

The association between CSN3 genotypes and milk production parameters in Czech Pied cattle

Kučerová Jitka^{*1} (jitka.k@seznam.cz), Němcová Eva¹, Štípková Miloslava¹, Jandurová Olga¹, Matějčíček Aleš² and Bouška Josef¹,

¹Research Institute of Animal Production, P.O.Box 1, 104 01 Prague 10, Czech Republic,

²University of South Bohemia, Czech Republic

Abstract

The aim of this study was to detect an association between CSN3 genotypes and milk production parameters. Data on CSN3 genotypes in Czech Pied sires and their breeding values for milk yield, protein yield, fat yield, protein content and fat content were available. Four genotypes (AA, AB, BB and BE) of kappa-casein marker (CSN3) were observed in the investigated population. Genotypes BB and BE were associated with higher protein content, but lower milk production. In contrast to this, genotypes AA and AB were associated with higher milk production, but lower protein and fat content. Further analyses using granddaughter design population structure will follow this study.

Introduction

Marker polymorphisms related to the performance parameters can be taken into account in selection procedures. Genetic markers explain a part of variance and improve the estimation of breeding value (Příbyl et Hruban, 2000). Marker for kappa-casein (CSN3) is one of the relevant markers related to milk production. The objective of this study was to detect an association between CSN3 genotypes and milk production parameters in Czech Pied cattle.

Material and methods

120 sires were genotyped for marker CSN3 using a method PCR-RFLP. Breeding values for milk, protein and fat yield and protein and fat content were available. Relation between marker CSN3 and milk production parameters was analysed in SAS 8.2 by use of GLM:

$$y_{ijklm} = \mu + C_i + bR_j + bP_k + O_l(C_i) + e_{ijklm}$$

where y is the breeding value of sire,

μ is the average of observed breeding value,

C is the effect of CSN3,

bR is the regression on year of sire birth,

bP is the regression on share of Czech Pied breed in sires,

O(C) is the effect of grandsire with nested effect of CSN3,

e is the set of residual effects.

Results and discussion

Genotype AB was found as the most frequent (49,2 %). Detected frequencies of genotypes AA, BB and BE were 28.3 %, 19.2 % and 3.3 %, respectively. Observed allele frequencies were 0.53 (allele A), 0.45 (allele B) and 0.02 (allele E). Compared to this, Hanuš et al. (2000a) found frequencies of alleles A 0.65 and B 0.35 in observed population of Czech Pied cattle. Neubauerová (2001) reported that allele A tends to prevail especially in milk breeds and improves milk production. Freyer et al. (1999) found prevalent frequency of genotype AA (54 %), frequency of genotype AB 27 %, BB only 2 %, AE 13 %, BE 3 % and EE 1 % in German Holstein cattle. According to studies of Boettcher et al. (2004) and Caroli et al. (2004), allele B increases milk protein content and improves milk protein quality. A negative

effect of allele E on milk production parameters was reported by Ikonen et al. (1997) and Leone et al. (1998).

Sires with *CSN3* genotype AA significantly increased milk, protein and fat yield but decreased protein and fat content compared to the opposite tendency found in sires with genotype BE. Sires with genotype BB increased protein content as well as protein yield but slightly decreased milk and fat yield and fat content. The effects of genotype BB on higher protein content but lower milk yield as well as the effects of genotype AA on higher milk yield but lower protein content were reported by Žitný et al. (1996), Chrenek et al. (1998), Hanuš et al. (2000b) and Kučerová et al. (2004). Contrary to our study, Neubauerová (2001) did not find any significant association between *CSN3* genotypes and breeding values for milk production parameters.

Conclusions

Genotypes BB and BE were associated with higher protein content but lower milk production. In contrast to this, genotypes AA and AB were associated with higher milk production but lower protein and fat content.

Low protein yield was linked with genotype BE compared to genotypes AA, AB and BB.

The financial support from the project NAZV IG46086 is acknowledged.

References

- Boettcher, P. J.; Carovi, A.; Stella, A.; Chessa, S.; Budelli, E.; Canavesi, F.; Ghiroldi, S. et Pagnacco, G. (2004): Effects of Casein Haplotypes on Milk Production Traits in Italian Holstein and Brown Weiss Cattle. *J. Dairy Sci.*, 87, 4311-4317.
- Caroli, A.; Chessa, S.; Bolla, P.; Budelli, E. et Gandini, G. C. (2004): Genetic structure of milk protein polymorphism and effects on milk production traits in a local dairy cattle. *J. Anim. Breed. Genet.*, 121, 119-127.
- Chrenek, P. et al. (1998): Milk production characteristics of Slovak Pied cows in relation to κ -casein genotype. *J. Farm. Anim. Sci.*, 31, 9-12.
- Freyer, G.; Liu, Z.; Erhardt, G. et Panicke, L. (1999): Casein polymorphism and relation between milk production traits. *J. Anim. Breed. Genet.*, 116, 87-97.
- Hanuš, O.; Beber, K. et Kopecký, J. (2000a): Milk protein variants and characteristics of milk and cows. In.: Collection of Scientific Papers „Breeding, nutritional and technological aspects of milk production and duality“. Rapotín, 47-49.
- Hanuš, O.; Beber, K.; Čermák, V.; Kopecký, J. et Jedelská, R. (2000b): Types of milk proteins influence performance, longevity and health of cows. In.: Collection of Scientific Papers „Breeding, nutritional and technological aspects of milk production and duality“. Rapotín, 53-57.
- Ikonen, T.; Ojala, M. et Syvaaja, E.L. (1997): Effects of composite casein and beta-lactoglobulin genotypes on renneting properties and composition of bovine milk by assuming an animal model. *Agricultural and Food Science in Finland*, 6, 283-294.
- Kučerová, J.; Němcová, E.; Štípková, M.; Vrtková, I.; Frelich, J. et Bouška, J. (2004): The relation between genetic marker *CSN3* and milk production parameters in Czech Pied cattle. *Proceeding Book from the 11th International Congress on Biotechnology in Animal Reproduction*, 153-157.
- Leone, P. et al. (1998): Effects of the CASK E variant on milk yield indexes in Italian Holstein Friesian Bulls. *26th Conference of ISAG, Animal Genet.*, 29, 63.
- Neubauerová, V. (2001): Detekce genetických markerů a možnosti jejich využití u skotu a dalších kopytníků. Thesis. University of South Bohemia. České Budějovice, 211.

Příbyl, J. et Hruban, V. (2000): QTL in animal breeding. In: Collection of Scientific Papers "Present and Perspectives in Animal Breeding ", ČZU v Praze, 9.11.2000, 122-132.

Žitný, J.; Trakovická, A.; Kúbek, A.; Michaličková, E. et Ostertag, I. (1996): Differences in milk efficiency of different kappa-casein genotypes of dairy cows of the Slovak Pied breed. Živ. výr., 41, 533-538.

Table 1: Basic statistics of breeding values for milk production parameters observed in sires

Parameter	Breeding values of sires for				
	Milk yield (kg)	Protein content (%)	Protein yield (kg)	Fat content (%)	Fat yield (kg)
\bar{x}	127	0,00	13	-0,05	2
s_x	423	0,12	15	0,19	18
Min.	-1162	-0,45	-42	-0,55	-55
Max.	1164	0,28	45	0,50	43

Figure 1: Average breeding values for milk production (kg) according to genotype groups of CSN3

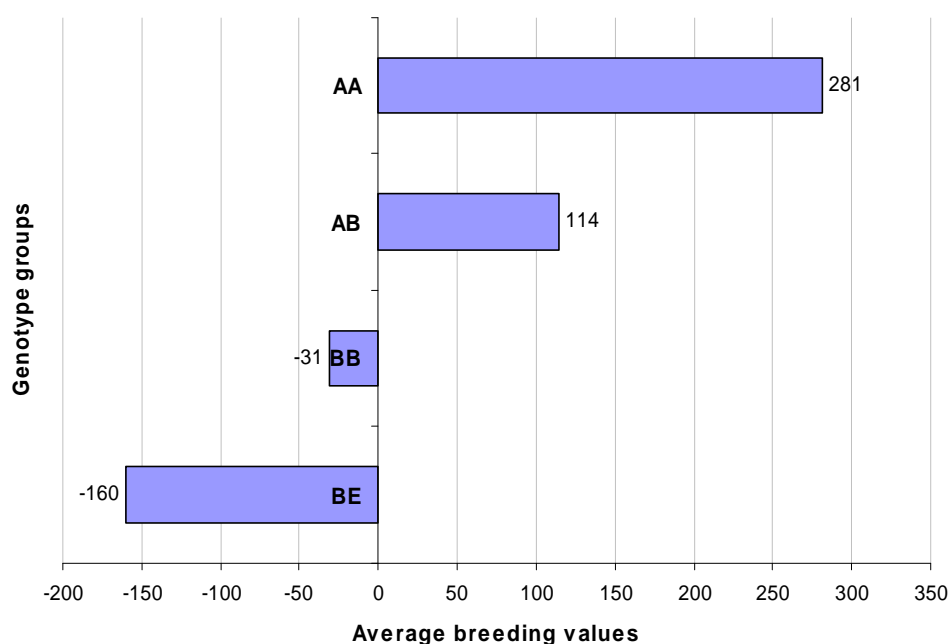


Figure 2: Average breeding values for protein and fat yield according to genotype groups of *CSN3*

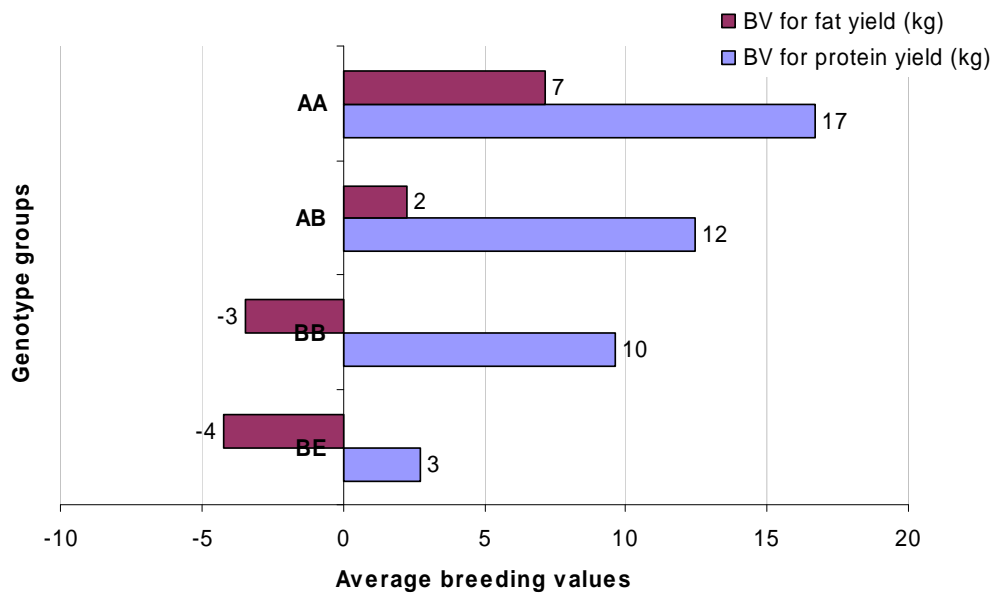


Figure 3: Average breeding values for protein and fat content according to genotype groups of *CSN3*

