A Robertsonian translocation in swine

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Summary

A cytogenetic survey was carried out on fattening male and female pigs of different lines in a local herd and on A.I. boars of German Democratic Republic. In 5 of 66 fattening animals and 4 of 461 A.I. boars analysed, a Robertsonian translocation was observed. The translocation was identified as a 13/17 fusion translocation by the G-banding technique. These animals were translocation heterozygotes with 2n = 37, XX or XY, t+ (13 q, 17q). The results obtained are discussed.

Key words : Fusion translocation, swine.

I. Introduction

Robertsonian translocations are distributed widely in cattle. This type of chromosome aberration was first reported as the 1/29 translocation which resulted from a centric fusion between the largest and the smallest autosomes in the Swedish Red and White cattle (GUSTAVSSON & ROCKBORN, 1964). Since then, the same or other types of Robertsonian translocation have been found in many cattle breeds. In contrast, in swine
reciprocal translocations are more common. So, in the last ten years several types of reciprocal translocations were described (Popescu & Legault, 1979; Förster et al., 1981; Golisch et al., 1982; Gustavsson et al., 1983; Popescu et al., 1983). The first information about the occurrence of Robertsonian translocations in swine (Sus scrofa) was given by Miyake et al. (1977). In this report we present information about a Robertsonian translocation in swine observed in a local unselected pig herd and in A.I. boars in GDR.

II. Materials and methods

Blood samples were obtained from 66 female and male animals of different lines raised on a fattening farm of the southern region of GDR, and from 461 A.I. boars of a breeding station. Each sample (2.0 ml) was incubated at 37 °C for 48 hrs in 8.0 ml Parker medium supplemented with 0.1 ml PHA (Welcome) and antibiotics. 1 ml of 0.001 p. 100 colchicine solution was added at 2.0 hrs before the termination of culture. After hypotonic treatment with 0.56 p. 100 KCl solution, cells were fixed in methanol-acetic-acid (3 : 1), and air dried on a slide glass. Trypsin G-band- (Seabright, 1971) and C-band-techniques (Sumner, 1972) were applied for additional identification of the chromosomes. A total of 30 metaphase cells from each animal was analysed with the ordinary non-banding method. Those with abnormal complements were further studied by G-banding methods.

III. Results

The present chromosome analysis was based on the G-band karyotype established by the standard system of Reading (Ford et al., 1980).

Table 1

Results of cytogenetic analysis in a local herd of fattening pigs of different cross breeding lines in GDR.

| Chromosome number and karyotype | Number of animals | Percentage (%) |
|-------------------------------|------------------|----------------|
| 2n = 38, XY or XX, normal      | 61               | 92.4 (83 ; 97) |
| 2n = 37, XY or XX, t+ (13 q, 17 q) | 5                | 7.6 ( 3 ; 17) |
| Total                         | 66               | 100.0          |

Confidence intervals are given at the level 0.95.

Out of the 66 unselected pigs, examined, 61 had the normal karyotype with 2n = 38, XY or XX (table 1). The remaining 5 animals showed a reduction of the diploid number from 38 to 37 chromosomes in all the cells studied. An additional large submetacentric chromosome was observed in their metaphases (fig. 1). Based on the
Giems stained metaphase chromosomes and karyotype of a male pig (2n = 37, XY, t+(13q, 17q)).

Fig. 1

G-banded metaphase chromosomes and karyotype of a female pig (2n = 37, XX, t+(13q, 17q)).

Fig. 2
results of karyotype analysis with G-banding technique it was confirmed that the large submetacentric chromosomes resulted from Robertsonian translocation between the largest acrocentric chromosome and a small acrocentric chromosome. The short and long arms of the fusion chromosome were identified by their sizes and characteristic G-banding patterns (see fig. 1, 2, 3) as chromosomes of the pairs 13 and 17. The 13/17 translocation also was characterized by having a single centromeric block (fig. 3) of C-banded constitutive heterochromatin. The detailed karyotype analysis of the fattening pigs examined is summarized in table 1. The translocation carriers were heterozygotes with the karyotype of $2n = 37$, XX or XY, $t^{+} (13\ q, 17\ q)$.

TABLE 2

*Frequency of the 13/17 Robertsonian translocation in A.I. boars in GDR.*

| Breeds                     | Number of animals analysed | Number of animals with a normal karyotype $2n = 38$, XY | Number of animals with a translocation karyotype $2n = 37$, XY, $t^{+} (13q, 17q)$ |
|----------------------------|----------------------------|-------------------------------------------------------|----------------------------------------------------------------------------------|
| Landrace                   | 62                         | 58                                                    | 4 (6.4%) ($3$ ; $15$ )                                                           |
| « Edelschwein » (Yorkshire)| 276                        | 276                                                   | 0                                                                                 |
| Crossbreeding lines        | 123                        | 123                                                   | 0 ($0.0$ ; $1.0$)                                                                 |
| Altogether                 | 461                        | 457                                                   | 4 (0.9%) ($0.3$ ; $2.2$)                                                          |

Confidence intervals are given at the level 0.95.
The frequency of the chromosomally aberrant animals was 7.6 p. 100 with a confidence interval (at the level 0.95) of (3.0 ; 17.0). Probably this frequency is not representative for the common pig population in GDR.

Because of the lack of information about the family connections of the fattening pigs analysed and with the aim of verifying the estimate of frequency 461 A.I. boars of different races were cytogenetically investigated. The results of this population survey are shown in table 2. The mean frequency of translocation carriers in the A.I. boars analysed is about 8 times lower than in the unselected animals from a local herd which had been studied. However, it is remarkable that this type of aberration was found only in A.I. boars of the Landrace. Among 62 Landrace animals analysed, 4 boars showed a heterozygote 13/17 Robertsonian translocation. In the other breeds investigated, this translocation was absent. In an additional study of familial relationships it could be shown that all chromosomally aberrant A.I. boars have a common ancestor (see fig. 4).

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**Fig. 4**

*Pedigree of four A.I. boars with a heterozygote 13/17 Robertsonian translocation.*
IV. Discussion and conclusion

With the observed 13/17 Robertsonian translocation in swine this type of aberration could be described for the third time. The present results supply additional hints that Robertsonian translocation plays a role in the evolution of swine, similar to the previously mentioned 1/29 translocation in cattle (Gustavsson, 1979; Mayr, 1977). The 15/17 and 16/17 fusion translocations observed in European and Asian wild pigs respectively (Mayr et al., 1984; Troshina & Tikhonov, 1980) support this assumption.

The mean frequency of the Robertsonian translocation in the sample of A.I. boars was 0.9%. However, with the detailed analysis of breeds it could be shown that the 13/17 translocation probably is present only in the Landrace boars of GDR, and therefore occurs in this race more frequently. Considering its distribution and C-banding pattern, it may be suggested that the 13/17 monocentric translocation occurred earlier in the Landrace of GDR. The increased local use of an aberrant boar in artificial insemination can lead to higher frequency, as could be observed in the present study in a local herd of fattening pigs.

The frequency of 6.45 p. 100 of 13/17 translocation carriers among the A.I. boars of the Landrace indicates that possible effects on performance do not effect an elimination of the translocation from breeding animals.

All the carriers in this study were phenotypically normal, and no abnormality was observed on the condition of their health. To evaluate the potential effects of the 13/17 Robertsonian translocation on other economic traits, especially on fertility, further analyses are in progress now (see Golisch et al., 1986).

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