

## A fragile site in the X chromosome of the reindeer (*Rangifer tarandus* L)

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### INTRODUCTION

During an investigation on the heterochromatin in the reindeer (*Rangifer tarandus*) and some related species (*Alces alces*, *Odocoileus virginianus* and *Dama dama*) (Bianchi *et al*, 1990), bromodeoxyuridine (BrdU) was added to the lymphocyte cultures for the last 5 or 7 h of growth. In chromosome preparations from BrdU-treated cultures, a fragile site (fra) was observed in the X chromosome of the reindeer. The X chromosome of the reindeer is exceptionally large and very different from the X chromosome of other mammals. The fra site was located at the distal end of one of the arms of the nearly metacentric X chromosome. The fraX seemed to occur in a high proportion of the BrdU-treated cells. fra sites were not observed in any of the other species investigated. The accidental finding of fraX in the reindeer initiated a reexamination of slides from the reindeer cell cultures.

The chromosomes of 5 reindeer were investigated. Three subjects represented the semi-domesticated animal (*Rangifer tarandus tarandus*), two were wild forest reindeer (*Rangifer tarandus fennicus*). As has previously been shown (Gripenberg, 1984), no cytogenetic differences appear to exist between the two subspecies. BrdU-treated cultures and parallel cultures without BrdU were screened. Furthermore, a sixth reindeer (*Rangifer tarandus fennicus*) was investigated. In this case only chromosome preparations from cultures without BrdU were screened.

Lymphocytes were cultured according to standard methods. Pokeweed mitogen (PWM) was the preferred mitogen. BrdU (20 µg/ml) was added to the cultures for the last 5 or 7 h of growth. The chromosomes were G- and C-banded or stained with conventional Giemsa. Whenever possible, 100 metaphases were screened for the presence of fraX.

Figure 1 shows the G- and subsequently C-banded chromosomes of the reindeer (Gripenberg, 1984). The X chromosome is of 'the duplicate type' or comprises 9% of

the total complement, while the mammalian X usually comprises 5% of the genome (Fraccaro *et al*, 1968).

Measurements of the length of the arms of X showed Xq to be 58% of the total length of X; large blocks of heterochromatin – 25% of the chromatin of the X chromosome – were located in Xq (fig 1). The length of Xp was 42% of X; the fra site was at the distal end of Xp (fig 2). It is worth noting that without the acquisition of the heterochromatin, the relation between the arm lengths would be the opposite: the fraX arm would be the long arm, Xq.

Due to extensive structural rearrangements in the X chromosome of the reindeer, bands or segments homologous with the human X can hardly be identified on the basis of cytological characteristics.

Table I shows the frequency of fraX in cells from 6 animals treated with BrdU for 0, 5 or 7 h. In cultures without any added BrdU, cells with fraX were only occasionally observed. Addition of BrdU for the last 5 h of growth apparently enabled incorporation into a smaller proportion of the cells; fraX was observed at a frequency less than 50%. Cells incorporating BrdU for 7 h showed a fraX frequency in the range of 53–86% (mean 66%). In females, the fra site is expressed in both X chromosomes in about half of the cells. fraX is often seen in both sister chromatids.

**Table I.** Frequency of fraX cells induced by BrdU in 6 reindeer cell cultures during 0, 5 or 7 h of growth.

<i>Subject no<sup>a</sup></i>	<i>Sex</i>	<i>BrdU incorp (h)</i>	<i>No of cells screened</i>	<i>% fraX cells</i>
1	M	7	100	53
2	M	7	100	86
3	F	0	100	1
		7	100	59
4	F	5	20	45
5	F	0	70	0
		5	100	36
		7	40	67
6	F	0	100	2

<sup>a</sup> Subjects 1, 2 and 6 are wild forest reindeer, 3, 4 and 5 are semi-domesticated reindeer. M: male; F: female.

fra sites have so far been described in only a few animals (Sutherland and Ledbetter, 1988). In human cytogenetics, more than 100 fra sites are presently known and most of these sites are regarded as harmless chromosome variants. A notable exception is, however, the human fraX located at the distal end of the long arm (Xq27.3) and associated with transmissible mental retardation (Kähkönen, 1989).

The cytological similarity between the human fraX and the reindeer fraX is obvious. In reindeer, the original chromosome arm with the distal fra site could be either Xq or Xp as explained above. The human fraX is a rare folate-sensitive fragile

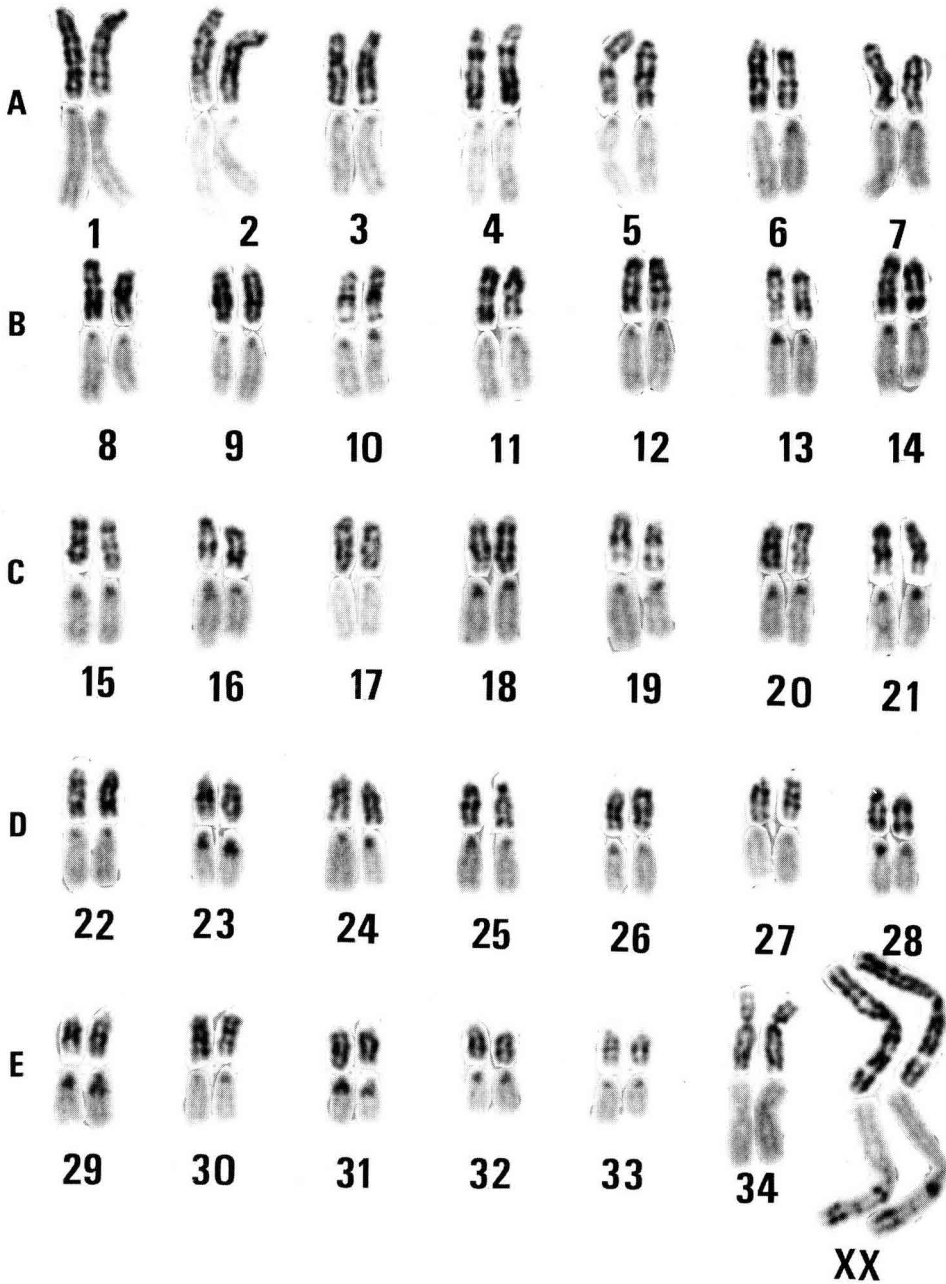


Fig 1. *Rangifer tarandus*, (70,XX). G- and subsequently C-banded chromosomes.



**Fig 2.** BrdU-induced fragile sites at the distal end of Xp. **a.** normal X chromosomes. **b.** fraX chromosomes.

site associated with mental retardation. The reindeer fraX, however, is a common BrdU-induced fra site. As far as it is known, the reindeer with fraX are normal in their appearance as well as in their behavior.

The biological significance of the fra sites is not known. It has, however, been pointed out that fra sites are cytogenetic manifestations of normal structural elements. Their significance for chromosome evolution has been discussed (Miró *et al*, 1987). fra sites can be regarded as recombinogenic hot spots causing inter- as well as intrachromosomal changes.

Whether the fra site of the reindeer X has contributed in some way to the structural changes in the chromosome arm in question is not known at present. However, the view that fraX in the reindeer is located at a site corresponding to the site of the human fraX seems possible. Homologies between human and reindeer X chromosomes remain to be further investigated.

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