# THE X CHROMOSOMES OF MAMMALS: KARYOLOGICAL HOMOLOGY AS REVEALED BY BANDING TECHNIQUES<sup>1</sup>

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#### ABSTRACT

A comparison of the Giemsa-banding patterns of the X chromosomes in various mammalian species including man indicates that two major bands (A and B), which are resistant to trypsin and urea-treatments, are always present irrespective of the gross morphology of the X chromosomes. This is true in all mammalian species with the "original or standard type" X chromosomes (5– 6% of the haploid genome) thus far analyzed. In the unusually large-sized X chromosomes the extra chromosomal material may be due either to the addition of genetically inert constitutive heterochromatin or to an X-autosome translocation. In these X chromosomes two major bands are present in the actual X-chromosome segment. Our data on C and G band patterns also support OHNO's hypothesis that the mammalian X chromosome is extremely conservative in its genetic content, in spite of its cytogenetic variability.

THE mammalian sex chromosomes vary considerably from species to species, in size as well as in morphology. The X chromosome, for example, may be the smallest or the largest element of the complement depending on the number and size of the autosomes present in a species. If many autosomes fused to form a small number of very large elements, as in the case of many species of bats, the X chromosome becomes the smallest element. On the other hand, if the autosomes are numerous and small, as in the case of the domestic *Bos* and the domestic dog, the X becomes the largest chromosome in the genome. It may be metacentric, submetacentric, or acrocentric. The Y chromosome is also extremely variable in size, ranging from a barely-discernible minute chromosome to second in size only to the X chromosome.

OHNO (1967) hypothesized that the mammalian X chromosome is extremely conservative in its genetic content, in spite of its cytological variability. This hypothesis is based on several facts: (a) many sex-linked genes, e.g., glucose-6phosphate dehydrogenase (G-6-PD), are found to be sex-linked throughout the class, whenever genetic data are available; (b) the "dosage compensation" phenomenon prevail in placental mammals so far studied; (c) the functional Xchromosome of mammals is similar in size relative to the entire complement, i.e.,

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it comprises approximately 5-6% of the haploid female genome (Ohno, Becak and Becak 1964).

The variability of the size of the X chromosome in mammals is, therefore, determined by two major factors: the number and size of the autosomes as mentioned earlier and the amount of extra chromosomal material attached to the X chromosome. Actually there are two systems whereby this ratio may change. The first, and probably the more common one, is the addition of the genetically inert constitutive heterochromatin material to the X chromosome, thus increasing the size of the X but not the genetic information. Many such examples have been described (SHARMA, PATHAK and RAY-CHAUDHURI 1970; WURSTER, SNAPPER and BENIRSCHKE 1971; PATHAK *et al.* 1973). The second category is the X-autosome translocation. When an autosome is translocated *in toto* or in part onto an X, the size of the X is naturally increased. Such cases are not common (FREDGA 1970), and usually these can be identified by the presence of the homolog of the translocated autosome which becomes an unpaired element, giving the males an odd diploid number.

The situation with the Y chromosome variation is very similar. The variation in size of the mammalian Y is due principally to the addition of constitutive heterochromatin. However, Y/autosome translocation has also been reported (MATTHEY 1966).

The chromosome banding procedures have significantly increased the resolution of cytological preparations. Comparison between the X chromosomes of related species by means of the fluorescence banding (Q bands), the Giemsa banding (G bands), and the reverse banding (R bands) revealed similar patterns, indicating a crude homology in terms of the arrangement of genetic material within the mammalian X chromosome (PEARSON *et al.* 1971; YOSIDA and SAGAI 1973; BORROW and MADAN 1973; PATHAK, HSU and ARRIGHI 1973; EVANS, BUCKLAND and SUMNER 1973; GROUCHEY *et al.* 1972; TURLEAU, GROUCHY and KLEIN 1972; LEJEUNE *et al.* 1973). However, in most of these studies, comparisons were made with a limited number of closely related species. Since we have in our laboratory a variety of frozen animal cells, we considered it worthwhile to compare the sex chromosomes of as many representative taxa as feasible by means of the G and C banding techniques. Table 1 contains a list of the mammalian species whose sex chromosomes have been studied by others using Giemsabanding.

#### MATERIALS AND METHODS

With a few exceptions in which primary cultures were used for cytological preparations, all others were recovered cultures frozen during the past decade in this laboratory. All cultures were grown in McCoy's 5a medium supplemented with 20% fetal bovine serum.

Harvest was made with a 1-hr Colcemid  $(0.05 \ \mu g/ml)$  treatment to arrest mitoses. Air-dried slides were made and the trypsin procedure (SEABRIGHT 1971) or the Urea procedure (SHIRAISHI and YOSIDA 1972) was used for G-banding. The slides were stained in 2% Giemsa mixture diluted in phosphate buffer.

In many species the cells were available from only one sex, either from a male or from a female individual. Since two X chromosomes of a female exhibit similar Giemsa-banding patterns, only one X has been reproduced in the figures.

#### RESULTS

## The X Chromosome

Table 1 lists the species of the mammals, with their taxonomic positions, whose sex chromosomes have been analyzed by G banding thus far. Although the bulk of the material came from the present investigation, the table also includes data obtained by other investigators. Figure 1 presents photomicrographs of the sex chromosomes of several representative types in respect to the X chromosome constitution. Figures 1a-f show the "standard type", (the "original type" of OHNO, BECAK and BECAK 1964), viz., the type with the X chromosome representing approximately 5–6% of the haploid genome. Figures 1g-k exemplify those containing additional heterochromatin; and Figures 11-m, those with autosomal translocation. In Figure 2, a number of diagrammatic representations are shown for comparison of the G bands patterns.

Generally, in the "standard type,,' the X chromosome exhibits two major bands and several minor bands when the preparations are treated with trypsin properly.



FIGURES 1a-m.—Giemsa-banding patterns of X chromosomes in various mammalian species. First row: "Original or standard type" metacentric, submetacentric, subtelocentric, and acrocentric X chromosomes, a. Atilax paludinosus (X), b. Ursus americanus (XY), c. Homo sapiens (XY), d. Xerus rutilus (XY), e. Sigmodon hispidus (XY), f. Clethrionomys rutilus (XY). Second row: Unusually large-sized X chromosomes where constitutive heterochromatin is added, g. Didelphis virginiana (XY), h. Microtus agrestis (X), i. Tylomys panamensis (XY), j. T. nudicaudus (XY), k. Balaenoptera acutorostrata (XY). Third row: Large-sized X chromosomes due to X-autosome translocation, l. Carollia perspicillata  $(XY_1Y_2)$ , m. Muntiacus muntjak  $(XY_1Y_2)$ .

# TABLE 1

A list of mammalian species whose sex chromosomes have been analyzed by Giemsa (G) banding

Species	References
Marsupialia	
Didelphidae	
Didelphis virginiana (Opossum)	This paper
Macropodidae	
Potorous tridactylus (Rat kangaroo)	This paper
Primates	
Tupaiidae	
Tupaia montana (Mountain tree shrew)	This paper
Urogale everetti (Mindanao tree shrew)	This paper
Cercopithecidae	
Macaca mulatta (Rhesus macaque)	STOCK and Hsu 1973
Cercopithecus aethiops (African green monkey)	Sтоск and Hsu 1973
Pongidae	
Pan troglodytes (Chimpanzee)	BOBROW and MADAN 1973; EGOZCUE, CABALLIN and GODAY 1973
Gorilla gorilla gorilla (Gorilla)	Egozcue <i>et al.</i> 1973; Sperling, Pers comm
Pongo pygmaeus (Orangutan)	Egozcue <i>et al.</i> 1973; Sperling, Pers. comm.
Homonidae	
Homo sapiens (Man)	SEABRIGHT 1972; This paper
Chiroptera	
Pteropidae	
Syconycteris australis	This paper
Phyllostomidae	
Artibeus lituratus (Big fruit-eating bat)	This paper
Carollia perspicillata (Seba's fruit bat)	PATHAK, HSU and UTAKOJI 1973
C. brevicauda	This paper
C. castanea (Allen's fruit bat)	This paper
Chiroderma villosum	This paper
Sturnira spp.	This paper
Anoura geoffroyi (Geoffroy's tailless bat)	This paper
Choeroniscus intermedius (Long-tailed bat)	This paper
Rodentia	
Heteromyidae	
Dipodomys microps (Kangaroo rat)	This paper
D. merriami	This paper
Cricetidae	
Gerbillus gerbillus (Gerbil)	This paper
Cricetulus griseus (Chinese hamster)	Kato and Yosida 1972; Какаті and Sinha 1972; This paper
Mesocricetus auratus (Syrian or "golden" hamster)	Popescu and DI PAOLO 1972
M. newtoni (Romanian hamster)	Voiculescu, Vogel and Wolf 1972
Microtus agrestis (Field vole)	Cooper and Hsu 1972; This paper
Clethrionomys rutilus (Alaskan red-backed vole)	I his paper
Peromyscus crinitus (Canyon mouse)	PATHAK, HSU and ARRIGHI 1973
<i>P. eremicus</i> (Cactus mouse) <i>Neotoma albigula</i> (White-throated wood rat)	Mascarello, Stock and Pathak 1974

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N. lepida N. micropus (Plains woodrat) N. phenax Tylomys panamensis (Climbing rat) T. nudicaudus Baiomys taylori (Pygmy mouse) Sigmodon arizonae (Arizona cotton rat) S. hispidus (Eastern cotton rat) Muridae Mus musculus (House mouse) M. poschiavinus (Tobacco mouse) Rattus norvegicus (Norway rat) R. rattus (Black rat) R. exulans R. muelleri R. fuscipes R. conatus R. sabanus Apodemus flavicollis Sciuridae Spermophilus columbianus (Ground squirrel) Xerus rutilus (African ground squirrel) Caviidae Cavia porcellus (Guinea pig) Lagomorpha Leporidae Lepus americanus (Snowshoe hare) Oryctolagus cuniculus (Domestic rabbit) Sylvilagus bachmanni (Brush rabbit) Carnivora Mustelidae Mustela putorius furo (Ferret) M. roxosa (Least weasel) Procyonidae Bassariscus astutus (Ring-tailed cat) Canidae Vulpes velox (Plains kit fox) V. macrotis (Desert kit fox) Canis familiaris (Dog) Ursidae Ursus americanus (American black bear) U. arctos (Grizzly bear) Viverridae Atilax paludinosus (Marsh mongoose) Herpestes edwardsi (Indian mongoose) Cetacea

Balaenopteridae Balaenoptera acutorostrata (Minke whale) B. physalus (Fin whale) MASCARELLO, STOCK and PATHAK 1974 MASCARELLO, WARNER and BAKER 1974 MASCARELLO, WARNER and BAKER 1974 Ратнак et al. 1973 Ратнак et al. 1973 This paper ZIMMERMAN and SIHVONEN 1973; This paper This paper SCHNEDL 1971; BUCKLAND, EVANS and SUMNER 1971; NESBITT and FRANCKE 1973; This paper **ZECH** et al. 1972 SCHNEDL and SCHNEDL 1972; UNAKUL and Hsu 1972 YOSIDA and SAGAI 1973 WOLF et al. 1972 This paper This paper This paper This paper This paper This paper FREDGA and MANDAHL 1973 This paper FREDGA and MANDAHL 1973

This paper This paper



FIGURE 2.—Diagrammatic figures of X chromosome Giemsa banding patterns in 1. Pan troglodytes, 2. Gorilla gorilla, 3. Pongo pygmaeus, 4. Cercopithecus aethiops, 5. Macaca mulatta, 6. Urogale everetti, 7. Tupaia montana, 8. Potorous tridactylus, 9. Syconycteris australis, 10. Artibeus lituratus, 11. Carollia castanea, 12. Chiroderma villosum, 13. Sturnira spp., 14. Anoura geoffroyi, 15. Choeroniscus intermedius, 16. Lepus americanus, 17. Oryctolagus cuniculus, 18. Sylvilagus bachmanni, 19. Mustela putorius, 20. M. roxosa, 21. Bassariscus astutus, 22. Vulpes velox, 23. V. macrotis, 24. Canis familiaris, 25. Herpestes edwardsi, 26. Balaenoptera physalus, 27. Tapirus terrestris, 28. Cervus elaphus, 29. Muntiacus reevesi, 30. Rangifer tarandus, 31. Bos taurus, 32. Capra hircus, 33. Ovis canadensis, 34. O. aries, 35. Ammotragus lervia, 36. Sus scrofa, 37. Gerbillus gerbillus, 38. Cricetulus griseus, 39. Mesocricetus auratus, 40. M. neutoni, 41. Peromyscus crinitus, 42. P. eremicus, 43. Neotoma albigula, 44. N. phenax, 45. Baiomys taylori, 46. Sigmodon arizonae, 47. Mus musculus, 48. M. poschiavinus, 49. Apodemus flavicollis, 50. Rattus

Species	References
Artiodactyla	
Cervidae	
Cervus elaphus (Red deer)	This paper
Muntiacus muntjak (Indian muntjac)	PATTERSON and PETRICCIANI 1973
	This paper
M. reevesi (Reeves' muntjac)	This paper
Rangifer tarandus (Caribou)	This paper
Bovidae	
Bos taurus (Cattle)	Schnedl 1972; Evans, Buckland and Sumner 1973; This paper
Capra hircus (Goat)	Evans, Buckland and Sumner 1973
Ovis canadensis (Wild sheep)	NADLER, HOFFMANN and WOLF 1973
O. aries (Domestic sheep)	EVANS, BUCKLAND and SUMNER 1973
Ammotragus lervia (Barbary sheep)	This paper
Suidae	
Sus scrofa (Wild pig)	Schnedl 1973
Perissodactyla	
Tapiridae	
Tapirus terrestris (Brazilian tapir)	This paper

TABLE 1-Continued

When the preparations are over-treated, the minor bands are no longer noticeable, but the two major bands usually persist. One of the major bands, hereafter referred to as Band A, is apparently composed of two dark sub-bands spaced closely to each other. This differentiation can be observed only in favorable cells. In most casses, these two sub-bands merge, forming a single broad band. The second band, referred to as Band B, is narrower than Band A but is always intensely stained with Giemsa.

Using the human X chromosome as a convenient example, Band A is located at the proximal portion of the long arm, straddling the zones Xq1 and Xq2, whereas Band B is located in the middle of the short arm, crossing zones Xp1 and Xp2 (PARIS CONFERENCE 1972). An additional broad band (Xq2.5) is found to locate at the distal portion of the human X chromosome. However, this particular band, like minor bands, is susceptible to trypsin overtreatment. In the SEABRIGHT paper (1972), her Figure 1 shows the major bands as well as the additional bands, but her Figure 5 shows only the two major bands.

In all mammalian species with the "standard type" X chromosomes thus far analyzed, the two major bands are present irrespective of the gross morphology of the chromosmes. However, the locations of these two major bands may vary. In acrocentric chromosomes, they are of course together in the same arm, with Band A more distal than Band B. In metacentric and submetacentric chromosomes, the two bands may be separated, one on each arm (Figure 2).

norvegicus, 51. R. rattus, 52. R. exulans, 53. R. muelleri, 54. R. fuscipes, 55. R. conatus, 56. R. sabanus, 57. Spermophilus columbianus, 58. Dipodomys microps, 59. D. merriami, 60. Cavia porcellus.

It is now very easy to demonstrate constitutive heterochromatin. The C band staining procedure (ARRIGHI and HSU 1971) can positively identify heterochromatic segments. In many species with large X chromosomes—e.g., the Chinese hamster, the Syrian hamster, Microtus agrestis, Tylomys panamensis, etc.—the long arm is invariably totally C band positive. In opossum X chromosome the heterochromatin is present in both arms (SINHA, KAKATI and PATHAK 1972). In M. agrestis and T. nudicaudus, a portion of the short arm (the proximal 1/4) is also heterochromatic. G band staining shows two distinct bands in the euchromatic segment of the giant X chromosome. A similar situation is found in other species possessing large X chromosomes. Thus, when the large amount of heterochromatin is disregarded, the euchromatic segments of these chromosomes are similar in G band patterns to those of the standard type (Figures 1a-f). Since in most of these cases heterochromatin is added, as one total chromosome arm, to the original acrocentric X, the two major G bands are invariably together in the same arm. An exceptional case is found in the Minke whale, Balaenoptera acutorostrata whose original X is a submetacentric and heterochromatin was added as an extension of the euchromatic long arm; the two major G bands are found in separate arms.

When an autosome is translocated onto an X chromosome, as in the cases of the Seba's fruit bat (*Carollia perspicillata*), and the Indian Muntjac (*Muntiacus muntjak*), the banding pattern of this translocated segment is expected to be the same as the free homolog or  $Y_1$ . This was found to be true (Figures 11–m). The real X chromosome is, therefore, represented by the short arm and the proximal segment of the long arm. The two typical G bands are both located in the short arm of the Muntjac, but are separated, one on each arm, in Carollia. FREDGA (1971) and COMINGS (1971) failed to demonstrate these two bands on the X chromosomes of the same Muntjac using the QM banding technique.

# The Y Chromosome

In C band preparations, the mammalian Y chromosome usually appears heterochromatic except when the Y is a tiny element. However, the intensity of staining of the Y chromosome varies from species to species and in some cases even from segment to segment within the same Y. The best example is probably the Caribou *Rangifer tarandus* (Figure 3). The Y chromosome contains distinct C banding pattern.

In G band preparations, longer Y chromosomes frequently show light and dark bands instead of being heavily stained throughout the lengths (Figures 1 and 2). Of particular interest is the Y of *Microtus agrestis*. COOPER and Hsu (1972) reported that in G band preparations the centromeric area, including the very short second arm, is lightly stained while the rest of the long arm is heavily stained. This staining characteristic of the Y chromosome has also been reported in several species of Tylomys (PATHAK *et al.* 1973). Whether this lightly stained area represents the functional portion of the Y remains to be confirmed. In C band preparations this tiny fragment is not always observed.



FIGURE 3.—A metaphase spread C band pattern of *Rangifer tarandus* chromosomes. Sex chromosomes (X-Y) showing segmental staining patterns.

### DISCUSSION

Although G band patterns of metaphases are not as precise as the banding patterns of the polytene chromosomes where the chromosomes are not only greatly stretched but also pair between the homologs, the G bands (or other banding techniques) are at least great improvements from conventional karvotyping. When closely related species are compared, or when a limited number of rearrangements occur within a karyotype, banding pattern analyses provide excellent information for determining homology. When phylogenetic relationships are remote and when the chromosome morphology changes a great deal as in the present report, comparing G bands for homology is unfortunately not as precise as one would like. There is no assurance that a band found in the X chromosme of one species is strictly homologous to a similar band found in the X chromosome of another species belonging to an entirely different order. Nevertheless, when a large sample of specimens is analyzed, and the similarity persists, the value of the data may be increased. The two trypsin-resistant bands of all Xchromosomes are always present, strongly suggesting that they are more or less the same. The banding patterns of the X chromosomes of various mammals add further credence to the hypothesis arrived at from genetic data that the evolution of the mammalian X has been conservative in terms of the retention of its genetic material. When the field of molecular cytogenetics advances to such a stage that the DNA composition of each band can be identified, the problem should be re-examined.

If the foregoing assumption is a reasonable one, then the basic mammalian X chromosome simply underwent a number of inversions, pericentric and para-

centric. Our data on C banding and G banding also support OHNO's contention that the basic X chromosome comprises of approximately 5–6% of the haploid mammalian genome. Larger X chromosomes contain either additional heterochromatic material or an autosomal segment translocated onto the X. Our only objection is to his terms "duplicate," "triplicate," etc., which at least imply that the genetic material is doubled, tripled, etc. Since constitutive heterochromatin is not known to carry genetic messages, addition of heterochromatin would mean only the addition of inert chromatin material. Furthermore, the addition of heterochromatin can be of various proportions, not necessarily in the exact doubling or tripling the length of the original X.

The Y chromosome may be a very small, microscopically barely perceptible body, in a number of mammals, notably in the opossum *Didelphus albiventris*, in many bats, and in many marmosets (Hsu and BENIRSCHKE 1967-1973). Since these species perpetuate without difficulty, it must be assumed that the tiny Y chromosomes of these species carry the necessary genetic information.

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