

# Flea-Borne Transmission Model To Evaluate Vaccine Efficacy against Naturally Acquired Bubonic Plague

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**A flea-to-mouse transmission model was developed for use in testing new candidate vaccines for the ability to protect against flea-borne plague. The model was used to evaluate a recombinant fusion protein vaccine consisting of the *Yersinia pestis* F1 and V antigens. After one to three challenges with *Y. pestis*-infected fleas, 14 of 15 unvaccinated control mice developed plague, with an average septicemia level of  $9.2 \times 10^8$  *Y. pestis* CFU/ml. None of 15 vaccinated mice developed the disease after similar challenges, and serological testing indicated that transmitted bacteria were eliminated by the immune system before extensive replication and systemic infection could occur. The transmission and development of disease in control mice correlated with the number of bites by blocked fleas but not with the total number of fleabites. The model provides a means to directly assess the efficacy of new vaccines to prevent naturally acquired bubonic plague and to study events at the vector-host interface that lead to dissemination and disease.**

*Yersinia pestis*, the bacterial agent of plague, remains an international public health concern. Recent plague outbreaks in India and parts of East Africa, where the disease had been dormant for decades, have raised fears of a new resurgence (6, 33). More troubling was the isolation of multidrug-resistant strains of *Y. pestis* during the ongoing plague epidemic in Madagascar (10, 12). Added to these concerns is the recognized potential of *Y. pestis* as a bioterrorism agent (19). The potential threat of plague outbreaks caused by naturally occurring or deliberately released antibiotic-resistant strains increases the urgency of possessing an effective plague vaccine, but none is available; the previously used killed whole-cell vaccine is no longer being produced.

*Y. pestis* is transmitted primarily by fleabite among its many rodent reservoir hosts. Transmission by direct contact or ingestion of infected tissues can occur in some cases, but the maintenance of plague in its natural environment is thought to depend on rodent-flea-rodent transmission cycles, and most human cases of plague also result from fleabites (21). During the first week after transmission, the bacteria disseminate from the peripheral fleabite site to the regional lymph nodes and produce a severe acute lymphadenitis, characterized by a swollen, painful lymph node called a bubo. Bubonic plague can be treated successfully with an appropriate antibiotic. If not treated, however, a fulminant, high-density, and usually fatal septicemia rapidly develops. Hematogenous spread to the lungs causes secondary pneumonic plague in ~5% of human cases (5).

New second-generation bivalent recombinant plague vaccines consisting of the *Y. pestis* F1 capsular and V antigens have shown promise in trials using rodent and primate animal mod-

els (13, 20). To evaluate protection against bubonic plague, vaccinated animals were challenged by subcutaneous (s.c.) or other parenteral injection of bacterial suspensions prepared from in vitro-grown cultures. There are important differences between this challenge model and natural challenge by fleas, however. To produce a transmissible infection in the flea, *Y. pestis* blocks the proventriculus, a valve that connects the esophagus to the midgut, by forming a large aggregate that is embedded within an extracellular matrix. When these blocked or partially blocked fleas attempt to feed, blood containing *Y. pestis* from the blocking mass is refluxed into the bite site (3, 15). It is in this flea-specific context, which is not duplicated by artificial challenge models, that *Y. pestis* exits the flea and enters the mammal.

For other arthropod-borne pathogens, it is known that the route of transmission can affect the initial interaction with the host immune system and vaccine efficacy. For example, the saliva of blood-feeding arthropods contains immunomodulatory components that inhibit complement activation, phagocytosis, T-cell proliferation, and cytokine secretion (24, 26). Notably, a vaccine that fully protected mice against needle-injected *Plasmodium* sporozoites was much less protective against mosquito challenge (31). Because of the unique phenotype and microenvironment of *Y. pestis* present at transmission and its potential to influence the immune response, we developed a model to explicitly evaluate the abilities of candidate plague vaccines to protect against fleabite challenge.

## MATERIALS AND METHODS

**Animal immunization.** Outbred, immunocompetent, hairless Crl:SKH1-*hr*BR mice (Charles River Laboratories, Wilmington, Mass.) were used in this study. Hairless mice were chosen because infected fleas could be more easily recovered from them after challenges. The F1-V recombinant fusion protein vaccine developed at the United States Army Medical Research Institute of Infectious Diseases (13) was evaluated in the flea challenge model. Fifteen 11-week-old female mice were immunized by s.c. injection of 0.2 ml of phosphate-buffered saline (PBS) containing 10  $\mu$ g of recombinant F1-V fusion protein adsorbed to

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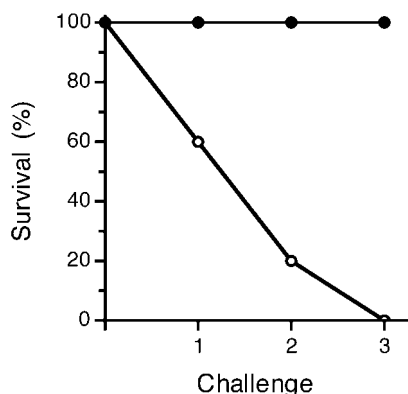


FIG. 2. Survival rates of vaccinated mice (solid circles) and control mice (open circles) after consecutive challenges with infected fleas. One control that survived two challenges (mouse C11 [Table 1]) was not given a third challenge.

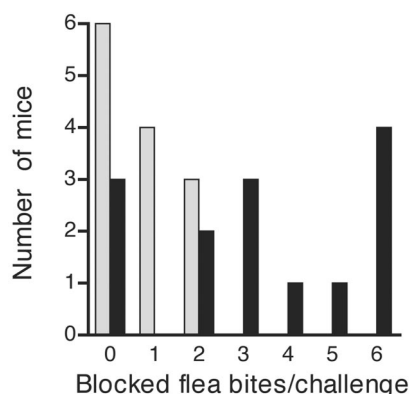


FIG. 3. Frequency distribution of numbers of control mice that survived (shaded bars) or developed disease (solid bars) after individual challenges that included zero to six bites by blocked fleas.

one surviving control mouse was challenged twice and had been fed upon by 47 fleas, 3 of which were blocked. Despite the inherent flea-to-flea variability in transmission, development of disease correlated with the number of bites by blocked fleas ( $P < 0.05$ ) but not with the total number of fleabites ( $P = 0.46$ ) per challenge (Fig. 3). When plague did occur, it developed rapidly in the control mice (50 to 101 h after challenge). The time to onset of symptoms did not correlate with either the total number of fleabites ( $P = 0.37$ ) or the number of bites by blocked fleas ( $P = 0.60$ ). Following euthanasia, *Y. pestis* levels in the peripheral blood, the spleen, or both tissues were determined for each symptomatic mouse. The average septicemia was  $9.2 \times 10^8$  *Y. pestis* CFU/ml (range,  $2.2 \times 10^4$  to  $5.6 \times 10^9$ ; median,  $2.4 \times 10^8$ ), and the average bacterial burden in the spleen was  $6.9 \times 10^9$  *Y. pestis* CFU (range,  $6.0 \times 10^4$  to  $4.0 \times 10^9$ ; median,  $1.0 \times 10^8$ ).

**Protective efficacy of F1-V vaccine.** The 15 vaccinated mice were challenged concurrently with the control mice, using the same cohorts of infected fleas. Vaccinated mice received two

or three challenges until each had experienced a minimum of four bites by blocked fleas and on average were challenged with more total bites and more bites from blocked fleas than the controls (Table 2). None of the vaccinated mice developed disease or showed any symptoms. The 15 vaccinated mice and the one surviving control mouse were euthanized 4 to 5 weeks after the final challenge. Cultures of blood and spleen from the control mouse, and of spleens from the 15 vaccinated mice, were all negative.

Pre- and postchallenge sera from three surviving vaccinated mice were used for immunoblots of whole-cell *Y. pestis* lysates. The sera used for Fig. 4 were from mouse V1 (Table 2), which was bitten by 24 fleas in the first challenge, none of which were blocked; by 52 fleas, 5 of which were blocked, in a second challenge 5 weeks later; and by 31 fleas, 6 of which were blocked, in the final challenge 2 weeks after the second. Thus, this mouse experienced bites from 11 blocked fleas, indicating a high probability of transmission. As expected, the paired sera both reacted with V and F1 antigens, the vaccine components. Few additional *Y. pestis* antigens were recognized in the post-challenge sera, collected 4 weeks after the final challenge.

TABLE 1. Incidence of plague in control mice after fleabite challenge

Mouse	No. of challenges	Total no. of fleabites (cumulative)	No. of blocked fleabites (cumulative)	Time to onset of symptoms (h)
C1	1	12	0	101
C2	2	51	3	55
C3	2	56	3	50
C4	1	13	0	53
C5	1	18	3	56
C6	1	22	5	77
C7	1	24	2	74
C8	1	16	0	74
C9	3	79	6	49
C10	3	87	8	75
C11	2	47	3	Survived
C12	2	61	5	99
C13	2	55	5	66
C14	2	58	6	72
C15	3	96	7	56
Mean $\pm$ SD	1.8 $\pm$ 0.8	46.3 $\pm$ 27.8	3.7 $\pm$ 2.5	68 $\pm$ 17

TABLE 2. Protection of vaccinated mice against fleabite challenge

Mouse	No. of challenges	Total no. of fleabites (cumulative)	No. of blocked fleabites (cumulative)	Time to onset of symptoms (h)
V1	3	107	11	Survived
V2	3	97	7	Survived
V3	3	101	6	Survived
V4	3	87	8	Survived
V5	2	67	6	Survived
V6	2	45	4	Survived
V7	2	43	11	Survived
V8	3	91	13	Survived
V9	3	69	7	Survived
V10	2	66	7	Survived
V11	3	103	13	Survived
V12	2	61	6	Survived
V13	2	60	5	Survived
V14	2	75	9	Survived
V15	2	60	4	Survived
Mean $\pm$ SD	2.5 $\pm$ 0.5	75.5 $\pm$ 20.9	7.8 $\pm$ 3.0	

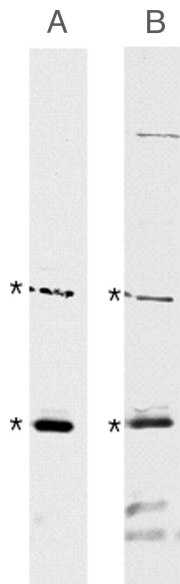


FIG. 4. Anti-*Y. pestis* antibodies present in prechallenge (A) and postchallenge (B) paired sera from vaccinated mouse V1 (Table 2) that survived three challenges with infected fleas over a 7-week period. Immunoblots of whole-cell lysates of *Y. pestis* grown at 37°C are shown. The asterisks indicate the positions of the 37-kDa V antigen and the 17.6-kDa F1 antigen.

## DISCUSSION

Potential new vaccine formulations designed to prevent disease caused by arthropod-transmitted pathogens are not often tested by natural infection routes, but there are good reasons for doing so. Transmission and subsequent infection caused by these agents depends on a complex coevolved interaction among the pathogen, vector, and vertebrate host that has not been well characterized for any arthropod-borne disease (23). However, vector-specific attributes are known to influence the outcome of infection, immune response, and vaccine efficacy for some arthropod-borne pathogens. For example, a component of sandfly saliva greatly enhances the infectivity of *Leishmania*, and normal infectivity of an arbovirus depends on both vector saliva and host factors at the bite site (8, 30). The 50% infective dose of *Plasmodium berghei* sporozoites recovered from infected mosquitoes and injected intravenously has also been shown to be 10 to 100 times greater than that of sporozoites delivered by mosquito bite. More importantly from the standpoint of vaccinology, the neutralizing ability of protective antibody and vaccine efficacy were greatly reduced in mosquito-borne compared to needle and syringe transmission models (31). Because the phenotype of the pathogen in the vector and the microenvironment of the deposition site in the skin are usually unknown and impossible to duplicate by needle and syringe inoculation of artificially cultured organisms, it seems prudent to include the natural transmission route in evaluating the efficacies of potential vaccines for arthropod-borne pathogens. Relying on artificial challenge alone may rule out candidate vaccines that may be perfectly adequate in the natural situation or falsely sanction others that are not, even though artificial challenge models indicate efficacy.

Previous trials of the recombinant F1-V fusion protein vac-

cine used in this study have demonstrated complete protection for vaccinated mice challenged by s.c. injection with  $>10^6$  wild-type *Y. pestis* organisms, a dose which is  $10^5$  times greater than the 50% lethal dose (13). Few data are available that address the number of *Y. pestis* organisms transmitted per fleabite, but an estimate of 11,000 to 24,000 bacteria transmitted by a single blocked *X. cheopis* flea has been reported (4). From the standpoint of sheer numbers, therefore, the challenge inocula delivered by the natural flea-borne route of transmission can be easily matched experimentally by needle injection, but factors specific to the bacteria-vector-host transmission triad are not. Arthropod-borne bacterial pathogens exhibit a specific phenotype in their vectors and produce vector-specific antigens. For example, *Borrelia burgdorferi* spirochetes transmitted by ticks display different surface antigens and induce a different immune response than do in vitro-grown, needle-inoculated spirochetes (11, 27). *Y. pestis* upregulates the expression of certain transmission-related genes in its flea vector that are downregulated or not expressed in the mammal (14, 21, 28). These genes enable *Y. pestis* to grow in a flea-specific biofilm-like phenotype in which a dense bacterial aggregate embedded within an extracellular matrix blocks the proventriculus (15, 18). The composition of the extracellular matrix associated with the biofilm-like growth of *Y. pestis* in the flea is unknown, but it is complex and appears to contain flea gut components in addition to bacterially derived material (B. J. Hinnebusch, unpublished data). The extracellular matrix associated with *Y. pestis* as it exits the flea and enters the mammal might provide initial protection against uptake or killing by phagocytes, as has been demonstrated for the extracellular matrices of other bacterial biofilms (7).

The feeding mechanism of fleas determines the initial site of infection and the ensuing dissemination route of *Y. pestis*. The mouthparts of *X. cheopis* are  $\sim 400$   $\mu\text{m}$  in length, which precludes s.c. injection. Flea saliva introduced into the intradermal bite site contains apyrase, an enzyme which acts to inhibit platelet and neutrophil aggregation, and may contain other immunomodulatory factors, as is true for the saliva of other blood-feeding arthropods (24–26).

Because of the unique phenotype and microenvironment in which *Y. pestis* is presented to the host by fleas, and its potential to affect immune surveillance and vaccine-induced protection, we developed a model to evaluate the abilities of plague vaccines to protect against natural challenge. Our results provide direct evidence that the F1-V recombinant fusion vaccine is able to protect against bubonic plague transmission in a real-world context. In this trial, the vaccine proved 100% effective in preventing flea-borne plague in mice after they had received challenges greater than those which resulted in disease in adjuvant-only control mice. Immunoblots of pre- and postchallenge sera demonstrated little evidence of seroconversion to other *Y. pestis* antigens in the vaccinated mice (Fig. 4), indicating that immunization eliminated the bacteria before they could extensively replicate, produce a disseminated infection, and stimulate a generalized immune response.

Several factors contributed to establishing a practical model. The mouse strain used was highly susceptible to plague and rapidly developed terminal septicemia, making it a sensitive animal model. The major disadvantage of natural challenge is that the dose cannot be controlled. Thus, an important aspect



of the model was to define the parameters of a sufficient challenge. Transmission by fleas is inherently variable, as has been noted consistently by previous investigators. Nearly half of the fleas clear themselves of infection even after feeding on a highly septicemic blood meal, and less than half of the fleas that are successfully colonized by *Y. pestis* develop proventricular blockage (4, 17, 22). Then, individual blocked *X. cheopis* fleas that attempt to feed on a susceptible host transmit plague only ~50% of the time (4), despite the fact that blocked *X. cheopis* fleas contain  $10^5$  to  $10^7$  *Y. pestis* organisms (16). The use of hairless mice made it feasible to simultaneously challenge with large numbers of infected fleas, which helped to overcome individual flea-to-flea variability and to increase the probability of transmission. The data reinforce the importance of the proventricular-blockage phenomenon in transmission by *X. cheopis*. Successful transmission to control mice correlated strongly with the number of bites by blocked fleas but not with the total number of fleabites (Fig. 3). Complete proventricular blockage was not absolutely required, however, because 3 of 15 (20%) control mice acquired plague even though none of the challenge fleas appeared to be blocked. This could be due to missed diagnosis of blockage or transmission by partially blocked fleas, which are also considered to be transmission competent (2). In our model, the vaccinated mice experienced bites from a minimum of four blocked fleas to ensure a high probability of transmission.

The natural challenge model provides a new tool to evaluate future second- and third-generation plague vaccines. In addition, very little is known about the early events after flea-borne transmission that lead to systemic invasion and disease. It has been hypothesized that *Y. pestis* is ingested and transported from the dermis to the regional lymph nodes by macrophages (29), but this has not been examined directly. The vector-borne transmission model should be useful in investigating how the flea-specific phenotype of *Y. pestis* and the microenvironment of the fleabite site influence the initial encounter with the host.

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