# The Mus musculus musculus type Y chromosome predominates in Asian house mice

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#### Summary

Using a mouse Y chromosomal repetitive sequence that differentiates between the Mus musculus musculus type Y chromosome and the M. m. domesticus type Y chromosome, we studied the Y chromosome in M. m. molossinus, M. m. castaneus and M. m. subspecies specimens recently trapped in Japan, Taiwan and China as well as Asian mice maintained at the Jackson Laboratory and Litton Bionetics. Here we report that the M. m. musculus type Y chromosome predominates in Asian house mice and that Japanese mice maintained at some laboratories may not represent typical M. m. molossinus.

# 1. Introduction

The isolation of mouse Y chromosomal DNA has been reported by several groups (Bishop et al. 1985; Eicher et al. 1983; Lamar & Palmer, 1984; Nallaseth & Dewey, 1986; Nallaseth et al. 1983; Nishioka & Lamothe, 1986, 1987). These DNA fragments provide powerful molecular tools with which to study the evolution of the mouse Y chromosome as some of them differentiate between the M. m. musculus and the M. m. domesticus types of Y chromosome (Bishop et al. 1985; Nishioka & Lamothe, 1986). It was found that many classical inbred strains have the M. m. musculus type Y chromosome (Bishop et al. 1985; Lamar & Palmer, 1984; Nishioka, 1987; Nishioka & Lamothe, 1986), despite the fact that the mitochondrial genome and most of the nuclear genomes of classical inbred strains originated from M. m. domesticus (Ferris et al. 1982; Moriwaki et al. 1982, 1985; Yonekawa et al. 1980, 1982). Since it is generally accepted that Japanese house mice (M. m. molossinus) contributed to the establishment of classical inbred strains (Festing & Lovell, 1981; Keeler, 1931; Potter, 1978), Bishop et al. (1985) predicted the presence of the M. m. musculus type Y chromosome in M. m. molossinus. Indeed M. m. molossinus specimens obtained from Litton Bionetics had the M. m. musculus type Y chromosome (Nishioka & Lamothe, 1986). We extended the survey to five M. m. molossinus specimens recently trapped in Japan and found the M. m. musculus type Y chromosome in all of them. In addition, M. m. castaneus trapped in Taiwan and

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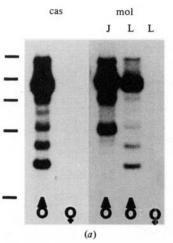
Thailand and two mice (M. m. subspecies) trapped in China also had the M. m. musculus type Y chromosome. Thus, the M. m. musculus type Y chromosome appears to be predominant in Asian house mice. Unexpectedly M. m. molossinus strains maintained at the Jackson Laboratory had the M. m. domesticus type Y chromosome.

# 2. Materials and methods

Mice: Ibred strains C57BL/6J and SWV (Staats, 1980) were obtained from the Jackson Laboratory (Bar Harbor, Maine) and Dr D. Trasler, McGill University, respectively. *M. m. molossinus* strains, MOLC/Rk, MOLD/Rk and MOLE/Rk were initially provided by Dr F. Biddle, University of Calgary. Subsequently we purchased these strains and MOLF/Ei from the Jackson Laboratory to confirm the observation. We also obtained *M. m. molossinus* specimens from Litton Bionetics (Kensington, Maryland) through Dr M. Potter of the National Cancer Institute. The localities of other Asian mice obtained from Dr Moriwaki, National Institute of Genetics, Japan are indicated in the figure legends.

Filter hybridization: DNAs isolated from liver were digested with Eco RI (Boehringer Mannheim Canada, Dorval, Quebec) and electrophoresed on a 0.8%agarose gel in a buffer consisting of 40 mM Tris (pH 8.0), 80 mM sodium acetate and 1 mM EDTA. DNA fragments were transferred to membrane filters (Gene Screen, New England Nuclear Canada, Lachine, Quebec) by the method of Southern (1975). Filters were pretreated with 50 µg/ml of denatured herring

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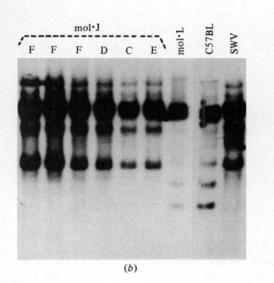


Fig. 1. Southern blot analysis of Asian mice maintained at two laboratories. About  $10 \mu g$  of DNAs were digested with Eco RI and electrophoresed on a 0.8% agarose gel. DNA fragments were transferred to membrane filters and hybridized to nick-translated AC11.

(a) cas: M. m. castaneus (Thailand) obtained from Litton Bionetics. mol. J: M. m. molossinus (MOLF/Ei) obtained from the Jackson Laboratory. mol. L: M. m. molossinus obtained from Litton Bionetics (sample number 19717). Solid lines indicate size markers from top to bottom 23 kb, 9.6 kb, 6.8 kb, 4.3 kb and 2.3 kb.

(b) mol-J: M. m. molossinus (C, D, E and F stand for strains MOLC/Rk, MOLD/Rk, MOLE/Rk and MOLF/ Ei, respectively). mol-L: M. m. molossinus (Litton Bionetics, sample number 19717).

C57BL/6J and SWV are used as controls for the *M. m. musculus* type Y chromosome and the *M. m. domesticus* type Y chromosome, respectively.

sperm DNA in a solution containing  $5 \times SSC$ (1 × SSC is 0.15 N-NaCl and 0.015 N-Na-citrate), 50% formamide, 1 × Denhardt's solution (Denhardt, 1966) and 0.1% SDS for 12–16 h at 41 °C. Filters were then hybridized to a [<sup>32</sup>P]-labelled Y chromosomal DNA

## 3. Results and discussion

AC11 is a mouse Y chromosomal repetitive sequence containing a Bkm-related sequence (Nishioka & Lamothe, 1986). In this study a portion of the clone deprived of the Bkm-related sequence was used as the molecular probe. When hybridized to male and female DNAs isolated from M. m. castaneus and M. m. molossinus, male lanes showed significant accumulation of AC11-related sequences (Fig. 1a), as previously observed in DNAs of M. m. musculus and M. m. domesticus (Nishioka & Lamothe, 1986), and M. m. castaneus had the M. m. musculus type Y chromosome. Since M. m. molossinus is an intersubspecific hybrid between M. m. musculus and M. m. castaneus (Yonekawa et al. 1986), the presence of the M. m. musculus type Y chromosome in M. m. molossinus was inferred. Unexpectedly, both types of the Y chromosome were found among M. m. molossinus specimens. Four strains, MOLC/Rk, MOLD/Rk, MOLE/Rk and MOLF/Ei, from the Jackson Laboratory had the M. m. domesticus type Y chromosome, whereas two specimens from Litton Bionetics had the M. m. musculus type Y chromosome (Fig. 1 and 2). We then extended our survey to 5 M. m. molossinus specimens recently trapped in Japan, 1 M. m. castaneus from Taiwan and 2 M. m. subspecies from China and found the M. m. musculus types Y chromosome in all of them (Fig. 2).

The Japanese mice (M. m. molossinus) maintained at Litton Bionetics were collected by Dr Hamajima of Kyushu University and sent to Dr M. Potter in 1967 from which 4 strains (MOLC/Rk, MOLD/Rk, MOLE/Rk and MOLF/Ei) were developed at the Jackson Laboratory using a single male as the founder (M. Potter, 1978, personal communications from M. Potter and M. Davisson). In view of the fact that M. m. musculus and M. m. molossinus are genetically closer to one another than either one to M. m. domesticus (Bonhomme et al. 1984; Ferris et al. 1983; Moriwaki et al. 1982; Sage, 1981; Yonekawa et al. 1986), the presence of the M. m. domesticus type Y chromosome in M. m. molossinus background should be considered exceptional.

House mice are commensal. They can spread through commerical traffic and introduce new genetic material into aboriginal populations. For example, a specimen with Robertsonian translocation, caught on Ogasawara Island, Japan was shown to be a hybrid between M. m. domesticus and M. m. molossinus (Moriwaki et al. 1984) and M. m. domesticus type mitochondrial DNA was found in a specimen caught in Kyushu, Japan (H. Yonekawa, personal communi-

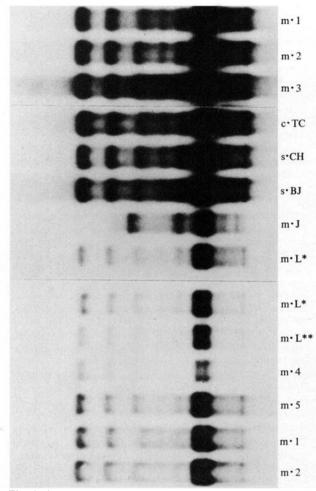


Fig. 2. Southern blot analysis of Asian wild mice. DNAs digested with Eco RI were electrophoresed on a 0.8% agarose gel, transferred to a membrane filter and hybridized to nick-translated AC11.  $m \cdot J :: M. m.$  molossinus strain MOLF/Ei from the Jackson Laboratory.  $m \cdot L :: M. m.$  molossinus from Litton Bionetics. \* Sample number 19717. \*\* Sample number 20597. The localities of other Asian mice are as follows:  $m \cdot 1$  (M. m. molossinus), Izumi, Japan;  $m \cdot 2$  (M. m. molossinus), Mishima, Japan;  $m \cdot 3$  (M. m. molossinus), Anjo, Japan;  $m \cdot 4$  (M. m. molossinus), Nagoya, Japan;  $m \cdot 5$  (M. m. molossinus), Chiba, Japan;  $c \cdot TC$  (M. m. castaneus), Taichung, Taiwan;  $s \cdot CH$  (M. m. subspecies), Changchun, China;  $s \cdot BJ$  (M. m. subspecies), Beijing, China.

cation). SF/CamEi was established from mice caught in a coal mine in California (Staats, 1980), but has the *M. m. musculus* type *Y* chromosome which was probably introduced from an Asian mouse brought to California (Nishioka, 1987). Therefore, a likely origin of the *M. m. domesticus* type *Y* chromosome in the *M. m. molossinus* strains is a *M. m. domesticus* male brought to Kyushu, although the possibility is not completely excluded that the introduction might have occurred in the U.S.A. In this context it is relevant to point out that SK/CamEi which was established from mice caught on Skokholm island, off Pembrokeshire (Staats, 1980) has probably been mixed with SF/Cam as originally suggested by Ferris *et al.* (1982). Subsequently the *M. m. musculus* type Y chromosome

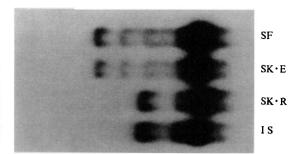


Fig. 3. Southern blot analysis of inbred strains. DNAs digested with Eco RI were electrophoresed on a 0.8% agarose gel, transferred to a membrane filter and hybridized to nick-translated AC11. SF; SF/CamEi; SK-E; SK/CamEI; SK-R; SK/CamRk; IS; IS/CamEi.

was found in SK/CamEi, despite its geographical origin (Nishioka, 1987). The presence of the M.m. domesticus type Y chromosome in another SK/Cam strain (SK/CamRk) further supports the above possibility (Fig. 3). In addition, Fig. 3 shows the presence of the M.m. domesticus type Y chromosome in IS/CamEi which was established from a cross between M.m. praetextus (M.m. domesticus) caught in Israel and a M.m. musculus composite stock (Staats, 1980).

This report is an extension of our previous survey that classified 39 inbred strains into two categories; strains with the M. m. musculus type Y chromosome and those with the M. m. domesticus type Ychromosome (Nishioka, 1987). It is clear that the M. m. musculus type Y chromosome predominates in Asia, while in America and Western Europe the representative Y chromosome is M. m. domesticus type. However, because of their commensalism with human beings, foreign house mice are introduced into certain local populations at an apparently high frequency. Our study demonstrated that DNA probes such as AC11 are useful molecular tools not only for population studies but also for confirmation of the origins of established inbred strains.

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