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Title

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Journal

Proceedings of the Vertebrate Pest Conference, 16(16)

ISSN

0507-6773

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Publication Date

1994

THE ROLE OF PREDATORS IN THE ECOLOGY, EPIDEMIOLOGY, AND SURVEILLANCE OF PLAGUE IN THE UNITED STATES

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ABSTRACT: Predators play important roles in the ecology, epidemiology, and surveillance of plague in the United States. Most predators are accidental hosts of plague and, with the possible exception of grasshopper mice (Onychomys spp.), are not important sources of infection for feeding fleas. However, predators undoubtedly do play an important role in the natural cycle of plague by transporting infected fleas between different populations of plague-susceptible rodents. Predators are known to be at least accidental hosts for 40 of the 50 flea species that have been found to be naturally infected with plague in the U.S. Carnivores, including domestic cats, also play an important epidemiological role and have been sources of infection for 24 human plague cases since 1970. Serosurveillance of rodent-consuming carnivores is currently the most cost-effective method of monitoring plague in the western U.S. During the 1990s, these surveys have allowed CDC and other public health agencies to both identify plague risks for humans living in endemic regions and document the spread of plague into areas where it had not been identified previously.

Proc. 16th Vertebr. Pest Conf. (W.S. Halverson & A.C. Crabb, Eds.) Published at Univ. of Calif., Davis. 1994.

INTRODUCTION

Yersinia pestis, the causative agent of plague, is maintained in nature as a flea-borne disease of rodents. Plague probably was first introduced into the United States around 1900 by rat-infested ships entering the port of San Francisco, California (Barnes 1982). Epizootics occurred in local rat populations as the disease spread from the port area to other regions of the city. These epizootics were the source of infection for 121 human cases between 1900-1904 (Link 1955). Within a few years after its introduction, plague passed from San Francisco's urban rat populations into the wild (sylvatic) rodents of the surrounding countryside. Once plague became established in rodents other than Rattus spp., it spread rapidly across the western U.S. and by 1940 had been identified as far east as the western edge of the Great Plains (Eskey and Haas 1940).

Plague continues to exist in scattered foci throughout much of the western U.S., although more than 90 percent of the human cases occur in New Mexico, Arizona, Colorado, and California (Figure 1). The ecological relationships of plague in these established foci are extremely complex and each regional focus typically has its own characteristic flea vectors and highly susceptible mammalian rodent hosts (epizootic hosts) (Pollitzer and Meyer 1961, Barnes 1982, and CDC unpublished data). Epizootics usually occur every few years in major foci and often cause mortality in excess of 80 to 99% among different epizootic hosts (Barnes 1982). Many of these epizootic hosts are associated with important flea vectors and form regional host/flea epizootic complexes. These complexes are important for regional amplification of plague and increase the chances that Y. pestis will spread to other rodent species or accidental hosts, such as humans, lagomorphs, or carnivores. The most important epizootic hosts (and their fleas) are discussed in Barnes (1982) and include various ground squirrels (Spermophilus spp.), antelope ground squirrels (Ammospermophilus leucurus), prairie dogs (Cynomys spp.) chipmunks Geographic Distribution of Human and Animal Plague in the Western United States by County of Origin, 1970-1993

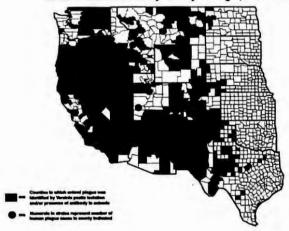


Figure 1. Plague surveillance map showing the distribution of human cases and counties where plague-positive mammals or flea pools have been identified.

(Tamias spp. and Eutamias spp.), and woodrats (Neotoma spp.). Other rodent species, such as Peromyscus maniculatus (deer mice) and Microtus californicus (California voles), have populations that differ in their resistance to plague-induced mortality. Rodents from resistant deer mouse or vole populations usually survive infection, but become bacteremic and can serve as sources of infection for feeding fleas. These animals are referred to as maintenance, or enzootic, hosts and are believed, by some, to be critical for maintaining plague during

interepizootic periods (Quan and Kartman 1962, Poland and Barnes 1979).

ROLE OF PREDATORS IN THE ECOLOGY OF PLAGUE

Mammalian predators can become infected with Y. pestis after consuming infected prey or being bitten by infected rodent fleas. With the notable exception of grasshopper mice (Onychomys spp.), which are discussed separately below, virtually all of these predators are members of the mammalian order Carnivora. Most North American carnivores, including canids, mustelids, procyonids, and ursids, apparently survive plague infections, as indicated by either experimental studies or the large number of seropositive animals found in plagueendemic areas (Rust et al. 1971a, Barnes 1982). Mortality among felids, however, is high and there are numerous records of fatal plague infection in domestic cats and bobcats (Tabor and Thomas 1986, Eidsen et al. 1988, 1991, Gasper et al. 1992, CDC unpublished records).

Although carnivores are susceptible to plague infection, they are only accidental hosts and probably play little or no role as sources of infection for feeding fleas. They do, however, play an important ecological role by transporting plague-infected fleas from one area to another (Poland and Barnes 1979). Carnivores can become infested with plague-infected fleas while consuming prey or exploring rodent nests or burrows. Fleas that have lost their normal rodent hosts during a plague epizootic are especially likely to attack a passing carnivore. Some fleas, such as the prairie dog flea Oropsylla hirsuta, actually congregate near burrow entrances after their hosts die and wait for new hosts to appear. When the next host is a carnivore, such as a coyote, the flea can be transported many kilometers before leaving its temporary host. A brief review of records from different treatises on North American Siphonaptera and unpublished CDC records indicates that a wide diversity of fleas at least accidentally infest predators, including a number of rodent fleas that are important plague vectors (Tables 1 and 2) (Hubbard 1947, Hopkins and Rothschild 1953, 1962, 1966, and 1971, Stark 1958, Campos 1971, Wittrock and Wilson 1974, Traub et al. 1983, Holland 1985). Each of the flea species listed in these tables has been reported to be naturally infected with plague (Pollitzer 1961 and CDC unpublished records). It should be noted that 40 of the 50 North American flea species reported to be naturally infected with plague are represented in these tables. Although some of these fleas, such as Pulex irritans or Echidnophaga gallinacea, are poor plague vectors, other species listed in these tables are extremely important, including many of the Oropsylla and Thrassis species.

Members of the genus Onychomys, especially the northern grasshopper mouse, O. leucogaster, play a unique and potentially important role in the interepizootic maintenance and dissemination of plague in North America. With the exception of the Pacific Coast region, the range of the northern grasshopper mouse overlaps much of the known distribution of plague in the United States. Grasshopper mice are unique among North American cricetine rodents in that they are true omnivores (Flake 1973, Landry 1970). In addition to a diet of seeds

and insect material, these mice will kill and consume other mice. This behavior exposes them to potentially infected prey and Thomas et al. (1989) demonstrated that 35% of O. leucogaster consuming experimentally infected mice became infected. Grasshopper mouse populations have been shown to vary in their susceptibility to plague, with populations from plague endemic areas being highly resistant to plague-induced mortality (Holdenried and Ouan 1956, Marchette et al. 1962 and Thomas et al. 1988a). Thomas et al. (1988a) found that first generation mice, reared from O. leucogaster parents collected in an area of Colorado where a plague epizootic had occurred recently, were nearly 12,500-fold more resistant to plague-induced mortality than similar offspring from Oklahoma mice collected in a non-endemic area. These data strongly suggest that selection among grasshopper mouse populations during plague epizootics results in a high degree of resistance.

Another important characteristic of grasshopper mice is their ability to serve as temporary or permanent hosts for a great diversity of fleas. Besides becoming infested with fleas while consuming rodent prey, grasshopper mice are likely to be exposed to fleas of other rodents while invading their burrows. It also has been hypothesized that grasshopper mice occupy the burrows of other rodents rather than digging their own (Bailey 1931, Thomas et al. 1988a). In reviewing published flea collection records from O. leucogaster, Thomas (1988b) documented collections representing 57 nominate species of fleas from these mice. Of these, 27 are known to be naturally infected with plague (Table 3). Pleochaetis exilis (formerly Monopsyllus exilis) is rarely found on hosts other than grasshopper mice, most of the fleas collected from O. leucogaster are normally associated with other rodents such as Dipodomys spp., Spermophilus spp., Peromyscus spp. and geomyids. The collections cited above indicate that there is a significant degree of flea transfer between grasshopper mice and other sympatric rodents. Given that a male O. torridus may utilize a home range as large as 3.71 ha during the breeding season (Frank and Heske 1992), it is likely that grasshopper mice play a role in disseminating plagueinfected fleas among rodent populations.

Raptors also have adequate opportunities to become infected with plague, but previous studies indicate that birds are resistant to infection and it is doubtful that predatory birds play any role as sources of infection for feeding fleas (Pollitzer 1954). Raptors, however, have been reported to be infested with potential plague vectors and could transport infected fleas from one susceptible rodent population to another (Hubbard 1947 and Holland 1985). The importance of raptors as transport hosts for plague-infected fleas has received little attention and should be investigated.

ROLE OF PREDATORS IN THE EPIDEMIOLOGY OF PLAGUE

Three hundred and nineteen cases of human plague have occurred in the United States since 1970 (CDC unpublished records). The probably source of infection was determined for 210 of these 319 cases. Most (77.1%) of these 210 cases were associated with various sciurid species and their fleas (Figure 2). Twenty-three

Table 1. Potential plague vectors infesting Felidae and Canidae. Compiled from a list of 50 potential flea vectors from the U.S. that have been found to be naturally infected with plague (Pollitzer 1961 and CDC unpublished data).

FELIDAE

Felis (domestic cats): Ctenocephalides felis, Echidnophaga gallinacaea, Euhoplopsyllus glacialis, Foxella ignota,

Oropsylla bruneri, Pulex irritans

Lynx: Cediopsylla inaequalis, Ctenocephalides felis, Echidnophaga gallinacaea, Euhoplopsyllus

glacialis, Foxella ignota, Pulex irritans, Thrassis petiolatus

CANIDAE

Canis (domestic dogs): Cediopsylla inaequalis, Ceratophyllus ciliatus, Ctenocephalides canis, C. felis,

Euhoplopsyllus glacialis, Echidnophaga gallinacaea, Hoplopsyllus anomalus, Nosopsyllus

fasciatus, Oropsylla bruneri, O. montana, Pulex irritans, P. simulans

Canis (wild Canis): Ctenocephalides canis, Echidnophaga gallinacaea, Euhoplopsyllus glacialis, Eumolpianus

eumolpi, Hystrichopsylla dippei, Megabothris abantis, Oropsylla hirsuta, O. idahoensis, O.

labis, O. rupestris, Pulex irritans, Pulex simulans, Thrassis acamantis

<u>Urocyon</u>: Ctenocephalides canis, C. felis, Echidnophaga gallinacaea, Pulex irritans

Vulpes: Ctenocephalides canis, C. felis, Euhoplopsyllus glacialis, Megabothris abantis, Oropsylla

bruneri, Pulex irritans, P. simulans, Thrassis bacchi

Table 2. Potential plague vectors infesting Mustelidae and Procyonidae. Compiled from a list of 50 potential flea vectors from the U.S. that have been found to be naturally infected with plague (Pollitzer 1961 and CDC unpublished data).

MUSTELIDAE

Gulo: Euhoplopsyllus glacialis, Oropsylla idahoensis

Martes: Aetheca wagneri, Ceratophyllus ciliatus, Epitedia wenmanni, Euhoplopsyllus glacialis, Foxella ignota,

Hystrichopsylla dippei, Nosopsyllus fasciatus, Orchopeas sexdentatus, Thrassis stanfordi

Mephitis: Ceratophyllus ciliatus, Echidnophaga gallinacaea, Eumolpianus eumolpi, Hystrichopsylla dippei,

Orchopeas sexdentatus, Oropsylla montana, Pulex irritans, Pleochaetis sibynus, Thrassis acamantis

Mustela: Aetheca wagneri, Catallagia decipiens, C. wymani, Ceratophyllus ciliatus, Ctenocephalides felis, Epitedia wenmanni, Euhoplopsyllus glacialis, Eumolpianus eumolpi, Foxella ignota, Hoplopsyllus anomalus,

Hystrichopsylla dippei, Malaraeus telchinum, Megabothris abantis, Megarthroglossus divisus, Nosopsyllus fasciatus, Neopsylla inopina, Orchopeas leucopus, O. sexdentatus, Oropsylla bruneri, O. hirsuta, O. idahoensis, O. labis, O. montana, O. rupestris, O. tuberculata chin Tusto, O. t. tuberculata,

Peromyscopsylla hesperomys, Pulex irritans, Thrassis acamantis, T. bacchi, T. pandorae, T. stanfordi

<u>Spilogale</u>: Anomiopsyllus nudatus, Atyphloceras echis, Ceratophyllus ciliatus, Ctenocephalides canis, Ctenocephalides felis, Hoplopsyllus anomalus, Orchopeas sexdentatus, Oropsylla montana

Taxidea: Echidnophaga gallinacaea, Hystrichopsylla dippei, Pulex irritans, Pulex simulans, Oropsylla bruneri, O.

idahoensis, O. labis, O. rupestris, O. tuberculata cynomuris, O. t. tuberculata, Thrassis acamantis, T.

bacchi, T. pandorae

PROCYONIDAE

Procyon: Ceratophlyllus ciliatus, Ctenocephalides felis, Pulex irritans

Bassariscus: Echidnophaga gallinacaea, Orchopeas leucopus, O. sexdentatus

Table 3. Potential plague vectors infesting Onychomys leucogaster. Compiled from a list of 50 potential flea vectors from the U.S. that have been found to be naturally infected with plague (Pollitzer 1961 and CDC unpublished data).

Aetheca wagneri, Anomiopsyllus spp., Catallagia decipiens, Echidnophaga gallinacaea, Epitedia stanfordi, E. wenmanni, Euhoplopsyllus glacialis, Foxella ignota, Hoplopsyllus anomalus, Malaraeus telchinum, Megarthroglossus divisus, Meringis shannoni, Opisodasys keeni, Orchopeas leucopus, O. sexdentatus, Oropsylla hirsuta, O. idahoensis, O. labis, O. montana, O. tuberculata cynomuris, O. t. tuberculata, Peromyscopsylla hesperomys, Pleochaetis exilis, Pulex irritans, Thrassis bacchi, T. francisi, T. pandorae, T. petiolatus

additional cases occurred in individuals handling infected rabbits (Sylvilagus spp.) and another person acquired plague while skinning an infected antelope (Antilocapra americana). The remaining 24 cases were associated with handling infected carnivores (Table 4). Fifteen of these cases occurred in pet owners or veterinary personnel exposed to infected domestic cats. Two cases occurred in members of a New Mexico Pueblo tribe who killed and skinned an infected gray fox to obtain its pelt for use in a religious ceremony. The remaining seven carnivore-associated cases occurred in hunters and trappers skinning infected furbearers.

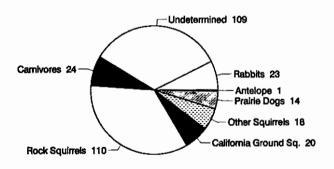


Figure 2. Mammalian sources of infection for human plague cases. Animals were considered to be sources of infection when person became infected after handling an infected member of that species or when a certain species of animal was thought to be the most likely source of infected fleas for a given case.

Table 4. Carnivores as sources of human plague infection in the United States (1970-1993) (CDC unpublished records).

Source	Total number of cases
Felis catus	15
Lynx rufus	4
Canis latrans	2
Urocyon cinereoargenteus	2
Taxidea taxus	_1
Total	2 4

Mortality for the 24 cases associated with carnivores was higher (25.0% of cases fatal) than for the other 295 cases (13.6% of cases fatal) reported from 1970 through 1993. This increased mortality is undoubtedly related to the disproportionately high percentage of carnivoreassociated cases that are either primary septicemic plague or primary pneumonic plague (20.8% and 16.7% of 24 cases, respectively). Among the 295 cases where carnivores were not identified as sources of infection, only 12.2% were primary septicemic plague and less than 1% were primary pneumonic plague. Primary septicemic and primary pneumonic plague are both considered more dangerous than the bubonic form of the disease because they progress more rapidly and are more difficult to diagnose (Poland and Barnes 1979). The higher incidence of primary septicemic plague among carnivore-associated cases is due to persons cutting themselves while skinning animals or exposing existing cuts and abrasions to plagueinfected body fluids or tissues. Persons skinning potentially infected carnivores or game animals always should wear thick latex gloves to protect themselves from these exposures. Each of the four carnivore-associated primary pneumonic cases were exposed while handling infected domestic cats. These animals also have been sources of infection for bubonic and primary septicemic cases, and persons should be cautious when handling sick cats from plague-endemic areas.

Pets also can increase human plague risks by bringing infected fleas into homes (Pollitzer 1954, Archibald and Kunitz 1971). Studies in the U.S. have demonstrated that dogs frequently are infested with a variety of fleas, including those potential plague vectors listed in Table 1. Archibald and Kunitz (1971) provided direct evidence that pets can transport infected fleas into human dwellings. These authors collected infected Pulex irritans from a plague patient's dog on the Navajo Reservation in Arizona (Archibald and Kunitz 1971). Although P. irritans is a relatively poor plague vector, dogs in the Southwest plague focus also have been found infested with important plague vectors, such as Oropsylla montana (CDC unpublished data). Persons living in plague-endemic areas should treat pets with insecticidal flea powders and prevent these animals from roaming free in areas where they are likely to be exposed to plague-infected animals or fleas. Trappers and hunters handling infected

furbearers also could be exposed to infected rodent fleas, although this has yet to be demonstrated to be a source of infection.

PREDATORS AS SENTINEL HOSTS

Analyzing serum samples from mammalian predators, especially carnivores, that consume rodent prey is a powerful means of detecting Y. pestis in rodent populations (Rust et al. 1971, Cruickshank et al. 1976, Willeberg et al. 1979, Taylor et al. 1981, Barnes 1982, Hopkins and Gesbrink 1982, Smith et al. 1984, Nelson et al. 1985, Isaacson 1986, Clover et al. 1989). Mammalian predators that survive plague infection often develop high antibody titers that persist for as long as four to eight months (Barnes 1982). Sampling even a few rodentconsuming carnivores, such as coyotes, can be roughly equivalent to sampling hundreds of rodents for evidence of plague infection. Carnivore serosurveillance has been used extensively since the 1970's to document the occurrence of plague throughout much of the western U.S., including recent (1992-1993) identifications of epizootic activity in counties of eastern Montana, western North Dakota, and western Nebraska where plague has not been identified previously (CDC unpublished data). The samples collected from western Nebraska represent the first identifications of plague in that state since surveillance activities began in the early 1900s. Carnivore serosurveys are especially recommended when: 1) vast areas must be sampled, 2) investigators want to determine if plague has spread to areas where it has not been detected previously, or 3) plague has not been observed in local rodent populations for many years and it is suspected that the disease may have disappeared from the region.

In North America, the carnivores that have been used most frequently for serosurveillance are domestic dogs, coyotes, and badgers (Rust et al. 1971, Willeberg et al. 1979, Barnes 1982, Hopkins and Gesbrink 1982, Messick et al. 1983, Smith et al. 1984, Nelson et al. 1985). Seropositive foxes, bears, weasels, skunks, martens, fishers, raccoons, ring-tailed cats, domestic cats, bobcats, and mountain lions also have been reported (Barnes 1982, Smith et al. 1984, Zielinski 1984, Clover et al. 1989, and unpublished CDC records).

When carnivores in plague-endemic areas are sampled during interepizootic periods, it is typical to find a small percentage (often < 5%) of animals seropositive at any given time. The percentage of seropositives usually increases dramatically, however, during, or immediately following, an epizootic. Such an increase in seropositives should serve as an early warning of increased plague risks for humans. Studies on the Navajo Reservation in the southwestern U.S. have demonstrated that serosurveillance of domestic dogs is a sensitive indicator of both increased epizootic activity and increased human plague risks. When the percentage of seropositive dogs increased significantly, the level of epizootic activity in local rodent populations also was greater than normal. Increased numbers of canine seropositives and increased epizootic activity also corresponded with increased numbers of human plague cases during these years (Barnes 1982).

Although most serosurveillance programs concentrate carnivores, grasshopper mice also might be appropriate sentinel hosts in some situations. During a 1984 plague epizootic in Weld County, Colorado, fleas were collected from five species of rodents (Spermophilus tridecemlineatus, Perognathus hispidus, Dipodomys ordii. Peromyscus maniculatus and Onychomys leucogaster). None of the rodents collected were seropositive for plague, but Y. pestis was isolated from four species of fleas collected from these same rodents (Meringis parkeri, Pleochaetis exilis, Thrassis fotus, Meringis hubbardi). Among these four flea species, only M. hubbardi (one flea collected from S. tridecemlineatus) was not collected from O. leucogaster. More than 450 fleas were collected the following year (1985) from these same rodent species, but none were plague positive. Moreover, among 85 rodent sera tested only seven were positive for plague antibody. Each of these seven positive samples were from O. leucogaster. If only grasshopper mice had been used for plague surveillance during this epizootic, all but one infected flea species would have been identified, and serologic evidence of the epizootic would have been observed.

Serosurveillance of predators will continue to be an essential part of plague surveillance programs in the U.S. because the technique is both highly sensitive and costeffective. The existing serosurveillance program is a collaborative effort involving CDC, state, and local public health officials, as well as animal damage control specialists, wildlife biologists, and others who work with wild or domestic carnivores or other suitable sentinel hosts. Unfortunately cost considerations limit the amount of serological testing that can be done at CDC, but we encourage participation in the program. Samples for these serosurveys can consist of either serum separated from whole blood or drops of blood dried on special paper sampling strips (Nobuto strips) (Wolff and Hudson Nobuto strips are especially useful for field workers who do not have access to centrifuges or refrigeration. Another advantage is that only about 0.2 ml of blood are required to coat a Nobuto strip with sufficient blood for serological testing. After the strips have been coated with blood, they are allowed to dry before being placed in mailing envelopes that have the appropriate collection data written on the outside. These mailing envelopes are then sent to a central laboratory, such as CDC, for processing. The small amount of blood required for testing can be collected from dead carnivores by opening the body cavity or cutting a large vein. Domestic dogs can be bled from the cephalic vein in the leg without adverse effects, although they should be properly restrained and muzzled to protect both the dog and its handlers. CDC can provide both mailing envelopes and Nobuto strips for appropriate serosurveillance projects.

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