Original article

Sequential GTG-RBA banding pattern in prometaphase chromosomes of cattle (Bos taurus L)

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Summary – A sequential GTG–RBA banding procedure, performed for the first time on the same prometaphase chromosomes of cattle, is presented with the aim of establishing correlations between G and R bands. The results of the present investigation contributed to the establishment of new standard GTG and RBA-banded karyotypes at prometaphase level, useful for the precise identification of chromosomal abnormalities, comparative cytogenetics and gene mapping in the species *Bos taurus* L.

sequential banding / GTG-bands / RBA-bands / standardization / cattle

Résumé – Une nouvelle technique de mise en évidence séquentielle des bandes G et R sur les chromosomes prométaphasiques du bovin (Bos taurus L). Une nouvelle technique de mise en évidence séquentielle de bandes G et R est présentée pour la première fois sur les chromosomes prométaphasiques du bovin dans le but d'établir des corrélations entre les bandes G et R. Les résultats de cette étude ont contribué à l'établissement de nouveaux caryotypes standards en bandes G et R au niveau prométaphasique. Cette étude est utile pour l'identification précise des anomalies chromosomiques, pour les études de cytogénétique comparée et pour la localisation des gènes de Bos taurus.

bandes séquentielles / bandes G / bandes R / standardisation / bovin

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INTRODUCTION

The Reading conference on the standardization of banded karyotypes of domestic animals (Ford *et al*, 1980), undoubtedly one of the most important steps in the history of animal cytogenetics, provided for the first time the 'standard' G-banded karyotype of cattle (*Bos taurus* L) as well as that of other domestic species. Despite its excellent quality, the bovine standard G-banded karyotype soon revealed some limitations for chromosome identification, due mainly to the degree of contraction of the chromosomes used for the standard and, secondarily, to an intrinsic feature of bovine chromosomes, common to all the Bovidae, *ie* G-negative centromeres and telomeres.

These problems have made it necessary to improve the Reading standard by using less contracted chromosomes and to adopt alternative banding techniques such as RBA and QFQ for standardization of new karyotypes which could be used more widely by the scientific community.

This paper reports the results of the sequential GTG-RBA banding technique performed for the first time on the same prometaphase chromosomes of cattle in order to establish correlation between the GTG and the RBA bands. These results provided an important part of the material used for the discussions at the 2nd International Conference for the Standardization of Banded Karyotypes in Domestic Animals, held at Jouy-en-Josas (France) the 22–26th of May, 1989 (ISCNDA, 1989).

MATERIAL AND METHODS

Peripheral blood, drawn from the jugular vein of 5 young bulls of the Italian Friesian breed, was cultured at 37.5° C for 72 h in RPMI 1640 medium (Flow, Dutch modification) supplemented with 10% fetal calf serum (Gibco), 0.1% L-glutamine and 0.1 ml of pokeweed mitogen (Gibco). 6.5 h before harvesting cells, 5'-bromodeoxy-uridine (BUdR, Sigma) at a final concentration of 20 μ g/ml and 5.5 h later colcemid solution (Gibco, final concentration of 0.03 μ g/ml) were added. In order to facilitate the spreading of the prometaphase chromosomes the cell suspension was subjected to a stronger hypotonic shock than usual (0.05 M KCl) at 37.5° C for 20 min and fixed with methanol-acetic acid solution (3:1) for 1 h, centrifuged, fixed again and left overnight in the refrigerator. Air-dried slides were prepared.

Sequential GTG-RBA banding procedure

The air-dried slides, 3–5-d old, were treated for GTG banding according to Lin *et al* (1977); soon after the Giemsa staining, the slides were flooded with phosphate buffer (pH = 7), covered with a coverslip and examined with a Leitz Dialux under bright field optics. The best G-banded prometaphase spreads were selected and photographed with a Kodak microfilm 1454. After microphotography, the coverslip was removed, the slide was destained gently in 30% ethanol for 10 min, washed in distilled water, air-dried and stained with an acridine orange solution (0.2% in phosphate buffer) for 15 min, washed again in tap water, mounted in the same buffer and sealed with paraffin (Di Berardino *et al*, 1979). The prometaphase

spreads previously examined for GTG banding were relocated and photographed again for RBA banding with the same Kodak microfilm 1454. Kodabrome F2M and F3M papers were used for printing GTG and RBA banded prometaphase spreads, respectively.

RESULTS

Figure 1 (A and B) shows, respectively, a GTG-banded prometaphase spread of cattle (2n = 60, XY) and the same spread sequentially stained for RBA-banding. In order to verify the correspondence between the Reading standard and the prometaphase G-banding pattern, individual chromosomes from figure 1A were arranged, side by side, with the G-banded chromosomes of the Reading standard, as shown in figure 2. From this figure it is possible to verify the great advantage and usefulness of using prometaphase instead of metaphase chromosomes, especially for the identification of the smallest autosomes ranking from pairs No 21–29. All the G-banded prometaphase chromosomes fit very well to the Reading standard, thus providing more information for a definite characterization and identification of individual chromosomes of the species. However, because of chromosome contraction of the G-banded cattle chromosomes reported by the Reading standard, it is quite difficult to distinguish among the chromosomes Nos 25, 27 and 29; hence, it is not fully evident that the actual prometaphase pairs correspond to the ones of the Reading standard.

In order to examine in detail the correlation between the GTG and the RBA banding pattern of individual prometaphase chromosomes of cattle, figure 3 (A, B and C) was prepared in which the GTG-banded prometaphase chromosomes from figure 1A (b and d) are compared with the sequentially stained RBA-banded chromosomes (a and e) and with the 'direct' RBA banded chromosomes (c). From this figure it is possible to verify the correct correspondence between G and R bands in almost all of the chromosomes, including the X and Y sex chromosomes.

DISCUSSION

The sequential GTG-RBA banding procedure, performed for the first time on chromosome of cattle, is suitable for a specific characterization of individual chromosomes of this species. Previous contributions on the RBA banding pattern in cattle chromosomes (Popescu, 1975; Gustavsson and Hagelthorn, 1976; De Giovanni *et al*, 1979; 1988; Di Berardino *et al*, 1979, 1983, 1985a, 1985b; Di Berardino and Iannuzzi, 1982) reported karyotypes which were based, as far as possible, on the Reading G-banded standard karyotype, but a direct correlation between G and R bands has not so far been reported. Recently, a G- and R-banding comparison of cattle prometaphase chromosomes arranged according to the Reading system has been reported (Iannuzzi, 1990) but without use of a sequential G-R banding procedure.

The present investigation was carried out in order to make correlations between G and R bands on the same chromosome preparation, thus providing the necessary information for the definition of new standard GTG and RBA banded karyotypes for the species *Bos taurus* L.



Fig 1. (A) GTG-banded prometaphase spread of cattle (2n = 60, XY); (B) the same spread sequentially stained for RBA-banding.

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Fig 2. Correspondence between the GTG-banded chromosomes of the Reading standard (at the center of each pair) and the GTG-banded prometaphase chromosomes of figure 1A (right and left).

In the sequential procedure reported here, the BUdR incorporation necessary to achieve the RBA banding did not seem to affect the G-banding pattern. Also the trypsin treatment used for GTG-banding did not produce significant effects on the quality of the RBA-banding pattern in almost all of the chromosomes. Therefore, this procedure could also be used for the standardization of GTG and RBA-banded



Fig 3. (A, B, C), Comparison of the GTG-banded prometaphase chromosomes of figure 1A (b and d) with the sequentially stained RBA-banded chromosomes of figure 1B (a and e) and the direct RBA-banded chromosomes (c).

karyotypes of other domestic species for which extensive cytogenetic material is already available.

Other ways using G- and R-banded marker chromosomes, such as centric fusions, translocations and nucleolus organizer chromosomes of Bovidae, as well as the biarmed chromosomes of related species, could provide additional information for a definitive characterization of the banding pattern of individual chromosomes of the species *Bos taurus* L.

The results of the present paper contributed to the definition of the 'standard' GTG and RBA-banded karyotypes and idiograms of cattle at the prometaphase level (ISCNDA, 1989) useful for precise description and identification of numerical as well as structural chromosomal aberrations, comparative cytogenetics and gene mapping.

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