with favourable renneting properties, partly due to their association with the high κ-casein concentration in the milk. The effect of the κ-casein E allele on renneting properties was unfavourable as compared with that of the κ-casein B allele, and possibly also with that of the κ-casein A allele. Results for the effect of the κ-casein E allele on renneting properties need to be confirmed with a larger data set. The β-LG genotypes had no effect on renneting properties but they had a strong effect on the protein composition of milk.

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Kaseiinien yhdistelmägenotypyyppien ja β-laktoglobuliinin genotypyyppien vaikutus maidon juoksettumisominaisuuksiin ja koostumukseen

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Tutkimuksen tarkoituksena oli selvittää κ- ja β-kaseiinien yhdistelmägenotypyyppien ja β-laktoglobuliinin genotypyyppien vaikutusta maidon juoksettumisominaisuuksiin sekä maidon yleis- ja valkuaisainekoostumukseen. Tutkimuksen aineisto koostui Helsingin yliopiston Viikin opetus- ja tutkimustilan 59 ayrshirelehmän 174 maitonäytteestä ja Suitian opetus- ja tutkimustilan 55 friisiläislehmän 155 maitonäytteestä. Maitonäytteitä kerättiin lehmiltä kolme kertaa lyypsykauden aikana; kuukauden, kolmen kuukauden ja viiden kuukauden kuluttua poikimisesta. κ- ja β-kaseiinien yhdistelmägenotypyyppien ja β-laktoglobuliinin genotypyyppien vaikutusten selvittämiseen käytettiin eläinmallia, jossa kiinteinä tekijöinä olivat poikimakerta, lyypsykauden vaihe, vuodenaika, κ- ja β-kaseiinien yhdistelmägenotypyyppit ja β-laktoglobuliinin genotypyyppit. Satunnaisina tekijöinä mallissa olivat eläimen additiivinen geneettinen vaikutus, eläimen pysyvä ympäristö ja jäännöstekijä.

Kaseiinien kytketyyneisyyden vuoksi havaintojen lukumäärä eri κ- ja β-kaseiinien yhdistelmägenotypyypeissä vaihteli selkeästi. Ayrshirelehmillä maidon juoksettumisominaisuudet olivat parhaimmat κ- ja β-kaseiinien yhdistelmägenotypyyppillä ABA_A₃, ABA_A₁ ja AAA_A₃ ja friisiläislehmillä yhdistelmägenotypyyppillä ABA_B, AAA_A₁, AAA_A₂, ABA_A₂ ja ABA_A₂. κ-kaseiinin E-alleeli vaikuttii epäedullisesti maidon juoksettumisominaisuuksiin B-alleeliin ja luultavasti myös A-alleeliin verrattuna. Juoksettumisominaisuuksiltaan parhaimmilla yhdistelmägenotypyypeillä oli yhteys maidon korkeaan κ-kaseiinipitoisuuteen kummallakin rodulla. Maidon κ-kaseiinipitoisuus vaikutti edullisesti maidon kiinteytymisäikään sekä juoksettuman kiinteyteen kummallakin rodulla. Osa yhdistelmägenotypyyppien vaikutuksesta maidon juoksettumisominaisuuksiin johtui siten niiden vaikutuksesta maidon kaseiinikoostumukseen. β-laktoglobuliinin genotypyyppillä ei ollut vaikutusta maidon juoksettumisominaisuuksiin. β-laktoglobuliinin genotypyyppit vaikuttivat kuitenkin maidon valkuaisainekoostumukseen. β-laktoglobuliinin AA genotypyyppi oli yhteydessä maidon korkeaan heraproteiini- ja β-laktoglobuliininpitosuuteen ja BB genotypyyppi korkeaan kaseiinipitosuuteen sekä kaseiinilukuun.