# Ecology of plague in Africa: response of indigenous wild rodents to experimental plague infection

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The Mastomys natalensis species complex, subdivided into genetically distinct species having diploid chromosome numbers 2n = 32 and 2n = 36, is a reservoir for several zoonoses including Lassa fever and plague. This report describes a study to determine whether these sibling species and three other rodent species have different potential as reservoirs for plague. It was found that M. natalensis (2n = 32) was significantly more resistant to experimental plague infection (50% survived inoculation with 120 000 Yersinia pseudotuberculosis subsp. pestis) than was M. coucha (2n = 36) (none of which survived doses of 190 Y. pseudotuberculosis subsp. pestis). In descending order of resistance were M. natalensis, Aethomys chrysophilus, M. coucha, Tatera leucogaster and A. namaquensis. No A. namaquensis survived inoculation of 10 or more plague bacilli.

Previous reports on susceptibility to plague or other infections, which were based exclusively on findings in the universally distributed laboratory-bred Mastomys, are thus not necessarily applicable to the M. natalensis species as a whole but probably only to M. coucha. The Y. pseudotuberculosis subsp. pestis fraction-1 passive haemagglutination test appeared to be relatively insensitive in that only 5 out of 47 animals surviving experimental plague infection showed specific antibodies 6 weeks after challenge.

The geographic distribution of human plague in southern Africa corresponds closely with that of the plague-susceptible species, M. coucha, while the resistant species, M. natalensis, predominates in areas where human plague has not been recorded. The role of A. namaquensis in the ecology of plague needs to be carefully studied and its possible importance in plague research should be investigated further.

The Mastomys natalensis species complex, one of Africa's most prevalent wild rodents, plays an important role in the natural cycle of several zoonoses, including Lassa fever and plague.

Although Lassa fever is known to occur only in West Africa, the rodent host is widely distributed over most of the continent. Monath (1) suggested several possible biological explanations for this discrepancy, including differences between subpopulations of Mastomys in their susceptibility to the virus. With regard to plague, much of the experimental work on Mastomys was done before it was appreciated that different responses by sibling species to Yersinia pseudotuberculosis subsp. pestis and other pathogenic agents might be significant. The colonies of this rodent, maintained in many laboratories around the world, are descended from the original colony still maintained at the South African Institute for Medical

Research, i.e., the 2n = 36 species (2). In South

Hallett (5), in 1970, found high Y. pseudotuberculosis subsp. pestis fraction 1 antibody titres in Mastomys sera and stated that this was unexpected but that similar results reported from Kenya might indicate that Mastomys had acquired genetic resistance in certain hyperenzootic areas. However, in 1953, Davis (6) had remarked on the occurrence of a limited plague focus in Morocco, where, in contrast to other African plague foci, Mastomys appeared not to be involved in the plague cycle.

The suggestion by Matthey (7) in 1966 that *M. natalensis* consisted of at least two genetically different species was unfortunately ignored until 1977

Africa, this animal became the preferred routine laboratory animal for the study of plague because of its consistently high susceptibility to very low doses of Y. pseudotuberculosis subsp. pestis. In 1968, Davis et al. (3) stated that "in South Africa Mastomys is highly susceptible to plague and resistance to P. pestis has never been demonstrated". In contrast, other workers (4) concluded, on the basis of epidemiological as well as laboratory susceptibility studies, that "Arvicanthis and Mastomys from Rongai in Kenya were mostly highly resistant to P. pestis".

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when Lyons et al. (8) reinvestigated the subject. Taylor et al. (9) drew attention to the apparent discrepancy between the occurrence of human plague in Zimbabwe and the distribution of plague antibodies in dog sera, and they related this to the rodent distribution. They noted the close relationship between the areas of human plague and the distribution of Mastomys (2n = 36) (Fig. 1), as mapped by Green et al. (10). The latter also suggested the renaming of members of the M. natalensis species complex as M. natalensis (2n = 32) and M. coucha (2n = 36). Other species may be added as the taxonomy becomes clearer (9).

In view of these and other apparently contradictory observations on an animal that has become established as a common laboratory model for a wide range of studies, it seemed appropriate to compare the susceptibility to a major natural pathogen, Y. pseudotuberculosis subsp. pestis, of the two sibling species of Mastomys and of other major rodent species in southern Africa.

To avoid confusion, reference to reports of work carried out on *Mastomys* species before the publication by Green et al. (10) will not include the species name. *Mastomys* used in this study are identified by their diploid chromosome numbers. *Aethomys chrysophilus* is also known to be a species complex in Zimbabwe with chromosome diploid numbers of 44 and 50 (11), but only the latter (2n = 50) has been recorded in animals from the capture site used in this study (D. H. Gordon, personal communication, 1982).

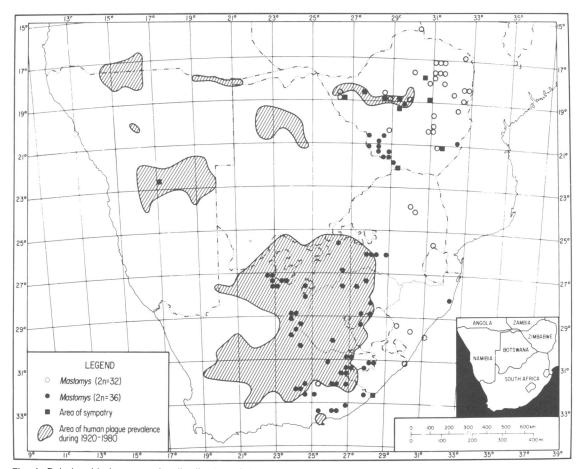


Fig. 1. Relationship between the distribution of the two chromosomal species of *Mastomys* and the prevalence of human plague in southern Africa from 1920 to 1980 (data partly from Green et al. (10) and Taylor et al. (9)). Note that karyotyping of *Mastomys* has been carried out on those from South Africa and Zimbabwe, but not yet on those from Botswana and Namibia.

### MATERIALS AND METHODS

### Animals

All rodents were trapped live in Zimbabwe in an area where plague has not been known to occur (17°55' S, 30°45' E) and Mastomys were typed according to their electrophoretic haemoglobin pattern, which is quite distinct for each of the chromosomal types (12). Serum was obtained from each animal to establish its plague antibody status prior to challenge. The animals were dusted with insecticide, kept in quarantine for some time, and then forwarded to South Africa for the challenge studies. Animals in the first batch were held in individual cages, but those in subsequent batches were kept in pairs, not necessarily of different sex. A further quarantine period was observed during which the animals were again dusted with insecticide. They were fed with standard mouse cubes and given water ad libitum. After inoculation with Y. pseudotuberculosis subsp. pestis, the animals were held in a Vickers' flexible film, negative-pressure animal containment isolator, fitted with air inlet and exhaust through HEPA filters.

The species used were: Mastomys (2n = 32) (the multimammate mouse), Mastomys (2n = 36), Tatera leucogaster (the bushveld gerbil), A. chrysophilus (the African or red veld rat), and A. namaquensis (the golden or Namaqua rock rat). Ten animals of each species were used per challenge dose. They were observed twice daily and autopsies were performed on the animals that died, when tissue was taken from heart, lung, spleen, and liver for culture. Survivors were sacrificed and exsanguinated 6 weeks after challenge for determination of plague antibody.

# Challenge strain

Y. pseudotuberculosis subsp. pestis strain SAIMR/F329/68, isolated in 1968 from a flea during a bubopneumonic plague epidemic in Lesotho, was used. It is characterized by the presence of a relatively stable pigment factor. Prior to challenge, the isolate was grown in brain-heart infusion broth and passaged through white mice and through laboratory-bred Mastomys (2n = 36). Dilutions were made in peptone water and challenge was by subcutaneous inoculation into a hind leg of 0.2 ml of the designated dilution. Control animals were inoculated with diluent only.

Viable plate counts were done immediately before and after each batch of inoculations. For the counts, 1-ml aliquots of the  $10^{-5}$  and  $10^{-6}$  dilutions were divided among five blood agar plates per dilution. These were incubated at 28 °C for 48 h, after which the colonies were counted and the mean doses received by the animals were calculated.

# Serology

The presence of antibodies against the Y. pseudo-tuberculosis subsp. pestis fraction 1B antigen was determined by means of the passive haemagglutination test using a microtitration method (13). Pre-and post-challenge sera of survivors were tested in parallel.

### RESULTS

It was noted that *Mastomys* (2n = 32) was considerably more aggressive and excitable than *Mastomys* (2n = 36).

The results of the study are summarized in Table 1, from which it may be concluded that *Mastomys* (2n = 32) is highly resistant to experimental plague infection, in that 50% of animals survived challenge with 120 000 *Y. pseudotuberculosis* subsp. *pestis*. On the other hand, *Mastomys* (2n = 36) was found to be highly susceptible, no animals having survived a dose as low as 190 bacilli. With increasing dose, the interval between inoculation and death progressively decreased.

A. namaquensis was extremely plague-sensitive, a dose of 10 bacilli having killed all inoculated animals. It also showed the shortest interval between inoculation and death, the median value of 3.0 days with a dose of 1900 bacilli being equalled only by Mastomys (2n = 36) when exposed to a dose of 120 000 bacilli.

The Y. pseudotuberculosis subsp. pestis fraction 1 haemagglutination (PHA) tests, which were done on 155 adult animals used in the study, yielded negative results prior to plague infection. Of 47 survivors, only 5, i.e., 3 Mastomys (2n = 32) and 2 A. chrysophilus, had specific antibodies six weeks after infection. Titres ranged from 1:4 to 1:16 in four animals with one, A. chrysophilus, having a titre of 1:128; the animals had been given between 1900 and 120 000 bacilli. Ten suckling infants born to three A. chrysophilus females prior to inoculation were left with their parents after the latter were infected. In each family one parent died (two females and one male). The infants, who learned to fend for themselves, all survived and did not develop clinically apparent illness or demonstrable plague antibodies.

### DISCUSSION

The role of *Mastomys* as a natural disease host has probably been most intensively investigated in relation to the epidemiology of bubonic plague. The simple technique used to distinguish *Mastomys* with chromosome numbers 2n = 32 and 2n = 36 made it

A. chrysophilus

A. namaquensis

	No. of bacilli inoculated								
Mastomys (2n = 32)	10		1.9 × 10 <sup>2</sup>		1.9 × 10 <sup>3</sup>		1.9×10⁴	1.2 × 10 <sup>5</sup>	
					2/10	(8.5)	0/10	5/10	(4)
Mastomys (2n = 36)	1/10	(6)	10/10	(7.5)	9/9	(4)		10/10	(3)
T. leucogaster	7/10	(7)	10/10	(6)	8/8	(4.5)			

Table 1. Results of subcutaneous inoculation of virulent *Y. pseudotuberculosis* subsp. *pestis* isolate SAIMR/F329/68 in 155 African wild-trapped rodents<sup>a</sup>

10/10 (4)

4/9

(3)

possible to capture and type the relatively large number of animals needed to carry out a plague susceptibility study. Although colonies of the two species are being successfully bred in our laboratory, wild-trapped rodents were used in preference to laboratory-bred animals for this preliminary study, to ensure that host factors would match closely those found in their natural environment.

10/10

(4.5)

In our study, we have shown that *Mastomys* (2n = 32) and *Mastomys* (2n = 36) differ significantly in their susceptibility to *Y. pseudotuberculosis* subsp. *pestis* in that the former is highly resistant whereas the latter is very sensitive (Table 1). *T. leucogaster*, the bushveld gerbil, which has long been believed to play an important role in southern African plague epidemiology, seemed rather more sensitive than *Mastomys* (2n = 36). *A. namaquensis* was the most sensitive of all while the response of *A. chrysophilus* was similar to that of *Mastomys* (2n = 32).

Both Aethomys and Mastomys are essentially arboreal rats, with tendencies to domesticity, and their role in plague is believed to be that of bridging the gap between the sylvatic reservoir (gerbils) and man (14). Our results, however, indicate that A. chrysophilus and A. namaquensis differ markedly in their susceptibility to plague and therefore may play different roles in the plague cycle.

It was also shown that an inverse correlation exists in the animals between degree of exposure and duration of survival after exposure. This probably reflects a dose-dependent incubation period.

Suckling infants of females with fatal plague infections did not become clinically ill, neither did a group of healthy control animals kept in the isolator throughout the study. In the absence of ectoparasites therefore, even intimate contact, as occurs between female animals and their suckling offspring, did not result in plague transmission. None of the animals had demonstrable haemagglutinating plague anti-

bodies prior to challenge and only 5 out of 47 survivors had demonstrable antibodies six weeks after challenge. In our experience, field studies of rodent serology appear to be rather fruitless as only a small percentage of animals can be shown to have antibodies (9). Our current results raise two possibilities: either the Y. pseudotuberculosis subsp. pestis fraction 1 passive haemagglutination test is not highly sensitive, or surviving animals eliminate bacilli by local defence mechanisms at the site of inoculation and a humoral antibody response does not occur.

5/10 (7)

8/10

An important result of this study is the demonstration of entirely different responses by Mastomys (2n = 32) and Mastomys (2n = 36) to a natural pathogen. These findings have important implications with regard to Mastomys as an experimental laboratory animal in various research fields (15). Especially in the African context, these results apply to *Mastomys* in its role as an ecological link in many zoonoses, such as plague (Fig. 1), Lassa fever, salmonelloses, African tickbite fever, and arbovirus infections. In this respect, a great deal of work needs to be done to redefine the relative roles of the two Mastomys species, not only with regard to their susceptibility to pathogenic microorganisms but also in relation to their behaviour, ectoparasite infestation, etc. The role of *Mastomys* as an experimental laboratory animal needs to be very carefully defined, as it is M. coucha (2n = 36) and not necessarily M. natalensis (2n = 32) which spontaneously develops adenocarcinoma of the stomach, and which has been found to be an ideal experimental host for organisms causing schistosomiasis and filariasis, and for many other pathogens.

Although most *Mastomys* colonies in laboratories throughout the world are probably derived from the original South African colony, which had the 2n = 36 chromosome number, a strong plea is made that laboratories should determine the diploid chromo-

 $<sup>^</sup>a$  Five animals in the  $1.9 \times 10^3$  dose group were killed by their cage-mates shortly after inoculation and are therefore excluded from the study. Results are given as number of deaths from plague/number of animals inoculated. Figures in parentheses give the median survival time (in days) of the animals that died.

some number of their colonies and identify the animals accordingly when publishing research findings. Finally, the epidemiological implications of these findings are probably applicable to medically important animal reservoir hosts in other parts of the world, indicating a need for reevaluation of old and current data and methodology.

# **RÉSUMÉ**

# L'ÉCOLOGIE DE LA PESTE EN AFRIQUE :

RÉACTION DE RONGEURS SAUVAGES INDIGÈNES À UNE INFECTION PESTEUSE EXPÉRIMENTALE

Une étude a été entreprise en vue de déterminer la sensibilité à l'infection pesteuse de plusieurs espèces indigènes de rongeurs sauvages dont on pense qu'elles jouent un rôle dans l'écologie de la peste en Afrique. Il s'agissait de Tatera leucogaster, d'Aethomys chrysophilus, d'Aethomys namaquensis ainsi que de membres du complexe d'espèces Mastomys natalensis. Les espèces du genre Aethomys sont essentiellement constituées de rats arboricoles à tendance domestique qui peuvent jouer un rôle analogue à celui des espèces du genre Mastomys en assurant la jonction entre les réservoirs de peste «sylvatique» (gerbilles et autres rongeurs sauvages) et l'homme. Les précédentes études ne prenaient pas en considération l'existence d'espèces jumelles de Mastomys morphologiquement analogues mais génétiquement distinctes, avec un nombre diploïde égal à 32 et à 36. espèces qui ont été respectivement rabaptisées Mastomys natalensis et Mastomys coucha. Il est établi que la colonie parente sud-africaine, utilisée dans le passé et dont des descendants ont été distribués à de nombreux laboratoires du monde entier, est constituée de M. coucha. Cette espèce s'est révélée si fortement et si régulièrement sensible à l'infection par Yersinia pseudotuberculosis subsp. pestis qu'elle est devenue l'animal de laboratoire classique tant pour le

diagnostic de la peste que pour la recherche.

La présente étude a permis d'obtenir deux types de données nouvelles. En premier lieu, on a constaté que A. namaquensis était nettement plus sensible à Y. pseudotuberculosis subsp. pestis que M. coucha et, en second lieu, on a observé que les espèces jumelles du complexe Mastomys sont nettement différentes quant à leur réaction vis-à-vis de Y. pseudotuberculosis subsp. pestis. M. coucha comme on pouvait s'y attendre, s'est révélé très sensible, en revanche M. natalensis était relativement résistant. Parmi les autres résultats obtenus, figure le fait que cette sensibilité accrue se caractérise par un laps de temps plus court entre l'exposition à une dose l'étale et la mort et par un taux de mortalité plus élevé. Aethomys chrysophilus forme également un complexe d'espèces, toutefois seules les espèces ayant un nombre diploïde de chromosomes 2n = 50 figuraient dans cette étude. Les variations aussi observées dans la réponse à un germe pathogène naturel montrent qu'il est nécessaire de revoir l'écologie d'un certain nombre de zoonoses dans les régions du monde où des complexes d'espèces leur servent de réservoirs.

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