

# NOR association in *Canis familiaris*

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## INTRODUCTION

The domestic dog has 78 chromosomes (Gustavsson, 1964). Because of the large number of chromosomes of almost similar size and morphology, a precise analysis of the karyotype is rather difficult (Howard-Peebles and Pryor, 1980; Manolache *et al*, 1976; Wurster-Hill and Centerwall, 1982). However, a G-banded karyotype of *Canis lupus* (Wayne *et al*, 1987), which is supported to be homologue to that of *Canis familiaris*, displayed detailed information about banding patterns and served as a guideline for the numbering system used in the RBG-banded karyotype (Poulsen *et al*, in press) and in the present study. Several authors have reported on the nucleolar organizer regions (NORs) and NOR associations in the karyotype of the domestic dog (Kopp *et al*, 1982a, b; Pathak *et al*, 1982). In this paper, the frequencies of NOR associations and the localization of NORs in the female karyotype of *Canis familiaris* are presented using sequential RBG-banding and the silver (Ag)-NOR staining technique.

## MATERIALS AND METHODS

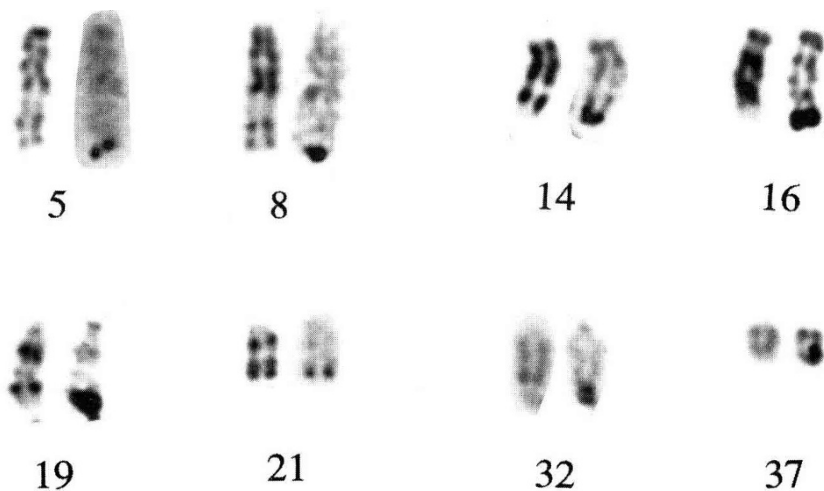
The donor animals were selected at random from the laboratory *Beagles* at the Institute of Biomedicine, Odense University. Peripheral blood samples from 3 female dogs were cultured and processed for RBG-band induction as previously described (Rønne 1985; Poulsen *et al*, in press). Ag-NOR staining was modified after Howell and Black (1980) and used to stain previously RBG-banded metaphases.

From each donor animal, 10 selected RBG-banded metaphase and prometaphase plates with good spreading and well-defined bands were photographed, registered and karyotyped according to Poulsen *et al* (in press). The selected metaphase and prometaphase plates were subsequently silver-stained to display active NORs and rephotographed after counter-staining with 3% Giemsa solution in Sørensen's phosphate buffer (pH 8.0) for 5 min. Comparison between donor animals showed the same NOR pattern for all 3 animals. Forty RBG-banded metaphase and

prometaphase plates selected at random from all 3 donor animals were photographed and registered. These cells were then silver-stained, counter-stained with Giemsa and rephotographed. Metaphase and prometaphase plates with corresponding R-band and NOR-staining were compared. The locations of active NORs and NOR associations were determined.

## RESULTS

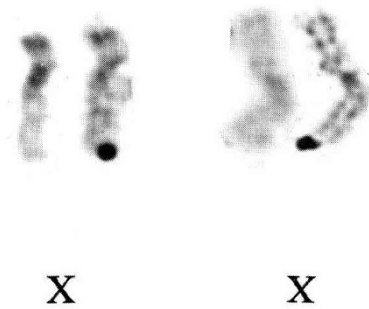
A total of 40 randomly selected metaphases were sequentially examined with RBG-banding and Ag-NOR staining. Eight autosomes, 5, 8, 14, 16, 19, 21, 32, 37, carry NORs in the telomeric regions (fig 1). The late-replicating X chromosome also showed an active telomeric NOR (Xq) in 10% of the examined metaphases (fig 2). Twenty-one metaphases displayed NOR association among the autosomes at a range of 1–3 associations per metaphase. The X chromosomes were not involved in any NOR association. As shown in table I, chromosomes 14 (38.46%) and 16 (19.23%) have remarkably high levels of involvement in NOR association. Typical NOR associations are shown in figure 3.



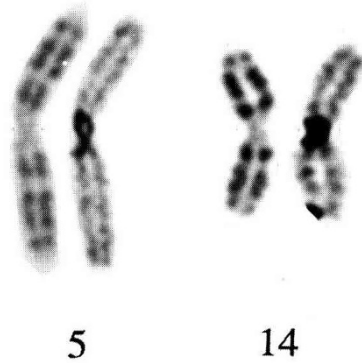
**Fig 1.** The 8 representative autosomes of the dog carrying active NORs after fluorescence plus Giemsa (FPG)-staining (left side) and subsequent Ag-NOR staining (right side).

## DISCUSSION

Several authors have described NORs in domestic dog (Kopp *et al*, 1982a, b; Pathak *et al*, 1982; Howard-Peebles and Howell, 1983). Since the dog karyotype is not standardized, the reported positions of NORs on dog chromosomes were inconsistent. Howard-Peebles and Howell (1983) published that the maximum number of NORs in the dog karyotype was 8, including the NOR on the Y chromosome. Using high-resolution RBG-banding (Poulsen *et al*, in press), NORs



**Fig 2.** The late-replicating X chromosomes of the domestic dog showing silver-positive NOR (right side) on the telomere of the q-arm.



**Fig 3.** Typical NOR associations observed in dog karyotypes (right side). When stained only with Giemsa (left side), they look like metacentric chromosomes.

**Table I.** The number and frequency of 26 identified NOR associations between autosomes.

<i>Chromosome number</i>	<i>5</i>	<i>8</i>	<i>14</i>	<i>16</i>	<i>19</i>	<i>21</i>	<i>32</i>	<i>37</i>	<i>Total</i>	<i>Frequency</i>
5	1			1				1	3	5.77
8			2		1	1	1		5	9.62
14		2	3 <sup>a</sup>	8 <sup>a</sup>		3	2	2	20	38.46
16	1		8 <sup>a</sup>			1			10	19.23
19		1				1			2	3.85
21		1	3	1	1				6	11.54
32		1	2						3	5.77
37	1		2						3	5.77
<b>Total</b>									<b>52</b>	<b>100.01</b>

<sup>a</sup> NOR association (14-14-16) was counted as 14-14, 14-16, 14-16.

on 8 different autosomes were observed (fig 1). In males, Pathak *et al* (1982) and Kopp *et al* (1982b) reported that the Y but not the X chromosome displayed active NORs. Late-replicating X chromosomes bearing silver grains were observed at a low frequency (10%) after Ag-NOR staining (fig 2). However, at this stage, further investigation of a larger population is needed to determine the presence and role of NORs on sex chromosomes in the dog.

As previously reported by Kopp *et al* (1982a), we observed a high incidence (50%) of metaphases with NOR association. NOR-associated chromosomes may look like metacentrics as shown in figure 3. Translocations in normal dog (Larsen *et al*, 1978, 1979; Mayr *et al*, 1986; Ma and Gilmore, 1971; Welling and Strandström 1988), dog cancer (Grindem and Buoen, 1986; Benjamin and Noronha, 1967; Oshimura *et al*, 1973; Else *et al*, 1982; Welling *et al*, 1988) and other abnormalities (Shive *et al*, 1965; Hare *et al*, 1967) were reported after using conventional Giemsa-stained or G-banded chromosomes. Especially in cancer studies (Mellink *et al*, 1989), metacentric chromosomes have been used as markers of neoplastic development. Without using NOR staining, however, there is a high risk of confusing NOR association with true translocation.

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