

THE EFFICIENCY OF PREDICTION OF PSE PORK MEAT ON THE BASIS OF HALOTHANE TEST AND HAL-GPI-A1BG-PGD HAPLOTYPING AND PCR/RFLP ANALYSIS

KURYŁ J.*, KOŁWIN-PODSIADŁA M.** and PRZYBYLSKI W.**

* Institute of Genetics and Animal Breeding, Polish Academy of Science, Jastrzêbiec, Mroków Poland. ** Agricultural and Pedagogical University, Faculty of Agriculture, Institute of Animal Breeding and Technology of Animal Production, Pig Breeding Department, Poland

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SUMMARY

The aim of studies was to examine the efficiency of prediction of PSE meat occurrence in German Landrace herd on the basis of HAL genotype defined by halothane test and HAL-GPI-A1BG-PGD haplotyping (Gahne and Juneja 1985) (Method 1) as compare to PCR/RFLP test (Fuji et al. 1992) (Method 2). A total 105 animals were evaluated. The HAL genotype within tested material identified by Method 1 was as follows: 30 NN (28,6%); 43 Nn (40,9%); 32 nn (30,5%) while by Method 2 - 25 NN (23,8%); 51 Nn (48,6%); 29 nn (27,6%). The genotype of 27 (25.7%) pigs was corrected by PCR/RFLP test. The PSE meat (qualified on the basis of pH₁ and R₁), was identified in 71,9% of animals of nn genotype defined by Method 1, whereas within group of the same HAL genotype but stated by Method 2 - 79,3% of pigs showed PSE meat post mortem. The meat quality rates (pH₁, R₁, meat colour) showed more profound changes within nn animals in favor of Method 2, but those differences found out insignificance.

Introduction

Various methods of predicting of meat quality in pigs have been described to reduce and eliminate the halothane sensitivity and PSE pork (see review Claes et al., 1992). Stress sensitivity to malignant hyperthermia induced by halothane is genetically determined. Malignant hyperthermia has been shown to be controlled by a recessive gene at a single autosomal locus HAL (Ollivier et al., 1975). The method of HAL genotype identification based on halothane test and typing of proteins polymorphism coded by genes linked to HAL locus was elaborated by Gahne and Juneja in 1985.

The ryanodine receptor of the sarcoplasmic reticulum calcium channel gene has been postulated to be the candidate for predisposition to porcine malignant hyperthermia. A single point mutation in ryanodine receptor gene (RYR 1) namely a cytosine (C) to thymine (T) at position 1843, led to an alteration in amino acid sequence from arginine at position 615 in the HAL^NHal^N pigs to cysteine in the HALⁿHalⁿ individuals. The genotyping test developed by Fuji et al. (1991) involved an amplification of RYR 1 gene fragment comprising 1843 nucleotide followed by HgiAI restriction endonuclease digestion. Electrophoretic separation of digested DNA fragments allowed HAL genotype identification (PCR/RFLP test). The association between HAL genotype determined by haplotyping method (Gahne and Juneja, 1985) and meat quality has been shown by many laboratories (Sellier, 1987).

A similar studies are actually realized using HAL genotyping by PCR/RFLP test (Pommier and Houde, 1993).

The aim of presented study was to examine the efficiency of prediction of PSE meat occurrence within German Landrace herd on the basis of HAL genotype defined by haplotyping as compare to PCR/RFLP test.

Material And Methods

The investigations covered 105 German Landrace pigs from one herd. The HAL genotype was identified by HAL-GPI-A1BG-PGD haplotyping (according to Gahne and Juneja, 1985) and by PCR/RFLP test (according to Fuji et al., 1991).

The pigs were killed by the live weight about 100kg. Meat quality was evaluated on the basis of value of parameters: pH_i and R_i measured in *M. longissimus dorsi* at 45 minutes post mortem. Meat brightness was determined using an apparatus Momcolor-D3098 with white standard.

PSE meat was classified on the basis of pH_i and R_i (Honikel and Fisher, 1977 in modification of Koæwin-Podsiad'a and Chmura-Janowiak, 1988).

Data were analyzed using classical procedures of one-way variance analysis and means were compared using Tukey test.

The association between pH_i , R_i values and PSE meat occurrence on one side and HAL⁺HAL⁺ genotype on other was calculated according to 2x2 containing table (Vögeli and Schwörer, 1982).

Results And Discussion

The HAL genotypes within tested material identified by HAL-GPI-A1BG-PGD haplotyping were as follows: 30 NN (28,6%), 43 Nn (40,9%), 32 nn (30,5%) while by PCR/RFLP test - 25 NN (23,8%); 51 Nn (48,6%); 29 nn (27,6%). The genotype of 27 pigs was corrected by PCR/RFLP test. The PSE meat (qualified on the basis of pH_i and R_i) was identified in 71,9% of animals of nn genotype defined by haplotyping whereas within by PCR/RFLP test - 79,3% of pigs - showed PSE meat post mortem (Table 1). About 20% of Nn animals developed PSE meat post mortem (Table 1). The necessity of explanation of this phenomenon is very significant for pig breeders.

The meat quality parameters (pH_i , R_i , meat colour) showed more profound changes within nn animals in favour of Method 2, but those differences found out insignificant on basis of variance analysis. The differences between R_i values determined for both NN and Nn pigs appeared to be insignificant when HAL genotype was typed by haplotyping method, whereas differed significantly between pigs typed by PCR/RFLP test (Table 1).

The linkage disequilibrium coefficient values shown in Table 2 confirmed the association between nn genotype and values of pH_i as well as PSE meat occurrence. However, the D_i value determined for HAL⁺HAL⁺ - pH_i pair (0,92) suggested more significant association between pH_i and HAL genotype defined by PCR/RFLP test.

The disagreement between HAL genotype and meat quality was observed regarding 3 animals. Two nn pigs showed normal meat quality ($pH_i=6,06$ and $R_i=1,037$, meat colour 19,96). Third pig of Nn genotype developed PSE meat post mortem ($pH_i=5,56$, $R_i=1,256$, meat colour 21,97) what was signalized in literature earlier (Koæwin-Podsiad'a et al., 1993).

The pH_i value determined for six Nn animals was below 6,0 (5,60 - 5,92) while the value of other parameters were characteristic for normal meat (R_i from 0,771 to 1,025; meat colour from 14,9 to 16,09). Altogether 9 animals (8,5%) showed more or less disagreement between HAL genotype and values of meat quality parameters. The presented results led us to conclusion that additional factors affected meat quality in pigs.

Conclusions

1. PCR/RFLP test appeared to be more efficient method for porcine meat quality prediction as compare to haplotyping test.
2. Additional factors probably affects the value of pH_i , R_i and meat colour leading to disagreement between HAL genotype and meat quality observed for some pigs.

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